

Genex Kidston Connection Project: Draft Environmental Assessment Report Powerlink Queensland

# Appendix J

# **Corridor Selection Report**



# PROPOSED GENEX KIDSTON CONNECTION PROJECT

# **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"



Version: 6.0

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# 1 PREFACE

Genex Power (Genex) is seeking to establish a combination solar and hydro pump storage power generation facility in the Far North Queensland region at Kidston, approximately 270km north-west of Townsville. The Kidston Solar Project is proposed to be co-located with the Kidston Pumped Storage Project and will be built on the tailings storage facility of the old Kidston gold mine.

Stage one of the project involves installation of a solar photo voltaic farm with a capacity of 50MW. Once a new transmission line has been built to the site, stage two of the solar project will be developed, which will expand the total capacity of the solar farm to 270MW<sup>1</sup>. Additionally, the Hydro Pump Storage Scheme seeks to have a generation capacity of 250MW<sup>2</sup>, providing a cumulative peak total generation capacity in excess of 500MW.

Powerlink has been commissioned by Genex to prepare a Corridor Selection Report (CSR), which identifies a preferred corridor and a preliminary alignment for the construction and operation of a new 275kV transmission line to connect the Kidston facility to Powerlink's existing transmission network, with a connection point being located near the existing Mt Fox communications site along the Ross to Chalumbin 275kV transmission line.

The goal of the CSR is to identify a preferred study corridor within the study area that, on balance, offers the most appropriate location for the proposed 275kV transmission line, taking into account social, economic and environmental factors. The preferred study corridor will be broad and of variable width. The CSR also identifies a preliminary alignment for the potential transmission line within the study corridor to guide further investigation and consultation in subsequent project stages as required.

Should the project to develop a new transmission line proceed, a comprehensive Environmental Impact Assessment and associated stakeholder and landholder consultation process would be required before any final decision on an alignment could be taken. Future consultation would build on the work undertaken and relationships established in the development of this CSR.

<sup>1</sup> Genex Power, The Kidston Solar Project, 2017, <u>http://www.genexpower.com.au/the-kidston-solar-project.html</u> (accessed 9 Jan 2017) <sup>2</sup> Genex Power, The Kidston Hydro Project, 2017, <u>http://www.genexpower.com.au/the-kidston-hydro-project.html</u> (accessed 9 Jan 2017)

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# 2 GLOSSARY

# 2.1 TERMS

Acronym	Definition
AHD	Australian Height Datum
ASS	Acid Sulfate Soils
CID	Community Infrastructure Designation
CLR	Contaminated Land Register
CSR	Corridor Selection Report
DCSR	Draft Corridor Selection Report
EIS	Environmental Impact Statement
EMR	Environmental Management Register
IDAS	Integrated Development Assessment System
km	Kilometres
kV	Kilovolts
LGA	Local Government Area
m	Metres
MNES	Matters of National Environmental Significance
MVA	Mega Volt Amp
MW	Megawatts
QLD	Queensland
RE	Regional Ecosystem
ToR	Terms of Reference

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# 2.2 ORGANISATIONS / AGENCIES

Acronym	Definition
Genex	Genex Power
Powerlink	Powerlink Queensland
DAF	Department of Agriculture and Fisheries
DEWS	Department of Energy and Water Supply
DEHP	Department of Environment and Heritage Protection
DNPSR	Department of National Parks, Sport and Racing
DNRM	Department of Natural Resources and Mines
DTMR	Department of Transport and Main Roads
Ergon	Ergon Energy
OCOG	Office of Coordinator General
BQ	Building Queensland

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# 3 EXECUTIVE SUMMARY

This report has been prepared in response to a request from Genex Power (Genex) to identify a suitable study corridor for a high voltage electricity transmission connection to its proposed combined solar/hydro pumped storage generation facility at Kidston, approximately 270km north-west of Townsville. To meet the load requirements, the proposed connection solution is a 275kV double circuit transmission line nominally to be constructed using steel towers. A substation with switching capability would be required along Powerlink's existing 275kV Ross to Chalumbin transmission line near Mt Fox (approximately 40km west of Ingham).

This CSR builds on the outcomes of a Draft Corridor Selection Report (DCSR) completed by Powerlink for Genex in May 2016, which assessed three possible corridor options between Mt Fox and Kidston. Based on the outcomes of the DCSR, one study corridor option has been excluded from for further assessment, having the highest potential environmental impact of the options assessed. This CSR therefore explores a study area based generally on the two study corridor options remaining from the DCSR.

In accordance with Powerlink's Project Engagement Model and seeking to achieve best practice project engagement, early and targeted consultation with stakeholders and landholders has been undertaken in development of this CSR. Engagement occurred during late 2016 and early 2017 and has provided meaningful insight into the study area to complement the desktop assessment. The process of undertaking engagement involved extensive travel through the study area, further enhancing Powerlink's understanding of the region. Powerlink also gained access to Ergon aerial survey data and undertook a helicopter fly over to better understand the project area.

The assessment undertaken in this CSR has indicated that study corridor option C is preferred (Figure 3). Study corridor option C is defined largely by existing Ergon transmission lines and reduces overall impact to the environment and landholders by virtue of co-locating linear infrastructure.

This CSR has not identified any constraints that would prevent the development of a potential 275kV connection within study corridor option C. The preference for study corridor option C was also confirmed by third party ecological advice, which assessed option C as having the lowest overall impact on matters of State and Federal environmental significance.

Within study corridor option C a preliminary alignment has been identified, which will serve as the basis for further assessment in subsequent project phases should the project proceed and Powerlink be appointed to undertake this work. The preliminary alignment is a 120m wide band around the existing Ergon transmission lines (60m either side of the existing Ergon assets), allowing flexibility for the future 275kV line to co-locate north or south of the existing Ergon lines on a 60m wide easement. A preference for locating the 275kV line north or south of existing lines is not possible at the current level of investigation.

Between approximately Greenvale and Conjuboy Ergon does not have existing lines. In this area the preliminary alignment is dictated by the location of an existing mining interest and the optimal path through large escarpments. The alignment width is maintained at 120m in this area for consistency and flexibility.

This CSR has also investigated the feasibility of a new substation at Mt Fox. It is outside the scope of this report to acquire a site or site option and the available terrain data is not detailed enough to define a firm site preference. However, initial landholder feedback and preliminary investigations indicate that a suitable site is likely to be available. This CSR therefore defines an area of interest for the substation which would be further assessed through a targeted site selection process supported by a terrain survey and more detailed landholder engagement.

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# 4 DESCRIPTION OF THE PROJECT AREA

The study area commences near Mt Fox, approximately 100km north-west of Townsville and extends in a westerly direction for approximately 180km to the township of Kidston within the Etheridge Shire. The study area intersects three individual local government areas (LGAs) – Hinchinbrook Shire Council, Charters Towers Regional Council and Etheridge Shire Council.

The proposed substation near Mt Fox would likely be located within the Hinchinbrook Shire Council LGA. The nearest town to the substation would be Mt Fox and the nearest population centre would be Ingham (approximately 40km north-east of Mt Fox). The proposed Kidston substation would be located within Etheridge Shire Council with the nearest town being Einasleigh (approximately 40km north of Kidston, population 202). Approximately 115km of the proposed transmission line would be located within Charters Towers Regional Council with the nearest town being Greenvale (population 150).

The topography within the project area ranges from flat low lying land to steep crossings of multiple ranges, including part of the Pelican Range (70km west of Mt Fox) and the Great Dividing Range (100km west of Mt Fox). Elevation throughout the project area generally ranges from 400-800m AHD. Constructability through these areas is a key consideration in the development of feasible options.

Existing infrastructure within the area includes an Ergon 66kV transmission line that runs 100km west from Mt Fox to Greenvale and another Ergon 132kV transmission line that runs from Ross to Kidston. Consideration of co-location opportunities with existing infrastructure corridors is a key consideration in the development of study corridor options within this report.

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# 5 APPROACH AND METHODOLOGY

# 5.1 CORRIDOR DEVELOPMENT PROCESS

This CSR is one step in a broader development process for high voltage transmission projects which in turn is a subset of the project development for the combined hydro / solar at Kidston. The general development process is outlined in Figure 1.



Figure 1 – Corridor Development Process

This CSR falls generally into steps 2 to 4 of Figure 1. It is important to note that Figure 1 represents a general process flow and the development process is tailored to individual projects.

In the case of Genex, the development processes is being undertaken in a staged manner. Each stage of the process represents a refinement of project scope and risk; allowing an opportunity to reassess the feasibility of the broader development. Sections 5.1.1 to 5.1.5 describe the stages of the corridor development process to date, including specific amendments to the study area during the development of this CSR. It is important to describe the approach to corridor development and refinement up front.

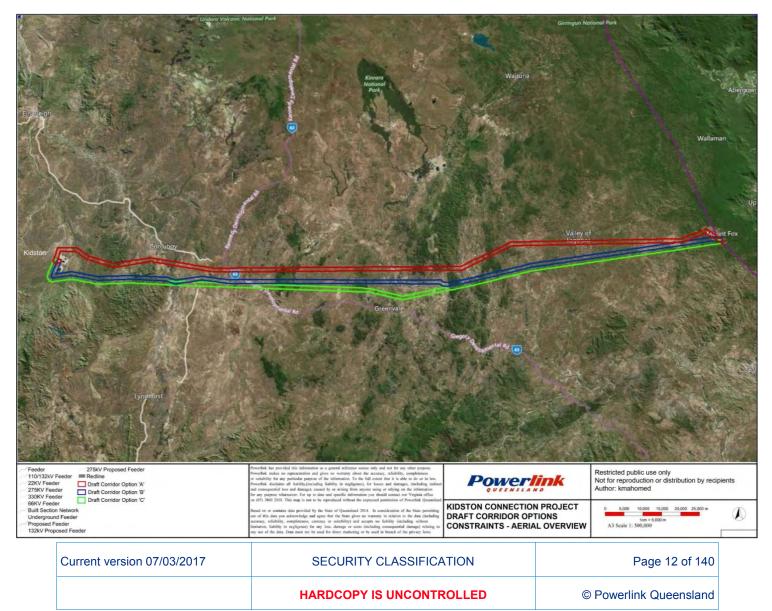
### 5.1.1 DRAFT CORRIDOR SELECTION REPORT

The DCSR was the first major step in the development process for Genex. The DCSR was completed in May 2016 and utilised publicly available datasets to identify three high level study corridor options for further assessment. The three study corridor options identified in the DCSR are shown in Figure 2.

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Figure 2 – DCSR Study Corridor Options





The three study corridor options were discussed and developed with input from a range of stakeholders, including local and State government. Feedback from these stakeholders ensured the outcomes of the desktop studies in the DCSR included local knowledge and that major opportunities and constraints in the region were identified. The DCSR concluded that study corridor option A was the most constrained and for this reason it was excluded from further assessment as part of this CSR.

Study corridor option A was the most constrained of the three corridors identified in the DCSR due to:

- Intersection of the Mt Fox section of Girringun National Park
- Intersection of the Newcastle Range Oaks Nature Refuge
- Proximity to Mt Fox Primary School
- Proximity to Kidston airstrip
- Highest relative intersection of steep terrain, major watercourse crossings, 'of concern' vegetation and potential habitat for protected species, including cassowary habitat near Mt Fox.

This CSR further explores study corridor options B and C as identified at the DCSR stage.

## 5.1.2 CORRIDOR SELECTION REPORT

The study area for this CSR is broadly defined by the outer bounds of study corridor options B and C from the DCSR, representing an iterative refinement of the study area. The study area for the CSR with study corridors B and C is depicted in Figure 3.

Figure 3 shows the study area expanded beyond the extent of the study corridors from the DCSR between Greenvale and Conjuboy. This is due to an amendment to the study area made during delivery of this CSR. A second amendment was also made at Mt Fox, refining the study area as greater understanding of the region was gained. The amendments are described in Sections 5.1.4 and 5.1.5.

This CSR has been developed based on a range of detailed desktop assessments, interrogation of existing agency data, aerial reconnaissance and stakeholder and landholder engagement. The following matters have been considered and informed by feedback from local representatives and landholders:

- Topography
- Hydrology
- Environment
- Native Title
- Cultural Heritage
- Tenure
- Local Government areas
- Residences
- Proximity to population

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- Land use
- Existing infrastructure
- State planning interests and other State policy.

It is also important to note that because the project development process for Genex is being undertaken in stages, Powerlink's scope of work is to compile the CSR only at this stage. Terms of Reference (ToR), EIS, easement acquisition or construction activities are not currently scoped. Powerlink has significant experience scoping transmission projects and is available to assist Genex with these activities if required.

### 5.1.3 CONNECTION POINT

Irrespective of the corridor selected, a substation with switching capacity near Mt Fox will be required along with a 275kV substation at Kidston. The switching functionality is required to maintain the high reliability of the 275kV network. Without the switching capability, any fault/outage on the Kidston feeder section would remove one of the Ross to Chalumbin feeders from service and severely restrict the power flow and reliability of high voltage electricity supply to Far North Queensland. Mt Fox is considered an appropriate location for the proposed substation as it is the closest viable point to Kidston along Powerlink's existing 275kV Ross to Chalumbin transmission line.

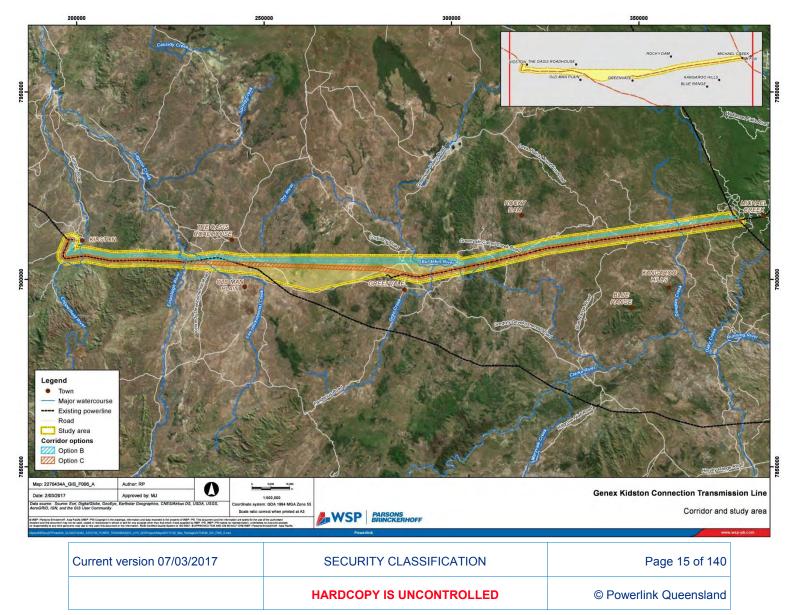
Other locations for the substation were considered, but discounted because they were much further away or were constrained in other ways, such as potential impacts to National Park along Powerlink's existing 275kV Ross to Chalumbin transmission line (including the existing Chalumbin Substation). A study area commencing at Mt Fox also provides opportunity to co-locate with the existing Ergon transmission lines, which most other locations would not.

One of the other options considered was a connection to the existing Ross to Chalumbin line where it meets the existing Ergon 132kV line to Kidston at around Hervey Range. This location does offer the benefit of colocation opportunities, but is up to 40km longer overall than a connection to Mt Fox resulting in potentially greater social and environmental impacts, as well as increased construction costs.

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Figure 3 – CSR Study Area and Study Corridors





# 5.1.4 STUDY AREA AMENDMENT (WEST OF GREENVALE)

During investigations and visits to the proposed study area and with landholders as part of preparing this CSR, it became evident that terrain approximately 25km west of Greenvale is challenging from a construction perspective and that the study area needed to be broadened to offer greater opportunity to avoid these areas.

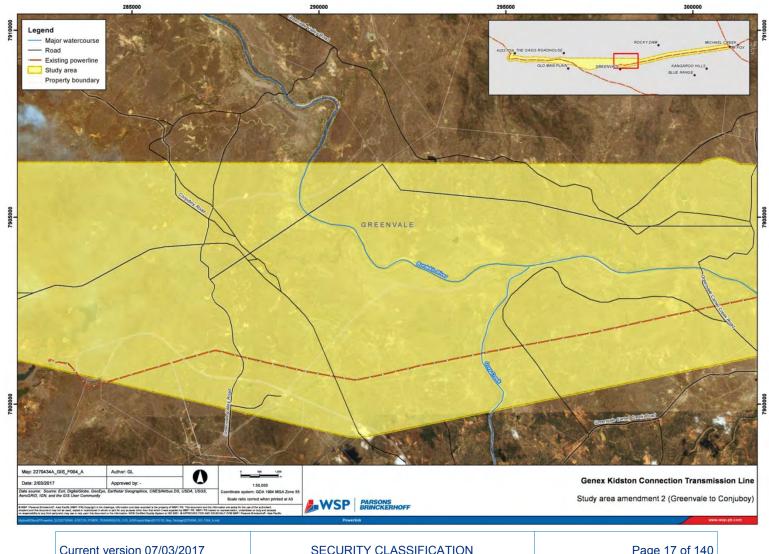
In this location the study area has been broadened significantly, as far south as the Gregory Development Road. Figure 4 shows the expanded study area. Importantly, the extension of the study area did not impact any additional new properties, with potential impacts limited to lots that were already intersected by the previous study area.

A preliminary alignment that avoided the most difficult terrain was also developed. The preliminary alignment deviates south of the initial extent of study corridor option C in this location, responding to the constraints identified in the field before reconnecting to the initial study corridor areas at Conjuboy. The preliminary alignment is discussed in Section 14.2.

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Figure 4 - Study Area Amendment 25km West of Greenvale



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### 5.1.5 STUDY AREA AMENDMENT (MT FOX)

The same refinement process was applied to the proposed study area around Mt Fox. As Powerlink travelled through the area and spoke with stakeholders and landholders it was evident that several of the properties impacted by the initial study area could be removed from the study area. The properties were removed for a range of reasons, including:

- The study area affected only a small part of the property and there was no real potential for Powerlink to identify a suitably sized substation site on the property
- The property backed directly onto Mt Fox Primary School (applies to 46 WU44).

Multiple properties located to the east of Powerlink's existing Ross to Chalumbin 275kV transmission line were also removed from the study area. Several factors contributed to this amendment:

- Additional investigations identified topographic and environmental constraints at Mt Fox that would make extending the line to the east of Powerlink's existing Ross to Chalumbin transmission line more challenging from a construction perspective
- The Liefway Nature Refuge occurred in the area and would not represent a viable location for a substation
- Reducing the size of the study area minimises social impacts
- Several opportunities for a substation site exist in more preferential locations under and west of Powerlink's existing Ross to Chalumbin transmission line.

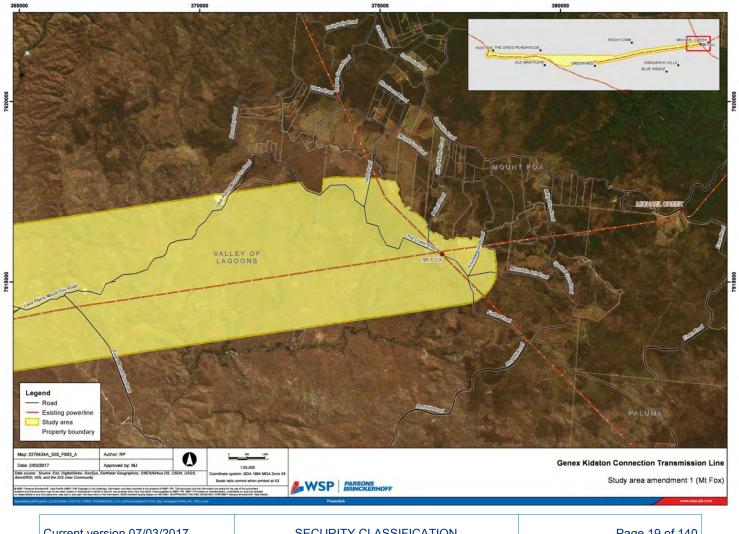
The proposed study area was therefore refined to be coincident with an unnamed creek immediately east of the existing Ross to Chalumbin transmission line. The creek line forms a logical natural boundary for the study area. Avoidance of a creek crossing is also considered beneficial where possible.

Figure 5 depicts the amended study area at Mt Fox. Properties removed from the study area were also contacted by phone and sent a letter to notify them of the change, and offer opportunities for further engagement if desired.

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#### Figure 5 – Amended Study Area at Mt Fox



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### 5.1.6 NEXT STEPS

This CSR will provide the basis for detailed investigations of the preferred study corridor and preliminary alignment by Powerlink in a future impact assessment process should the project proceed.

Genex has multiple process options available to progress the proposed connection project. Environmental and technical field investigations and further opportunities for stakeholder and landowner engagement will be integral parts of the process, irrespective of the approval path chosen.

Powerlink has significant experience in transmission line development in Queensland, including impact assessment, land acquisition, stakeholder and landholder engagement to support Genex in subsequent phases, whichever process is deemed preferable.

# 5.2 LIMITATIONS AND ASSUMPTIONS

This report has been prepared based on the following assumptions:

- The preferred study corridor and preliminary alignment are of variable width and are subject to localised amendment as constraints are further understood in subsequent assessment processes
- No ecological, geotechnical, resistivity, hydrological, water quality or other detailed field based technical assessment has been undertaken
- A 60m wide easement will be required for any future final corridor selected to accommodate a 275kV double circuit transmission line
- Access, laydowns and off easement property requirements are not known at present and therefore not assessable at this level of investigation.

Other assumptions and limitations are provided throughout the document where they are relevant to specific sections.

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# 6 INFRASTRUCTURE REQUIREMENTS

### 6.1 TRANSMISSION LINE

This study has been based on a potential requirement to provide a 275kV double circuit transmission line from Powerlink's existing high voltage network to Genex's generation facility at Kidston. Based on draft corridors developed to date, approximately 188km of transmission line would be required. Configuration of the transmission line would likely be double circuit steel lattice tower design constructed within a new 60m wide easement.

Underground construction is not considered viable where sufficient space for overhead infrastructure exists. Underground construction is not only significantly more costly than overhead construction, but the easement must be kept clear of vegetation. Overhead infrastructure can be carefully sited to span sensitive environmental and cultural features. In the event of a fault it also often takes longer to restore service on underground transmission lines, with fault identification more difficult than equivalent overhead transmission lines.

No network limitations have been identified that would prohibit connection of the generation facility at Kidston. Temporary workers camps, accommodation and offices are also likely to be necessary. The location and scale of those facilities will be defined during subsequent detailed impact assessment processes.

# 6.2 SUBSTATION – MT FOX

A substation with switching capability is necessary for the connection to the Ross to Chalumbin transmission line as it allows for electricity to be transferred to either Ross Substation (south of Mt Fox) or Chalumbin Substation (north of Mt Fox) as required. The switching functionality is required to maintain the high reliability of the 275kV network. Without the switching capability, any fault/outage on the Kidston feeder section would remove one of the Ross to Chalumbin feeders from service and severely restrict the power flow and reliability of high voltage electricity supply to Far North Queensland.

The proposed Mt Fox Substation would be located along Powerlink's existing double circuit 275kV Ross to Chalumbin transmission line. Actual site configurations and technical arrangements will need to be developed as part of any future design and consultation activities, but this CSR discusses high level site opportunities.

# 6.3 SUBSTATION – KIDSTON

A 275kV substation at Kidston, located at the Genex generation site, is required to transform the solar and hydro energy generated to a voltage suitable to transmit to the grid connection point at Mt Fox. Powerlink understands that the Kidston substation will be located on Genex owned land at Kidston, as close as possible to the generation facility.

The study area enters the western side of the Genex owned land at Kidston. Doing so maximises co-location with the existing Ergon transmission lines, increases the distance to residences east of the Genex facility, and avoids a nature refuge north-east of the Genex facility.

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# 7 ENGAGEMENT PROCESS

Powerlink undertook early and targeted consultation with stakeholders and landholders to inform this CSR, leveraging a depth of experience in project engagement. This consultation builds on that initially undertaken for the DCSR and seeks to ensure local knowledge helps inform this CSR.

# 7.1 STAKEHOLDER ENGAGEMENT

This CSR is underpinned by engagement with relevant Federal, State and Local government departments and elected representatives departments within the project area. The objective of stakeholder engagement at this point in the process is to gain meaningful understanding of opportunities, constraints and key issues in the study area.

Table 1 outlines the stakeholder groups consulted. The list of stakeholders is not intended to be definitive or exhaustive and it is highly likely that other stakeholders exist in the region that could add valuable insight to future project development processes as part of future, more detailed planning. Table 1 represents Powerlink's best endeavours during the CSR phase to identify interested stakeholders and seek out local knowledge.

#### Stakeholder FEDERAL GOVERNMENT DEPARTMENTS FEDERAL AND STATE ELECTED REPRESENTATIVES Environment and Energy (DoEE) Mr Andrew Cripps MP, State Member for Hinchinbrook STATE GOVERNMENT DEPARTMENTS/ AGENCIES Mr Shane Knuth MP, State Member for Dalrymple Energy and Water Supply (DEWS) Mr Robbie Katter MP, State Member for Mt Isa Mr Bob Katter MP, Federal Member for Kennedy Treasury PEAK BODIES **Building Queensland** Office of the Coordinator-General (OCOG) **Queensland Farmers Federation** Infrastructure, Local Government and Planning (DILGP) AgForce **Queensland Resource Council** Natural Resources and Mines (DNRM) Environment and Heritage Protection (DEHP) NQ Conservation Council National Parks, Sport and Racing (DNPSR) Northern Gulf Resource Management Group **Townsville Enterprise** Agriculture and Fisheries (DAF) LOCAL GOVERNMENTS **ABORIGINAL PARTIES**

Table 1 – Key	/ Stakeholders
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Stakeholder			
Hinchinbrook Shire Council	Gugu Badhun Aboriginal Corporation		
Charters Towers Regional Council	Tatampi Puranga Aboriginal Corporation		
Etheridge Shire Council			

Powerlink hosted a stakeholder workshop in Townsville on 21 November 2016. The workshop provided an opportunity for stakeholders to collaboratively explore the study area and provide insight on opportunities, constraints and key issues. Powerlink employed the services of an independent facilitator to guide discussion and record the outcomes of the workshop, which are summarised in Section 13.1. The outcomes of the broader stakeholder engagement process are also described in Section 13.2.

# 7.2 LANDHOLDER ENGAGEMENT

This CSR is also underpinned by valuable landholder engagement. All landholders potentially directly impacted by the study area were contacted and offered an opportunity to meet with a Powerlink representative at a time and location most convenient to them. Landholder engagement for the project commenced in late November 2016 and continued into late January 2017.

Landholder engagement at the CSR stage aims to engage the community potentially affected by a proposed project as early as possible. Early engagement provides valuable insights into the local area, affords landholders the opportunity to provide input as early as possible and has resulted in a number of refinements to the study area for the proposed Genex Kidston Connection Project (see Section 5.1).

A consultation blackout period of 20 December 2016 to 5 January 2017 was observed in accordance with Section 298 of the *Sustainable Planning Act 2009*, with the exception of one landholder meeting already organised on 21 December 2016. The blackout period is not considered mandatory for major projects but represents general good practice to minimise disturbance to landholders during the Christmas/New Year period.

More than 30 landholders were initially identified as potentially directly impacted by the initial proposed CSR study area (as defined by the process described in Section 5.1); however iterative refinement of the study area later removed a number of properties around Mt Fox.

All potentially impacted landholders in the study corridor took up the opportunity to meet with Powerlink and provide early input into the CSR. The majority of landholders met with Powerlink at face to face meetings and all known property representatives were able to be contacted by some means.

Key features of the landholder engagement process included:

- The majority of landholders in the study area assisted Powerlink by providing input to the CSR
- The scope of consultation was to seek landholders knowledge and input to the operation of their properties and conditions that could constitute constraints, opportunities and other information relating to the proposed Genex Kidston Connection Project
- Speaking with Powerlink and providing input to the CSR does not indicate landholder support for the project
- Powerlink provided landholders with opportunities to provide input through face-to-face meetings, phone conversations and email in an effort to minimise disruption

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- Most landholders agreed to face-to-face meetings to receive project information and provide their input to the CSR
- Meetings, email and phone were methods used to provide follow up information to landholders in response to requests for further information or responses to questions.

A summary of landholder feedback is presented in Section 13.7. No personal information has been included in this report due to privacy considerations. An overview of key landholder engagement statistics is provided in Table 2.

As a result of engagement, Powerlink also gained an enhanced understanding of the study area, both through local knowledge and by travelling the area extensively. In response to an improved understanding of the region and landholder input, several refinements were made to the proposed study area. These amendments are described in Section 5.1.

Purpose of Engagement Activity	Method of Engagement			Total Number of Engagement
	Face-to-face meeting	Phone contact	Email contact	Activities
Provide project information and obtain landholder input to CSR	14	27	2	43
Follow-up information and follow- up in response to landholder request	0	4	5	9
Make arrangements for future engagement opportunities including meetings	0	27	0	27
Provide advice and obtain landholder input on helicopter flythrough activities	1	8	5	14

#### Table 2 – Key Engagement Statistics

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# 8 RELEVANT BACKGROUND

# 8.1 PREVIOUS PROJECTS

Existing Powerlink infrastructure in the project area includes the 275kV Ross to Chalumbin Transmission Line and Mt Fox Communications Site. Powerlink does not currently have any other new projects in this area.

# 8.2 CURRENT / ONGOING ACTIONS / STUDIES / PROJECTS

It is understood that Genex is well advanced in its planning and preparation for the development the Kidston site. At time of writing, a number of key activities have occurred or are underway:

- The 50MW Kidston Solar Project (Phase 1) has secured funding, is under construction and will connect to Ergon Energy's existing distribution network
- The 250MW Kidston Pumped Storage Hydro Project is currently the subject of a Bankable Feasibility Study
- The 270MW Kidston Solar Project (Phase 2) is currently the subject of technical feasibility studies.

The proposed 275kV transmission line proposal, to which this report relates, is just one element of Genex's overall scheme. A 275kV connection would be required for the development of the Kidston Pumped Storage Hydro Project, as the existing Ergon Energy lines would not have adequate capacity to enable transmission of generated electricity to the National Electricity Market. The proposed 275kV transmission line will facilitate connection of the Kidston Solar Project (Phase 2) to the National Electricity Market if it proceeds.

# 8.3 HELLS GATE AND MT FULLSTOP DAM

Powerlink became aware through the CSR consultation process of a concept plan for a new dam at Hell's Gate, with associated pipeline and treatment infrastructure and management plans. Minimal information regarding the location of the proposed dam is available in the public realm, limited to overview maps that show it as a single point at a regional scale. The maps available also show another potential dam nearby, being Mt Fullstop.

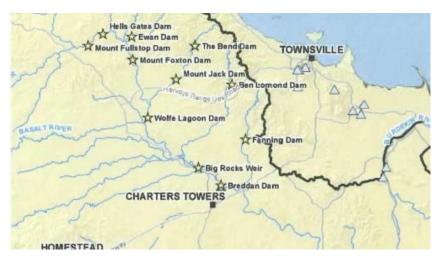
Powerlink understands the concept for the Hells Gate Dam is being investigated by Townsville Enterprise, who has released a tender for a feasibility study which gives a broad indication of the likely location (see Figure 6).

Given the limited information available, it is difficult at this stage to assess the potential for any incompatibilities between the proposals for these dams and the Genex Kidston connection, but further investigations are required to continue monitoring information about the dam's likely size and location.

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#### Figure 6 – Hells Gate Dam Regional Map



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# 9 REGULATORY FRAMEWORK

A range of planning and environmental approvals at the Commonwealth, State and local level will be required to develop the proposed transmission line. Applicable planning and environmental legislation and the approvals likely to be required are discussed in the following sections. The number of applicable legislative instruments is large, but the level of effect of each one varies. An overview of relevant legislation and its likely level of influence is provided in Table 3.

This CSR does not discuss or consider other regulatory approvals which may be required that are not associated with planning or environmental matters, for example investment approvals which may be required under the *Building Queensland Act 2015* or any relevant requirements under the National Electricity Rules.

Regulatory approval obligations vary depending on project value, the proponent and many other factors and will be discussed directly with Genex. Regulatory approvals not associated with planning or environmental matters do not differentiate corridor options and are not discussed further in this CSR.

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#### Table 3 – Approvals Overview

Commonwealth	State	Local
Native Title Act	Sustainable Planning Act 2009/ Planning Act 2016	Etheridge Shire Council
Environment Protection & Biodiversity Conservation Act 1999	State Development & Public Works Organisation act 1971	Charter Towers Regional Council
	Electricity Act 1994	Hinchinbrook Shire Council
	Electrical Safety Act 2002	Local Laws
	Nature Conservation Act 1992	
	Forestry Act 1959	
	Environmental Protection Act 1994	
	Environmental Protection Regulation 2008	
	Environmental Protection (Water) Policy 2009	
	Environmental Protection (Air) Policy 2008	
	Environmental Protection (Noise) Policy 2008	
	Aboriginal Cultural Heritage Act 2003	
	Vegetation Management Act 1999	
	Water Act 2000	
	Land Act 1994	
	Acquisition of Land Act 1967	

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Biosecurity Act 2014	
Stock Route Network Management Bill 2016	
Transport Infrastructure Act 1994	
Fisheries Act 1994	
Regional Planning Interests Act 2014	
Regional Plan	
State Planning Policy	
Infrastructure Plan 2016	
State Development Assessment Provisions	

	An approval under the Act is required/ likely to be required
	A process must be undertaken in accordance with, and to demonstrate compliance with the provisions of the Act
Approval is possibly required (dependant on parameters that are currently unknown/ undecided)	

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## 9.1 COMMONWEALTH

### 9.1.1 ENVIRONMENT PROTECTION AND BIODIVERSITY CONSERVATION ACT 1999

The Environment Protection and Biodiversity Conservation Act 1999 (EPBC 1999) regulates actions that have potential to impact on Matters of National Environmental Significance (MNES). MNES may include fauna and flora species, specific vegetative communities/ assemblages or geographic features. Triggers for assessment under the EPBC 1999 have the potential to occur within each of the two identified study corridor options within this CSR.

Where MNES may be impacted, a proponent refers the project to the Commonwealth Department of Environment & Energy (DoEE) for assessment of potential impacts. DoEE will make a declaration that the project is either:

- Not a controlled action meaning that DoEE has no requirement to assess the project further
- Not a controlled action 'particular manner' meaning that DoEE has no requirement to assess the project further providing that the action is untaken in accordance with conditions that DoEE will supply with the decision
- Controlled action meaning that DoEE wish to assess the project against the EPBC Act. There are several mechanisms available for that assessment dependent on the type of project.

Powerlink has procured third party ecological advice to better understand potential impacts to MNES. The assessment is described in detail in Section 11. A significance assessment will need to be undertaken in subsequent project phases to further refine risks around MNES and to inform a decision around referring the project to DoEE.

#### 9.1.1.1REFERRAL

Powerlink has significant experience and relationships in managing the EPBC referral process and designing solutions to minimise potential impacts to MNES. Powerlink projects have generally not been declared controlled actions due to our ability to optimise infrastructure siting to minimise terrestrial impacts.

Based on Powerlink's experience, it is recommended that referral of the project to the DoEE for further approvals in relation to the EPBC Act should occur early in subsequent stages of the project (e.g. Draft EIS phase).

This approach is recommended as field surveys will be necessary to provide adequate detail for DoEE assessment managers to develop sufficient understanding of any potential issues and, if necessary, develop and apply risk-appropriate controls.

Whilst every reasonable effort has been undertaken to assess potential impacts to MNES flora and fauna at the desktop level for this CSR, the broad scale nature of mapping over the largely undeveloped project area means field surveys will be important in future stages of the project to confirm any potential impacts and allow appropriate mitigation methods to be developed.

Preliminary scheduling by Powerlink indicates that sufficient time is available in Genex's schedule to refer the project and receive a decision under the EPBC Act, even if the project is declared a Controlled Action.

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Importantly, field surveys and a better understanding of the presence of threatened habitats and species, may also support an application for a lower level of assessment, such as a declaration of Not a Controlled Action - Particular Manner.

The optimal approach for the project is therefore to undertake field surveys prior to referring the project to DoEE, ensuring that potential impacts to MNES flora and fauna are understood and a proposal for an appropriate level of assessment under the EPBC Act can be formulated.

### 9.1.2 NATIVE TITLE ACT 1993

Native title recognises the traditional rights and interests to land and waters of Aboriginal and Torres Strait Islander people. Native title can be extinguished or suppressed by certain acts under the *Native Title Act 1993*.

Where Native Title has not previously been extinguished or suppressed, Powerlink typically undertakes a process under Section 24ka of the *Native Title Act 1993* before construction commences. Section 24ka suppresses the Native Title interest in the subject land for the lifetime of the infrastructure to be established and over the extent of the property interest established (easement).

The 24ka process forms part of the wider land/easement acquisition process and involves notification to applicable registered Native Title holders, claimants or suitable alternate parties. The timeframes involved in the process are generally managed within the broader approvals and acquisition schedule.

Notified parties are afforded objection and/ or a claim for compensation rights under the process. A similar process will be undertaken for the substation at Mt Fox, though native title would likely be extinguished over the site which requires exclusive possession rather than an easement.

Powerlink has well-established processes for, and significant experience in the management of Native Title in transmission line development and has established initial contact with the relevant Aboriginal Parties during the development of this CSR. The processes of other parties or organisations are not discussed in this CSR.

# 9.2 STATE

### 9.2.1 SUSTAINABLE PLANNING ACT 2009

The *Sustainable Planning Act 2009* (SPA 2009) coordinates the planning and development system in Queensland. It manages the processes by which most development takes place, the effects of development on the environment, and the coordination of planning and development across local and State Governments.

The SPA 2009 provides the approval process typically used by Powerlink for its transmission lines via the Community Infrastructure Designation (CID) process. It also provides the framework for a range of permits and approvals that may be required in addition to CID for certain activities, and where exemptions do not apply. It is noted that the SPA 2009 is due to be replaced by a new Act in mid-2017, which is discussed in Section 9.2.2.

### 9.2.1.1COMMUNITY INFRASTRUCTURE DESIGNATION (CID)

CID under the SPA 2009 is the typical overarching project approval pathway used successfully by Powerlink for many years. This process is best suited to transmission projects for a number of reasons.

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CID is quite flexible in terms of scheduling, owing principally to the fact that the SPA 2009 does not prescribe timeframes for most parts of the process and that process control is largely with the proponent. Most other processes in contrast are led by an assessment manager who controls the administration of the process in accordance with legislated timeframes.

CID offers a high degree of scalability and also offers some unique benefits, including but not limited to:

- No other approval under the relevant Planning Schemes is required
- If development under a designation is carried out by a public sector entity there are no infrastructure charges applicable
- Exemption from permits under the Vegetation Management Act 1999 for clearing on designated land
- Exemption from some offset requirements by virtue of existing permit exemptions which Powerlink has negotiated over time through consistent demonstration of best practice
- Approval is secured in perpetuity when easements are acquired
- Typically carries few or no conditions.

#### 9.2.1.2INTEGRATED DEVELOPMENT ASSESSMENT SYSTEM (IDAS)

IDAS is the process by which most development in Queensland is approved. Local governments publish Planning Schemes that contain most of the provisions for assessment under the IDAS process. For a project of this nature extending over three local government areas, the CID process is favoured over the IDAS process as it ensures a single, consistent and overarching project approval is achieved.

IDAS permit requirements may still apply for specific aspects of development and would be additional to the CID process. IDAS requirements arise from specific triggers in the SPA 2009, and not from exclusions in the major approvals process. For clarity, the IDAS permits would not be triggered by the relevant Planning Schemes, but by the SPA 2009 directly.

It is not feasible at this stage to define specific IDAS development permit requirements. The exact requirement for permits will depend on the constraints revealed during subsequent investigations and the construction solutions available for specific localities.

It is noted that multiple permits (where required) are typically sought under a single IDAS application. It is also important to note that these development permits are likely to be code assessable only, meaning that they do not require public consultation. Due to the low number of possible IDAS permit triggers and the likelihood that they will be code assessable only there is negligible risk to the projects schedule.

It is also noted that the emerging Planning Act 2016 (Section 9.2.2) will alter the requirement for IDAS permits where designation has been achieved. It is likely that the requirement for some IDAS permits will be removed entirely because the designated project will be made accepted development, exempt from all other permits rather than the planning scheme.

Powerlink has significant previous experience in the IDAS permit process and working closely with local councils.

### 9.2.2 PLANNING ACT 2016

The SPA 2009 is currently under review and as at February 2017, three new Acts have been passed:

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- Planning Act 2016
- Planning and Environmental Court Act 2016
- Planning (Consequential) and Other Legislation Amendment Act 2016.

A range of supporting instruments are currently open for consultation and the new Acts and supporting instruments are expected to come into force in mid-2017. The SPA 2009 will cease on the commencement of the new Acts.

Powerlink has been actively involved in consultation activities relating to the new Acts and supporting instruments and has provided detailed submissions to each stage of the consultation process to ensure it can continue to provide services which facilitate the efficient development of the electricity transmission network.

Certain procedural changes will occur in the CID process when the new Act comes into effect. These changes are however mostly administrative in nature and do not materially affect the broader assumptions behind the CID.

#### 9.2.3 STATE DEVELOPMENT AND PUBLIC WORKS ORGANISATION ACT 1971

The State Development and Public Works Organisation Act 1971 (SDPWOA) provides for State planning and development through a coordinated system of public works organisation, for environmental coordination, and for related purposes.

The SDPWOA provides an alternative approvals pathway for the transmission line project. Powerlink continues to work with Genex and the Office of the Coordinator-General to identify best-for-project approvals solutions.

### 9.2.4 ELECTRICITY ACT 1994

The *Electricity Act 1994* sets out the requirements which all electricity industry participants must follow to ensure a safe, efficient and reliable supply of electricity. It also requires that the supply of electricity is undertaken in an environmentally sound manner. Under Section 31(b) of the *Electricity Act 1994*, a transmission entity is required to properly take into account the environmental effects of its activities under the transmission authority.

The *Electricity Act 1994* and subordinate regulation specifies requirements for construction and operation of the electricity network (which includes substations). The *Electricity Act 1994* also identifies the types of activities associated with the construction and operation of a substation site that are exempt from approval.

Section 112A of the *Electricity Act 1994* makes clearing of native vegetation on freehold land exempt development if the clearing is for operating works for a transmission entity on land designated for the operating works by a Minister under the SPA 2009.

The provisions of the *Electricity Act 1994* apply to each corridor option in the same manner and therefore do not serve to differentiate corridor options in this report.

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#### 9.2.5 ELECTRICAL SAFETY ACT 2002

The *Electrical Safety Act 2002* seeks to prevent through regulation, the death, injury and destruction that can be caused by electricity. Accordingly, the purpose of this Act is to establish a legislative framework for:

- Preventing persons from being killed or injured by electricity; and
- Preventing property from being destroyed or damaged by electricity.

Any future design for the proposed transmission line and substations must satisfy the requirements of the *Electrical Safety Act 2002*. The provisions of the *Electrical Safety Act 2002* apply to each corridor option in the same manner and therefore do not serve to differentiate corridor options in this report.

#### 9.2.6 NATURE CONSERVATION ACT 1992

Much of Queensland's native wildlife is protected by legislation to ensure its survival and to protect biodiversity. All native birds, reptiles, mammals and amphibians are protected in Queensland under the *Nature Conservation Act 1992* (NCA), along with a limited range of invertebrates, freshwater fish and the grey nurse shark. All plants that are indigenous to Australia are protected.

A permit is required to clear near threatened, vulnerable or endangered (commonly known as EVNT) protected plants and their supporting habitat. This permit is separate to the CID and IDAS processes. It cannot be definitively known if EVNT flora will be impacted until field studies are complete. If EVNT flora are identified in the area impacted by the alignment, a separate approval to CID and IDAS will be required from the Department of Environment and Heritage Protection.

Due to the preliminary nature of this report, the provisions of the NCA do not serve to differentiate corridor options. Powerlink has existing relationships with DEHP and significant experience managing impacts to flora and fauna regulated by the NCA.

#### 9.2.7 FORESTRY ACT 1959

The *Forestry Act 1959* regulates the use of forest products such as timber on all State land including State forests, leasehold land and unallocated State land. It is unlikely that any State forest areas will be impacted by the project, with all State forest areas located to the east of Mt Fox. The *Forestry Act 1959* is therefore not discussed further.

#### 9.2.8 ENVIRONMENTAL PROTECTION ACT 1994

The *Environmental Protection Act 1994* (EP Act) provides the key legislative framework for environmental management and protection in Queensland. The EP Act utilises a number of mechanisms to achieve its objectives.

Section 319 places a general environmental duty on an organisation undertaking an activity not to cause, or be likely to cause, environmental harm unless all reasonable and practicable measures to prevent or minimise the harm are taken.

By undertaking this study and subsequent environmental impact assessment studies, and through the development and implementation of environmental management plans, the duty of care will be demonstrated. A general environmental duty of care will be applied to the project irrespective of the preferred corridor and the EP Act therefore does not serve to differentiate corridor options.

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#### 9.2.9 ENVIRONMENTAL PROTECTION REGULATION 2008

The *Environmental Protection Regulation 2008* outlines Environmentally Relevant Activities (ERAs) that have potential to release contaminants to the environment or cause environmental harm. ERAs are assessable development under the SPA 2009 and require a permit (generally under the IDAS framework) before proceeding.

Transmission line development typically does not constitute an ERA. If an ERA is revealed during subsequent assessments a permit will be sought for the activity as part of the IDAS process. No ERA is expected to occur on any of the available corridor options and the *Environmental Protection Regulation 2008* therefore does not serve to differentiate corridor options.

#### 9.2.9.1ENVIRONMENTAL PROTECTION (WATER) POLICY 2009

This policy describes and seeks to protect the ecological and social values of waterways across the State. The policy will be regarded during subsequent assessments and appropriate design solutions and management measures adopted to ensure compliance with the policy. The policy does not serve to differentiate corridor options based on the current level of assessment.

#### 9.2.9.2 ENVIRONMENTAL PROTECTION (AIR) POLICY 2008

This policy describes and seeks to protect air quality values across the State. It outlines specific thresholds for contaminant release that are particularly relevant for construction phase activities associated with a transmission line.

The policy will be taken into consideration during subsequent assessments when appropriate design solutions and management measures can be identified and adopted to ensure compliance with the policy. The policy does not serve to differentiate corridor options based on the current level of assessment.

#### 9.2.9.3ENVIRONMENTAL PROTECTION (NOISE) POLICY 2008

This policy describes outcomes for the acoustic environment across the State. It outlines specific thresholds for noise that are particularly relevant for construction phase activities associated with a transmission line.

The policy will be regarded during subsequent assessments when appropriate design solutions and management measures can be identified and adopted to ensure compliance with the policy. The policy does not serve to differentiate corridor options based on the current level of assessment.

#### 9.2.10 ABORIGINAL CULTURAL HERITAGE ACT 2003

The Aboriginal Cultural Heritage Act 2003 provides for the recognition, protection and conservation of Aboriginal cultural heritage, including the establishment of a register and database of Aboriginal cultural heritage, and processes for the management of activities that may harm Aboriginal cultural heritage. The Queensland Department of Aboriginal and Torres Strait Islander Partnerships (DATSIP) is responsible for administering the Aboriginal Cultural Heritage Act 2003.

A search of the Cultural Heritage Register for Aboriginal cultural heritage identified numerous records of indigenous cultural heritage significance located within the study area. Specific Aboriginal parties relevant to the respective corridor are identified within Section 10.2 of this report.

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Powerlink is responsible for carrying out activities in accordance with the Duty of Care under the *Aboriginal Cultural Heritage Act 2003* by taking all reasonable and practicable measures to ensure its activities do not harm Aboriginal cultural heritage.

Powerlink has established processes and frameworks for, and significant experience in the management of cultural heritage in transmission line development. The processes of other parties or organisations are not discussed in this CSR.

#### 9.2.11 VEGETATION MANAGEMENT ACT 1999

The Vegetation Management Act 1999 (VM Act) is the principle legislation governing the management of native vegetation in Queensland (except for non-woody vegetation governed by the Nature Conservation Act 1992).

The VM Act is administered by Department of Environment and Heritage Protection (DEHP) and rests on the Regional Ecosystem (RE) mapping created by the Queensland Herbarium. Remnant vegetation is used as the trigger under the VM Act for determining when clearing of native vegetation is considered assessable development and requires a permit.

Exemptions from the need to obtain a permit to clear native vegetation are afforded where clearing is on land designated for community infrastructure. Clearing of native vegetation on land other than that designated for community infrastructure is likely to require a development permit.

Powerlink also is afforded some exemptions for clearing vegetation under provisions contained within the *Electricity Act 1994*. These exemptions may not apply to other development or constructing organisations.

#### 9.2.12 WATER ACT 2000

The purpose of the *Water Act 2000* is to provide for the sustainable management of water and other resources. Under section 266 of the *Water Act 2000*, a riverine protection permit is required to:

- Destroy vegetation in a watercourse, lake or spring;
- Excavate in a watercourse, lake or spring; or
- Place fill in a watercourse, lake or spring.

The State publishes a range of self-assessable codes that exempt certain activities from the need for a permit. It is not feasible at this level of assessment to determine whether a permit is, or is not required for any specific aspect of the project. The detailed requirement for a permit under the Water Act 2000 will be determined in more detailed assessment and permits gained as required. The Water Act 2000 does not serve to differentiate corridor options based on the current level of assessment.

#### 9.2.13 LAND ACT 1994

The *Land Act 1994* provides a framework for the allocation of State land as leasehold, freehold or other tenure and its subsequent management. Unallocated State land can also be purchased under the *Land Act 1994* where applications for such are approved by the DEHP.

Sections 101 and 102 of the *Electricity Act 1994* provide exemptions to the *Land Act 1994* allowing electricity entities such as Powerlink to undertake necessary actions on road reserves and publicly controlled places

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such as Unallocated State land. The *Land Act 1994* does not serve to differentiate corridor options based on the current level of assessment.

#### 9.2.14 ACQUISITION OF LAND ACT 1967

The Acquisition of *Land Act 1967* enables Powerlink to acquire freehold land or an easement over land for electricity works. Powerlink may acquire land:

- By agreement if an agreement can be reached on compensation to be paid, land or an easement can be acquired as soon as the necessary transfer documents have been executed, or
- Compulsorily resumptions are made subject to the provisions of the Acquisition of Land Act 1967.

Powerlink's first preference is to negotiate acquisitions wherever possible and will make all reasonable attempts to reach voluntary agreement for easements and other tenure requirements. Depending on the individual circumstances, and in order to ensure the timely delivery of projects, Powerlink may also begin compulsory resumption processes in parallel.

Powerlink has significant experience and established processes for the acquisition of land and easements by both negotiation and compulsory means and seeks to provide fair and reasonable support for landholders impacted by these processes, including access to independent expert advice relating to their compensation claim as early in the process as possible (i.e. on release of Draft EIS when there is greater certainty about the location of new infrastructure).

The land acquisition processes of other parties and organisations are not described by this report.

Land tenure across the project is principally leasehold and freehold. Due to the large nature of the land holdings traversed by the various corridor options there is insufficient variation in potential requirements under the *Acquisition of Land Act 1967* to differentiate corridor options.

#### 9.2.15 BIOSECURITY ACT 2014

The Biosecurity Act 2014 commenced on 1 July 2016. It ensures a consistent, modern, and risk-based approach to biosecurity in Queensland. The Act provides comprehensive biosecurity measures to safeguard the economy, agricultural and tourism industries, environment and way of life, from:

- Pests (e.g. Wild dogs and weeds)
- Diseases (e.g. Foot-and-mouth disease)
- Contaminants (e.g. lead on grazing land).

The Act replaced the many separate pieces of legislation that were previously used to manage biosecurity. Powerlink's internal policies and procedures are being aligned with the provision of the new Act and subsequent impact assessment phases will consider the requirements of the Act in greater detail. The provisions of the Act do not serve to differentiate study corridor options.

# 9.2.16 STOCK ROUTE NETWORK MANAGEMENT BILL 2016

The Stock Route Network Management Bill 2016 was introduced to Parliament in November 2016, proposing a single, contemporary Act to better support the long term management of Queensland's stock

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route network. Several declared stock routes may be traversed with the study area, depending on the final corridor selected. All of the declared stock routes within the study area are likely to be road reserves.

The Act requires all reasonable steps to be taken to minimise impacts to the resources of the stock route network as described in local government stock route network management plans. A stock route management plan has been identified for Charters Towers Regional Council. Powerlink engaged with both Charters Towers Regional Council and Etheridge Shire Council regarding potential impact to stock routes in the stakeholder engagement workshop on November 21, 2016. The ability for transmission lines to span stock routes was generally agreeable to all parties and will serve as the basis for more detailed impact assessments in subsequent project stages. Where stock routes are of a width that cannot be spanned, careful structure placement will be important to avoid impacting the utility of the stock route.

#### 9.2.17 TRANSPORT INFRASTRUCTURE ACT 1994

During construction, temporary road closures and temporary accesses (relating to State controlled roads) may be required to permit activities such as delivery of construction materials on long and/or wide haulage vehicles. State controlled roads within the study area include the Gregory Developmental Road and Kennedy Developmental Road.

The primary permit generally required under the Act is a road corridor permit, granted by the Queensland Department of Transport and Main Roads. The nature of all permits and licenses will be confirmed with the Department during subsequent detailed impact assessment and consultation processes. Because interactions with State roads are largely administrative in nature (with limited or no terrestrial impacts) the *Transport Infrastructure Act* does not aid in the differentiation of corridor options in this report.

#### 9.2.18 FISHERIES ACT 1994

The *Fisheries Act 1994* provides a framework, in conjunction with the SPA 2009, for the regulation of fisheries, declared fish habitat areas and marine plants. A review of the State declared fish habitat area plans did not identify any declared fish habitat areas within the study area. The inland location of the study area means that it is very unlikely that any aquaculture sites or marine plants will be impacted by the project.

Requirements of the *Fisheries Act 1994* may apply to the project during the construction phase, as works or disturbances may be required within watercourses traversed by the transmission line easement, such as waterway barriers and crossings. These matters will be identified and discussed with key regulatory stakeholders in subsequent environmental impact assessment studies. At the current level of assessment, the *Fisheries Act 1994* does not serve to differentiate corridor options.

#### 9.2.19 REGIONAL PLANNING INTERESTS ACT 2014

The *Regional Planning Interests Act 2014* identifies and protects areas of Queensland that are of regional interest. In doing this, the Act seeks to manage the impact and support co-existence of resource activities and other regulated activities in areas of regional interest. The Act includes Strategic Cropping Land.

The proposed transmission line does not qualify as either a resource activity or regulated activity under the Act. The provisions of the Act therefore do not apply.

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#### 9.2.20 REGIONAL PLAN

The Etheridge Shire area is part of the Gulf Regional Development Plan (GRDP). The GRDP focuses on management of growth and development in the region. It is noted that whilst in place since 2000, the GRDP is not a statutory regional plan.

Notwithstanding, the Genex Kidston Connection Project is likely to be considered consistent with the GRDP, supporting the growth of sustainable industries in the region. Comprehensive impact assessment and consultation during future assessment processes will ensure an alignment that best balances social, environmental and economic considerations is selected, upholding the balanced development outcomes sought by the GRDP.

Compliance with the specific strategies in the GRDP should be investigated during future assessment processes. It is not feasible however to undertake this level of assessment for the purposes of this document, where site specific impacts and the alignment are not known. Due to the broad nature of the Regional Plan it is unlikely that any provisions will differentiate corridor options.

The Hinchinbrook Shire Council and Charters Towers Regional Council areas are presently not subject to a regional plan, however feedback received from the Department of Infrastructure, Local Government and Planning has indicated that both LGAs are proposed for inclusion within a proposed North Queensland Regional Plan, which is being prepared and scheduled for consultation in mid-2017.

#### 9.2.21 STATE PLANNING POLICY

The State Planning Policy (SPP), which commenced on 1 July 2014, contains 16 State interests that are important to protect and enhance through Queensland's continued development. It is a key component of the State's land use planning system that enables responsible development, contributing to a liveable, sustainable and prosperous Queensland.

Energy supply is a State interest in the SPP, supporting economic development across Queensland. The Genex Kidston Connection Project will meet community expectations for the supply of energy infrastructure and the promotion of renewable energy projects. Compliance with the other State interests in the SPP, such as biodiversity, planning for hazards, protection of agriculture and others will be assessed during future assessment processes.

It is noted that the SPP is currently in a consultation phase as part of the reform process for the *Planning Act* 2016. Powerlink is actively engaged in the consultation process and , seeking to ensure any proposed amendments do not compromise its ability to develop the transmission network in a safe and efficient manner to support renewable energy projects. The provisions of the SPP do not serve to differentiate corridor options.

#### 9.2.22 INFRASTRUCTURE PLAN 2016

The State Infrastructure Plan (SIP), released on 13 March 2016, outlines a new strategic direction for the planning, investment and delivery of infrastructure in Queensland. The SIP identifies what the State Government ultimately seeks from its infrastructure and how this can best be achieved. Importantly, it is designed to provide confidence and certainty to business, industry and the community by confirming the government's investment program over the next four years.

The SIP recognises the importance of energy supply to the ongoing development of the State and places significant focus on the development of renewables and diversification of the energy sector. The project will

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support this focus. Compliance with the other infrastructure classes in the plan will be assessed during future assessment processes, though it is unlikely to differentiate corridor options.

#### 9.2.23 STATE DEVELOPMENT ASSESSMENT PROVISIONS

The State Development Assessment Provisions (SDAP) set out the matters of interest to the State for development assessment, where the Chief Executive administering the SPA 2009 is responsible for assessing or deciding development applications. They are statutory in nature but are not used by local government.

The SDAP is prescribed in the *Sustainable Planning Regulation 2009*, and contains the matters the Chief Executive administering SPA 2009 (the chief executive) may have regard to when assessing a development application as either an assessment manager or a referral agency.

The SDAP is divided into modules. Given the inland nature of the project area it is likely that many of the modules will not be relevant. Compliance with those modules that are relevant will depend greatly on the specific alignment and construction solutions identified for the project. These matters will be assessed during future assessment processes and do not differentiate corridor options at the current level of assessment.

It is noted that the SDAP is currently in a consultation phase as part of the reform process for the *Planning Act 2016*. Powerlink is actively engaged in the consultation process and seeking to ensure any proposed amendments do not compromise its ability to develop the transmission network in a safe and efficient manner to support renewable energy projects. The SDAP do not serve to differentiate corridor options.

### 9.3 LOCAL

#### 9.3.1 PLANNING SCHEMES

#### 9.3.1.1 HINCHINBROOK SHIRE COUNCIL

The Hinchinbrook Shire Planning Scheme took effect in 2005 and remains in force. The Planning Scheme does not list a transmission line as a defined use, with no defined use for infrastructure of any scale provided. The default level of assessment for any use that is not defined by the scheme is impact assessment. No overlays are noted other than bushfire mapping, which is ubiquitous across the shire. The likely overarching project approval framework would provide exemptions to the project from specific approvals under the respective local government planning schemes. However, compatibility with planning scheme intent would be assessed through any future Environmental Impact Statement and ongoing consultation with the Council.

The underlying zoning for the area impacted by the project is rural. The rural zone code contains a range of provisions for development in the rural zone, including probable solutions for intended uses in the zone. It is likely the project will need to demonstrate alternative solutions for many of these criteria because the code was not written with major linear infrastructure or substations in mind.

A full assessment against local codes is not feasible at the current level of assessment, with site specific impacts and alignments unknown. The broad mapping that underpins the Planning Scheme does not assist in differentiation of corridor options.

Powerlink engaged directly with Hinchinbrook Council during late 2016 and early 2017 with consultation at this early stage not identifying any major issues of concern. Powerlink will continue to work closely with Council to address planning scheme requirements and other matters should it be engaged to progress the

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Genex transmission line connection project further. Feedback from the consultation undertaken to date is provided in Section 13.

#### 9.3.1.2 CHARTERS TOWERS REGIONAL COUNCIL

The Dalrymple Shire Council Planning Scheme took effect in 2005 and remains in force in the project area. Dalrymple Shire was amalgamated with the City of Charters Towers in 2008 to form Charters Towers Regional Council.

The Planning Scheme defines the project as a Major Utility and the underlying zoning in the project area is rural. The level of assessment for a Major Utility in the rural zone is impact assessment. No overlays are noted other than nearby resource tenures. The likely overarching project approval framework would provide exemptions to the Genex Kidston Connection Project from specific approvals under the respective local government planning schemes. However, compatibility with planning scheme intent would be assessed through any future Environmental Impact Statement and ongoing feedback sought from Council.

The rural zone code contains a range of provisions for development in the rural zone, including acceptable solutions for intended uses in the zone. It is likely that the project will need to demonstrate alternative solutions for many of these criteria because the code was not written with major linear infrastructure or substations in mind.

A full assessment against local codes is not feasible at the current level of assessment, with site specific impacts and alignments unknown. The broad mapping that underpins the Planning Scheme does not assist in differentiation of corridor options.

Powerlink engaged directly with Charters Towers Regional Council during late 2016 and early 2017, with consultation at this early stage not identifying any major issues of concern. Powerlink will continue to work closely with Council to address planning scheme requirements and other matters should it be engaged to progress the Genex transmission line connection project further. Feedback from the consultation undertaken to date is provided in Section 13.

#### 9.3.1.3 ETHERIDGE SHIRE

The Shire of Etheridge Planning Scheme took effect in 2005 and remains in force. The Planning Scheme defines a transmission line as Community Infrastructure. The level of assessment for Community Infrastructure in the rural zone is code assessment. Bushfire and Good Quality Agricultural Land overlays are noted, which trigger code assessment or lower under the scheme provisions. The likely overarching project approval framework would provide exemptions to the project from specific approvals under the respective local government planning schemes. However, compatibility with planning scheme intent would be assessed through any future Environmental Impact Statement and ongoing feedback sought from Council.

The rural zone code and overlay codes contain a range of provisions for development, including probable solutions for intended uses. It is likely the project will need to demonstrate alternative solutions for many of these criteria because the codes were not written with major linear infrastructure or substations in mind.

A full assessment against local codes is not feasible at the current level of assessment, with site specific impacts and alignments unknown. The broad mapping that underpins the Planning Scheme does not assist in differentiation of corridor options.

Powerlink engaged directly with Etheridge Shire Council during late 2016 and early 2017, with consultation at this early stage not identifying any major issues of concern. Powerlink will continue to work closely with Council to address planning scheme requirements and other matters should it be engaged to progress the

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Genex transmission line connection project further. Feedback from the consultation undertaken to date is provided in Section 13.

#### 9.3.2 LOCAL LAWS

Queensland local governments have adopted a diverse range of local laws, administered through the *Local Government Act 2009,* to assist them to effectively govern their respective areas. While the project is potentially subject to a range of State and Commonwealth legislative and regulatory exemptions, it is not exempt from local laws and a range of permits may be required.

Local laws likely to apply to the project include:

- Etheridge Shire Council Local Law No. 4 (Local Government Controlled Areas, Facilities and Roads) 2011
- Etheridge Shire Council Local Law No. 3 (Community and Environmental Management) 2011
- Charters Towers Regional Council Local Law No. 3 (Community and Environmental Management) 2011
- Charters Towers Regional Council Local Law No. 4 (Local Government Controlled Areas, Facilities and Roads) 2011
- Charters Towers Regional Council Local Law No. 7 (Aerodromes) 2011
- Charters Towers Regional Council Local Law No. 24 (Grids)
- Charters Towers Regional Council Local Law No. 5 (Parking) 2011
   Hinchinbrook Shire Council Local Law No. 3 (Community and Environmental Management) 2012
- Hinchinbrook Shire Council Local Law No. 4 (Local Government Controlled Areas, Facilities and Roads) 2012
- Hinchinbrook Shire Council Local Law No. 5 (Parking) 2012
- Hinchinbrook Shire Council Local Law No. 7 (Aerodromes) 2012.

Local laws will be included in subsequent detailed impact assessments, where site specific impacts are known and the requirement for local law permits can be more thoroughly assessed.

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## PROPOSED GENEX KIDSTON CONNECTION PROJECT

## **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"



### 10 CONSTRAINTS AND OPPORTUNITIES

#### 10.1 EXISTING TENURE & LAND USE

#### 10.1.1 TENURE

Land tenure in the project area is principally leasehold. This land is held by the State of Queensland and leased for specific purposes (grazing, agriculture, telecommunications etc.) for a specified period. These leases are generally for a long period, registered on title and, in many cases, offer the lessee exclusive rights over the land. There are also scattered freehold lots throughout the project area.

Land use is rural in nature, predominately grazing, though specific management practices are not known at the current level of assessment. More details will be discovered through future field reconnaissance and engagement with authorities and landholders in the region. The Planning Schemes note the land as rural in nature and no intent to depart from rural uses is evident.

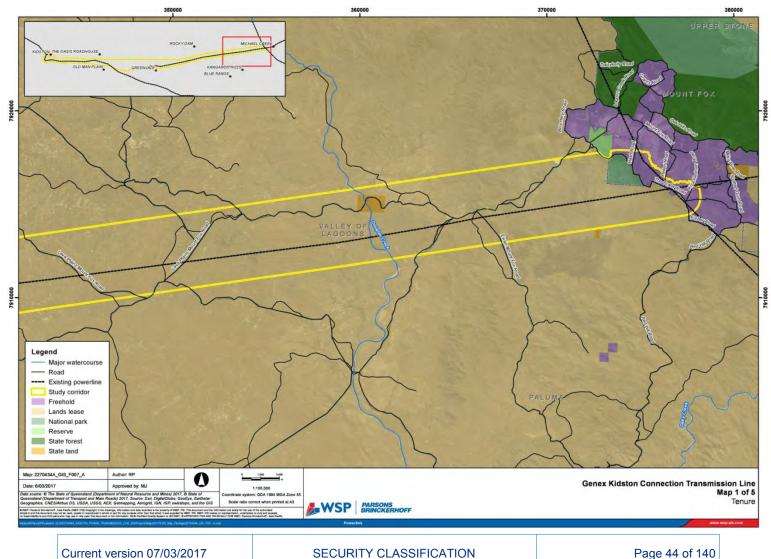
The project is not likely to impact heavily on existing tenure and land uses, which occur over broad areas and appear to be generally of low intensity. Potential conflicts with existing land use and tenure will be discovered through engagement with relevant authorities, with feedback utilised to refine corridor options. It is not possible to differentiate corridor options based on land management practices at the current level of assessment.

Maps depicting tenure and pastoral holdings are provided in Figure 7 and Figure 8.

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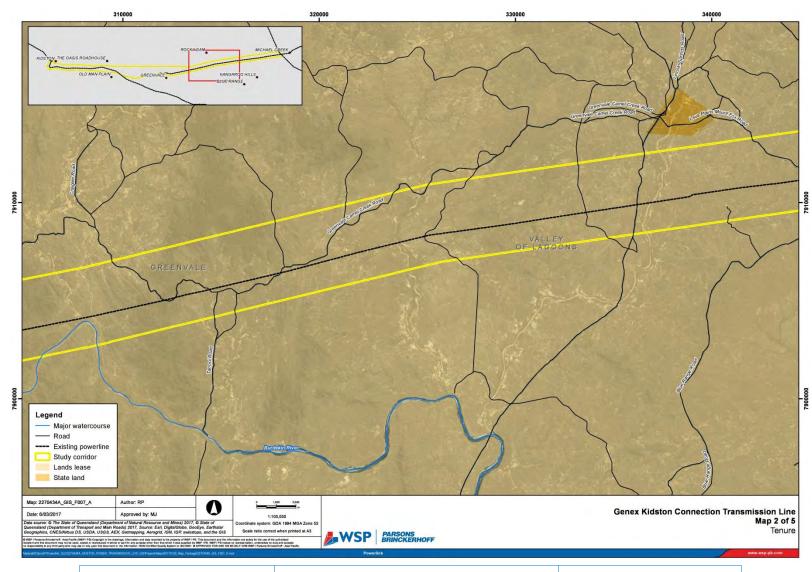


Figure 7 – Tenure



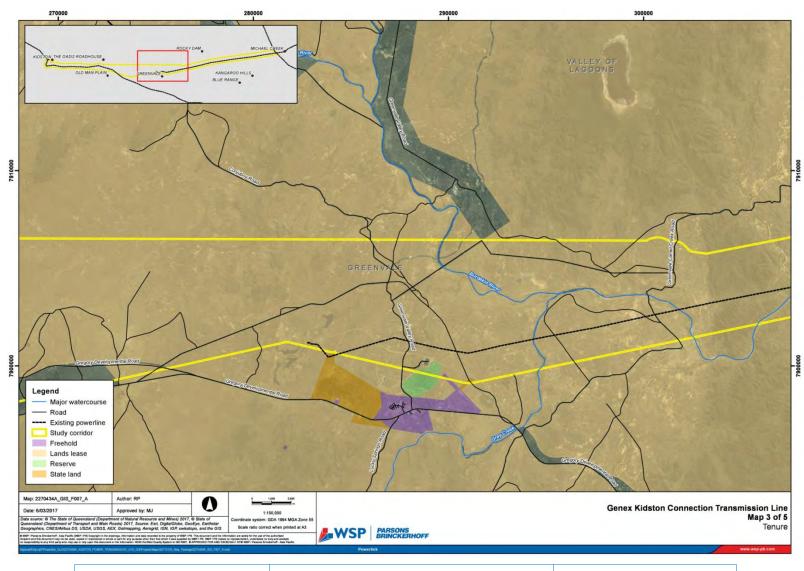
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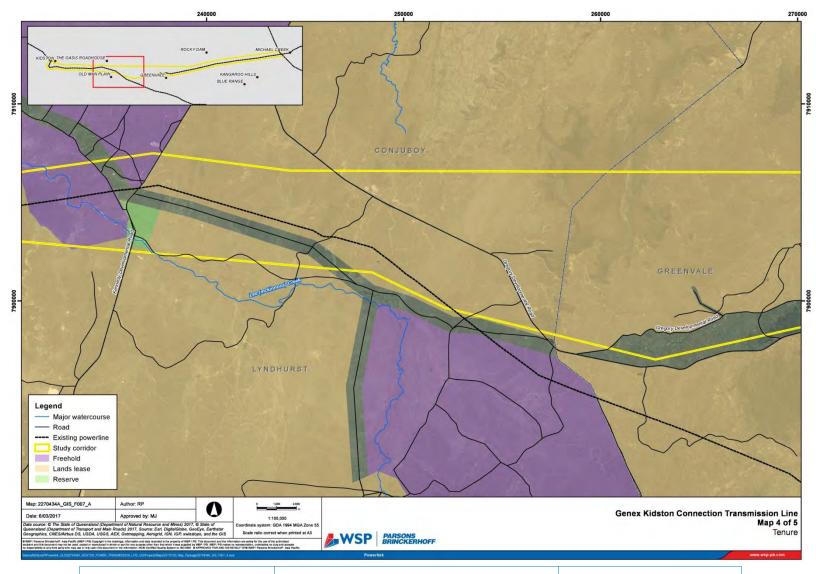
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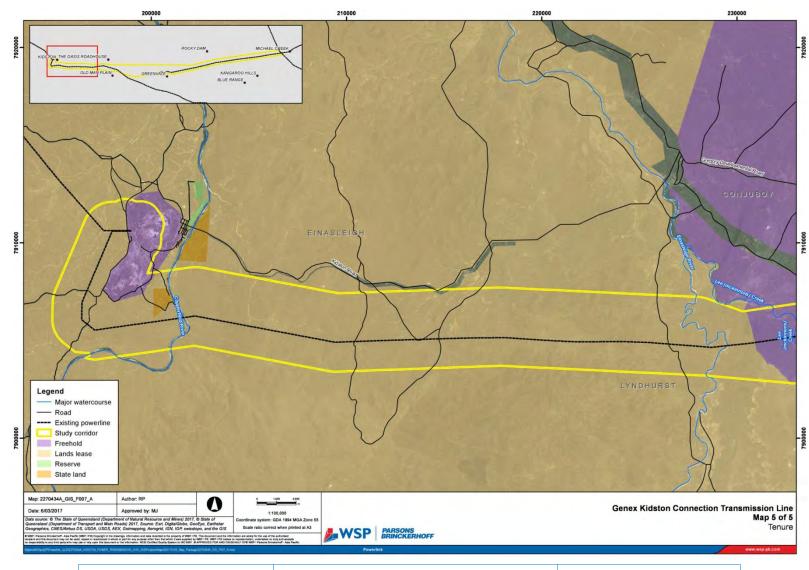
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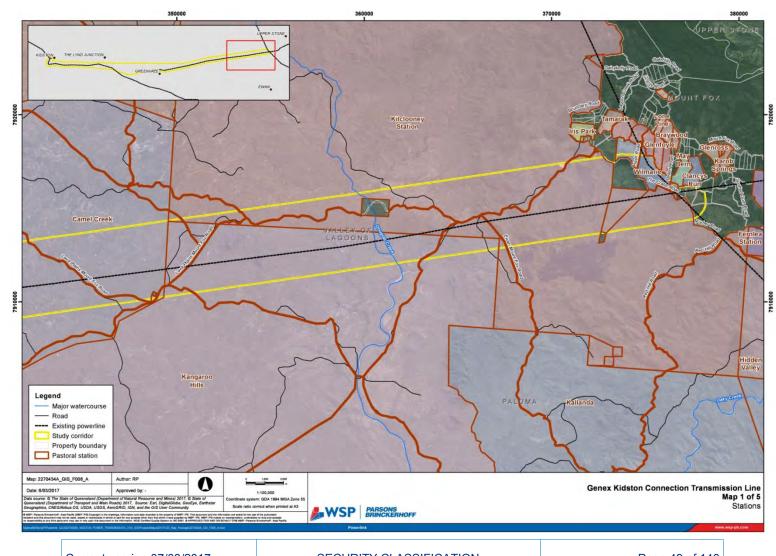




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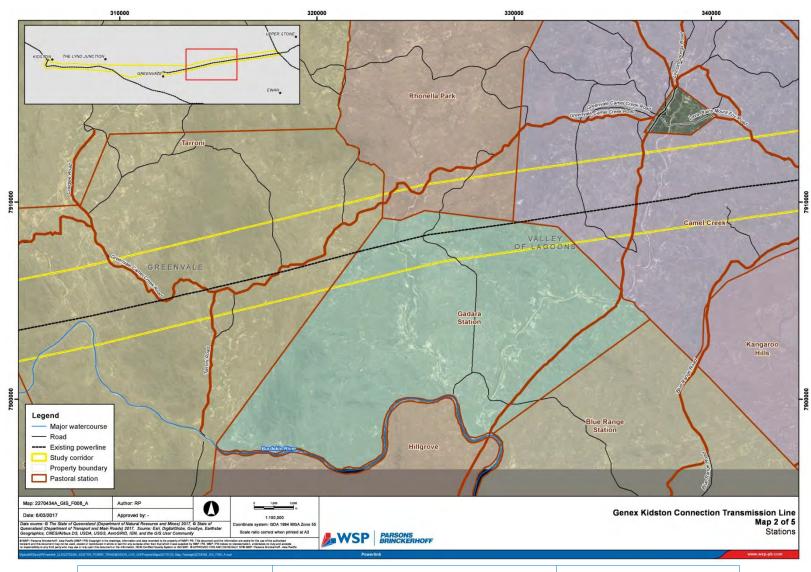


Figure 8 – Pastoral Stations



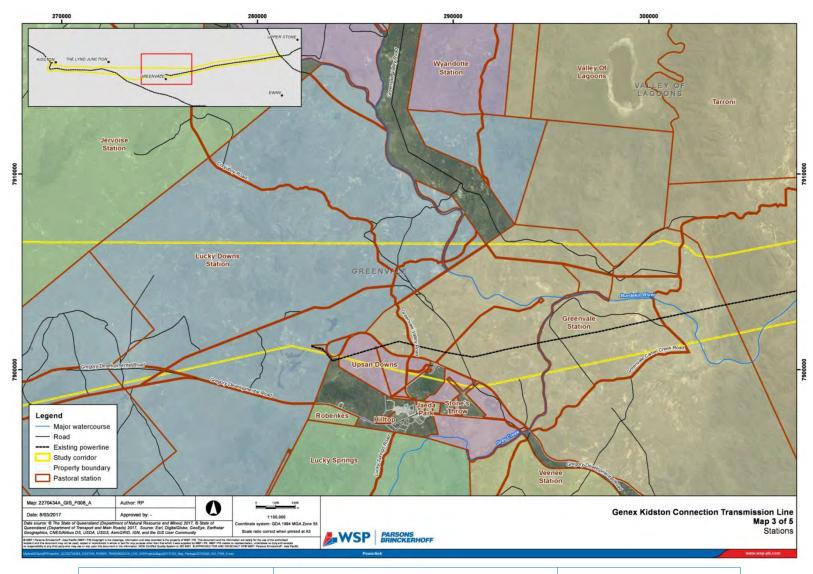
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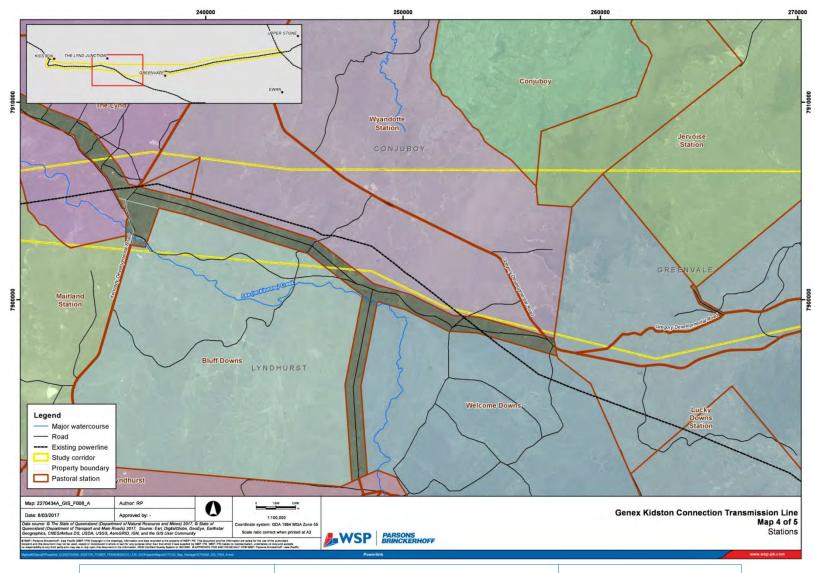
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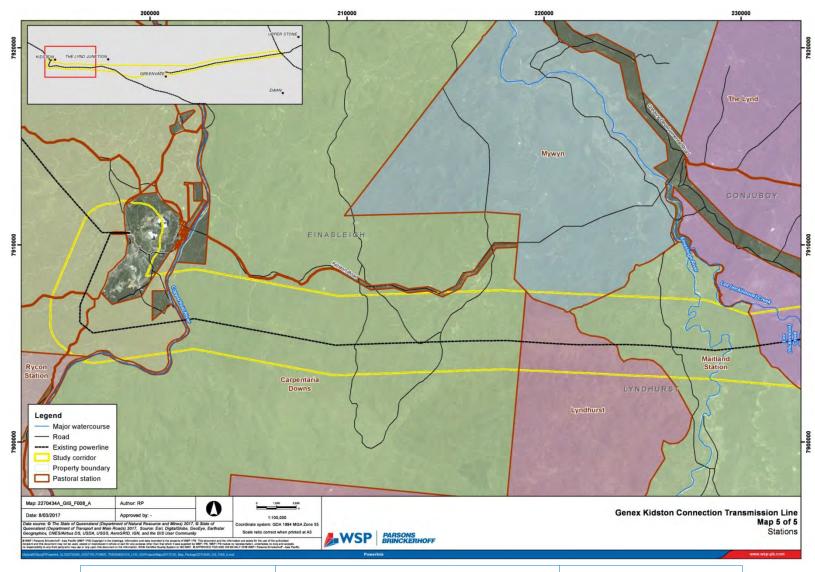
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#### 10.1.2 RESOURCE INTERESTS

Resource interests in Queensland are generally governed by the *Petroleum and Gas (Production and Safety) Act 2004* and *Mineral Resources Act 1989*. These acts afford resource explorers and producers certain rights above and below land. The resource interest holder is not required to also be the owner of the land over which the resource right exists. The rights are granted under various mechanisms dependent on the type of activity and maturity of any resource operation development.

Where infrastructure is proposed to cross or traverse a resource interest, consent from the respective authority holders pursuant to the respective governing legislation as discussed above will be required for the construction of transmission line, as well as any heavy vehicle and plant movements which have the potential to impact any existing infrastructure. Additionally, resource interest holders may be entitled to compensation as part of any land/easement acquisition process.

No interests under the *Petroleum and Gas (Production and Safety) Act 2004* have been identified within the study area. Several resource interests as governed by the *Mineral Resources Act 1949* have been identified. Consents from resource interest holders are not linked to the EIS or IDAS approvals processes and can be sought once a design and construction activities are more certain. Powerlink has significant experience in negotiating these consents through its previous work in the Surat and Bowen Basin regions.

It is noted that the primary resource interests identified are for exploration purposes. This is less likely to have a bearing on future alignment selection, unless exploration activities are well advanced and a possible footprint of mining development is known. Where a corridor interacts with Mining Leases, consideration will be given to specific activities of those operations.

Table 1 identifies the resource interests in the study area. Sections 10.1.2.1 to 10.1.2.3 provide some specific details for the mining leases identified. The Kidston mining lease is not discussed because it occurs on the Genex site.

At the current level of investigation there appears to be minimal impact to resource interests and ample opportunity to mitigate impacts through route selection given the vastness of the project area. Resource interests therefore do not differentiate study corridor options at the current level of investigation.

#### Table 4 - Study Area – Resource Interests

Permit Number	Status	Expiry	Principle Holder	
MINERAL EXPLORATION PERMITS				
EPM 12513	Granted	04/01/2019	SNOW PEAK MINING PTY LTD	
EPM 14107	Granted	25/08/2019	SNOW PEAK MINING PTY LTD	
EPM 14576	Granted	21/06/2017	CURTAIN	
EPM 17077	Granted	30/05/2019	ECHO RESOURCES LIMITED	
EPM 18877	Granted	13/02/2017	COPPER STRIKE LTD	
EPM 18987	Granted	24/09/2018	SUPERIOR RESOURCES LIMITED	
EPM 25259	Granted	07/04/2019	SNOW PEAK MINING PTY LTD	
EPM 25498	Granted	10/11/2019	SNOW PEAK MINING PTY LTD	
EPM 25659	Granted	20/04/2020	SUPERIOR RESOURCES LIMITED	
EPM 25691	Granted	07/04/2020	SUPERIOR RESOURCES LIMITED	

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Permit Number	Status	Expiry	Principle Holder
EPM 25834	Granted	05/01/2021	NORNICO PTY LTD
EPM 25865	Granted	14/12/2020	GREENVALE OPERATIONS PTY LTD
EPM 25934	Granted	08/06/2021	CORONATION RESOURCES LIMITED
EPM 26163	Application		LYZA MINING PTY LTD
EPM 26165	Application		SUPERIOR RESOURCES LIMITED
EPM 26326	Application		CAESAR GOLD MINES PTY LTD
EPM 26353	Application		LITHIUM AUSTRALIA NL
EPM 26356	Application		OAKDALE RESOURCES LIMITED
EPM 26360	Application		STURT EXPLORATION AND MINING PTY LTD
EPM 9323	Granted	12/04/2016	SNOW PEAK MINING PTY LTD
MINING LEASES			
ML 10368	Application		GREENVALE OPERATIONS PTY. LTD.
ML 1363	Granted		KELSO
ML 3347	Granted		KIDSTON GOLD MINES LIMITED
ML 6750	Granted		SUPERIOR RESOURCES LIMITED

#### 10.1.2.1 MLA10368 SCONI PHASE 1 PROJECT

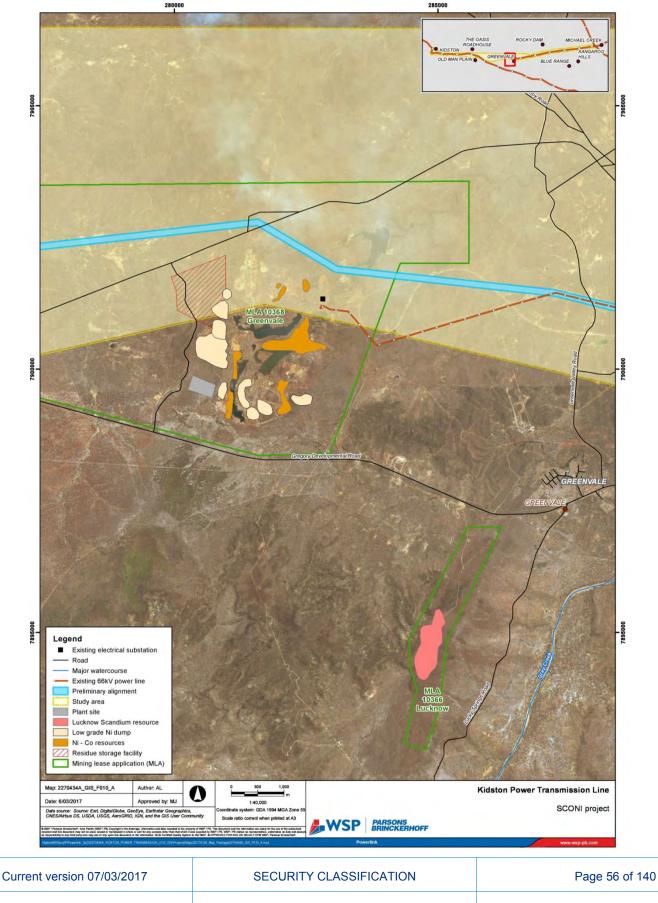
Approximately 6km west of Greenvale Township is the historical Greenvale Nickel mine, now operated by Metallica Minerals Pty Ltd as the SCONI project, with scandium, cobalt and nickel resources. An ASX release from March 2013 identified the following indicative layout for the operation (Figure 9).

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Figure 9 – SCONI Phase 1 Project



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The existing 66kV Ergon transmission line can be seen in Figure 9, representing the general path of study corridor option C. Metallica Minerals commenced the EIS process, lodging an initial advice statement and drafting a Terms of Reference consistent with the *Environmental Protection Act 1994 in 2012 and 2013 respectively*. The process has not been continued and is now registered as a withdrawn EIS process on the Department of Environment and Heritage Protection website, with the proponent voluntarily concluding the EIS process in 2015<sup>3</sup>.

There appears to be sufficient room between the resource areas to site a transmission line. Powerlink provided written project information to Metallica Minerals about its study corridor investigations in early 2017 and invited Metallica to participate in further discussions if desired.

#### 10.1.2.2 ML6750 'ONE MILE'

ML6750 is known as One Mile and based on exploratory drilling that contains low-grade copper, gold, zinc and silver values. Powerlink provided written project information to the One Mile about its study corridor investigations in early 2017 and invited them to participate in further discussions if desired.

#### 10.1.2.3 ML1363 'LUCKY DIP'

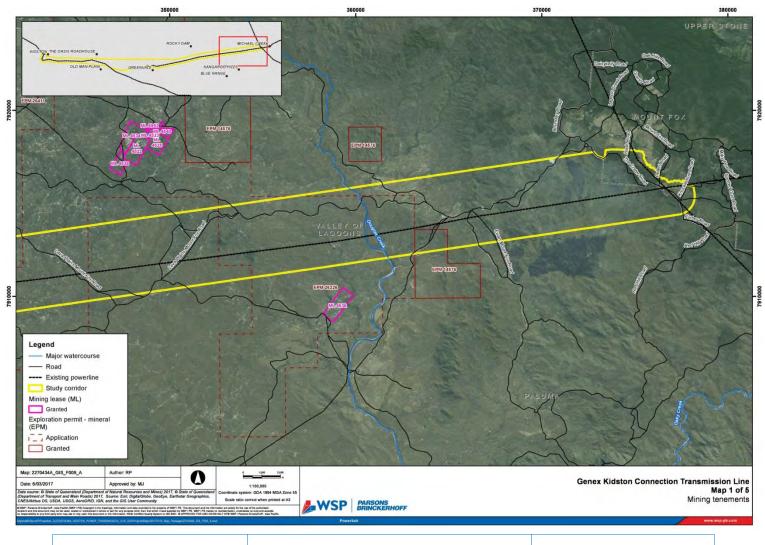
ML1363 is an older permit, lodged in 1966. Few details are recorded against the permit and it is very small at only 2 hectares. No contact details could be identified for the permit. Given its diminutive size and the vastness of the surrounding project area, the permit is readily avoided if necessary.

<sup>3</sup> https://www.ehp.qld.gov.au/management/impact-assessment/eis-processes/concluded.html

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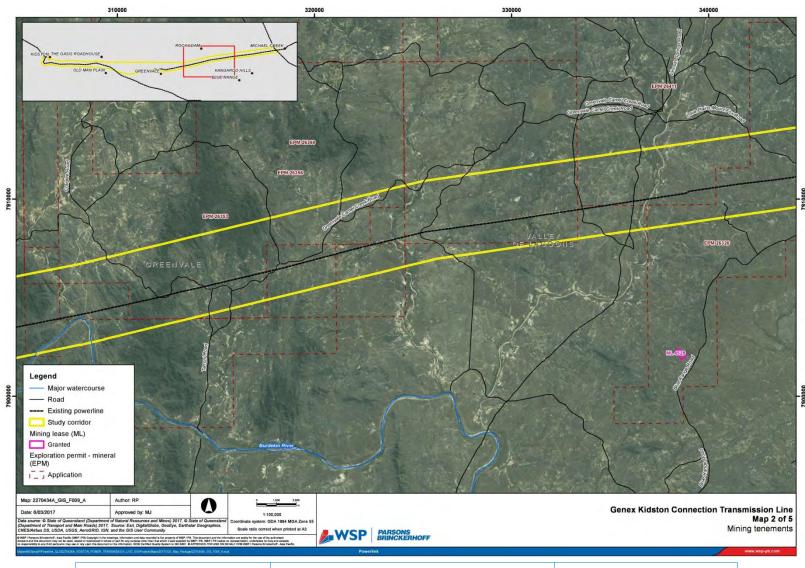


Figure 10 – Mining Interests



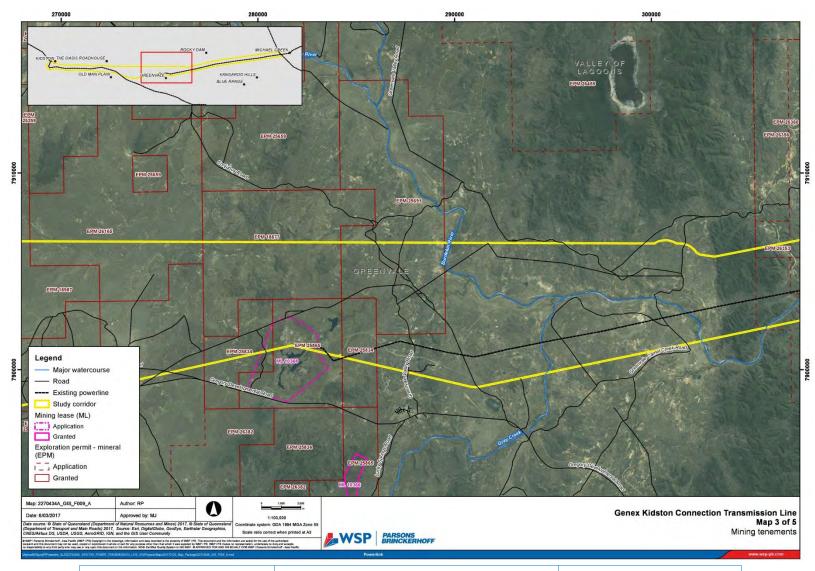
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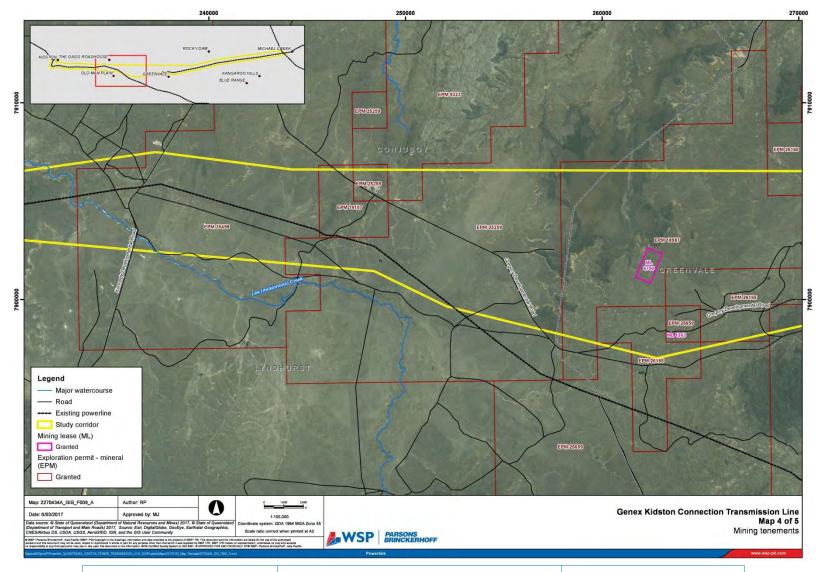
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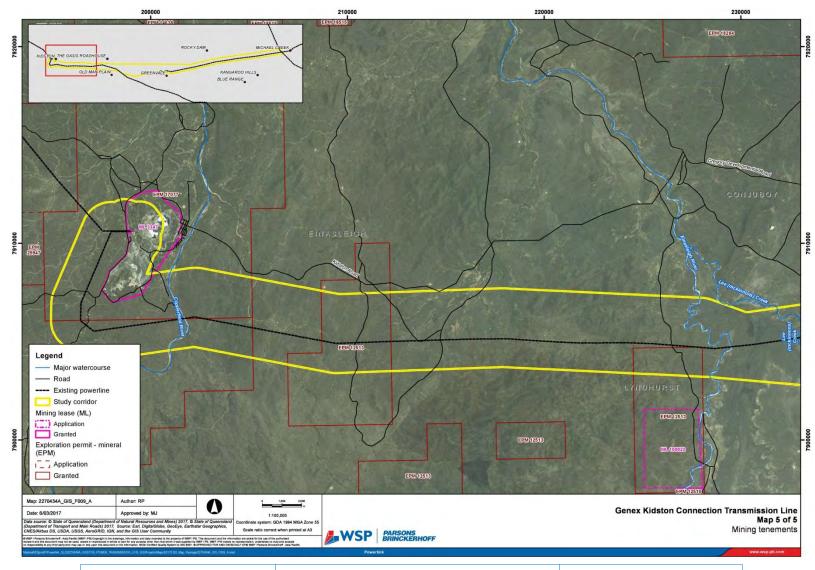
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#### 10.1.3 DEVELOPMENT APPLICATIONS

A list of active development applications is published online by the Hinchinbrook Shire and Charters Towers Regional councils. The address of each of the development applications was searched to determine whether any are likely to occur in the project area. No applications were identified in or near to the study area.

Existing development applications are not published on the Etheridge Shire Council website. Etheridge Shire Council attended the stakeholder workshop in November 2016 and no conflicting developments were noted at that time or in subsequent engagement with Council.

#### 10.1.4 ENCUMBERANCES

Few easements exist in the vicinity of the project, limited to existing Powerlink and Ergon power lines. These easements do not pose a constraint to the project. A number of strata rights exist across the project area. The exact nature of the land uses associated with these rights is unknown, though they appear to occur over state lands and stock routes.

Engagement during the impact assessment stage will be required to determine the specific nature of the many strata rights observed. These encumbrances do not serve to differentiate study corridor options at the current level of assessment and are considered low risk to the overall development of the proposed Genex Kidston Connection transmission line.

#### 10.1.5 EXISTING INFRASTRUCTURE

The project area contains little existing major infrastructure. Major roads include the Gregory and Kennedy Developmental Roads, which are intersected in the vicinity of Conjuboy. The proposed transmission line will be able to safely span over these roads.

Similarly the roads are unlikely to offer significant opportunity for co-location because they emanate from locations that offer insufficient access to the electrical transmission network and traverse through the populated areas in the region, such as Greenvale.

Existing Ergon transmission lines are noted, running parallel to the southernmost corridor option from Mt Fox to Greenvale and from an area around Conjuboy to Kidston. These lines operate at transmission voltages (66kV and 132kV). Existing cleared corridors for the Ergon lines may offer co-location opportunities for a new transmission corridor.

Co-location is beneficial in grouping together large, linear disturbances to the natural landscape. Fragmentation of vegetation is kept to a single area, minimising breaks in ecological connectivity in the area. Further benefits are derived in the potential for common access tracks, minimising both physical and social disturbance in the region.

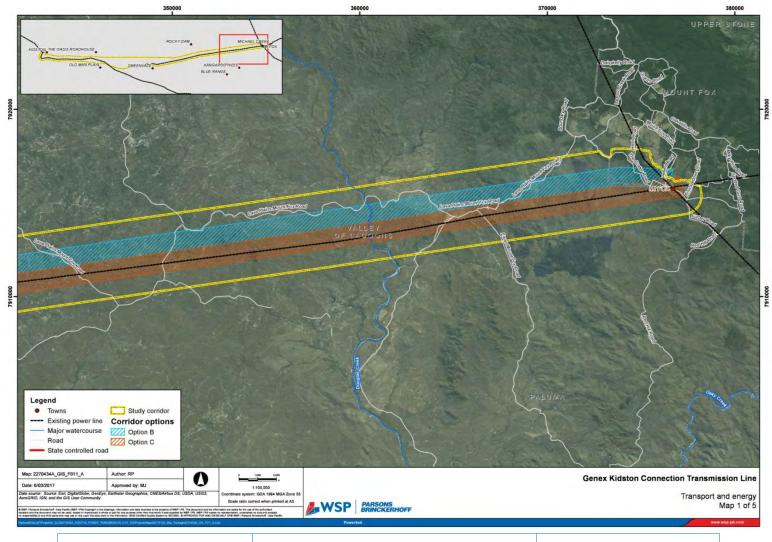
A number of small private airstrips, as well as larger facilities at Greenvale and Kidston were noted during travel through the study area. All corridor options are likely to be proximal to airstrips and potential impacts to airstrips and aerial activities such as mustering will be an important consideration in subsequent impact assessments. Airstrips do not serve to differentiate corridor options at the current level of investigation.

Figure 11 identifies existing infrastructure throughout the project area.

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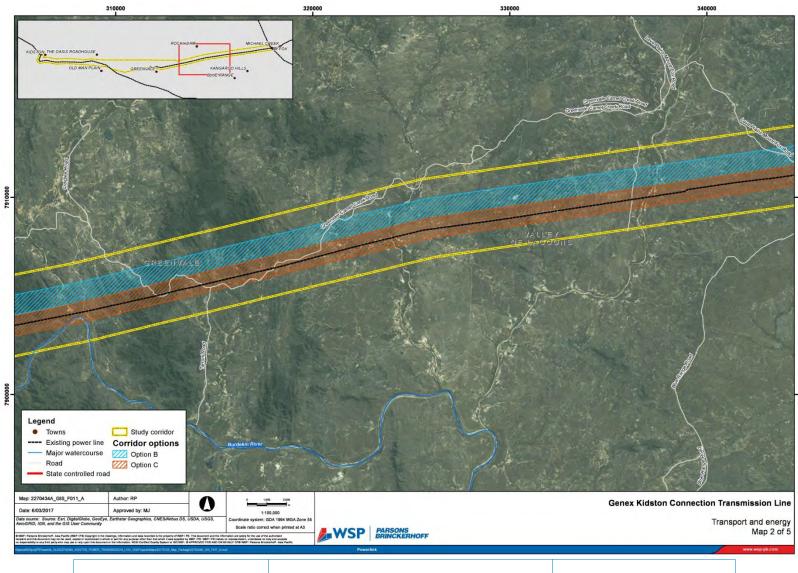


Figure 11 – Transport and Energy Infrastructure



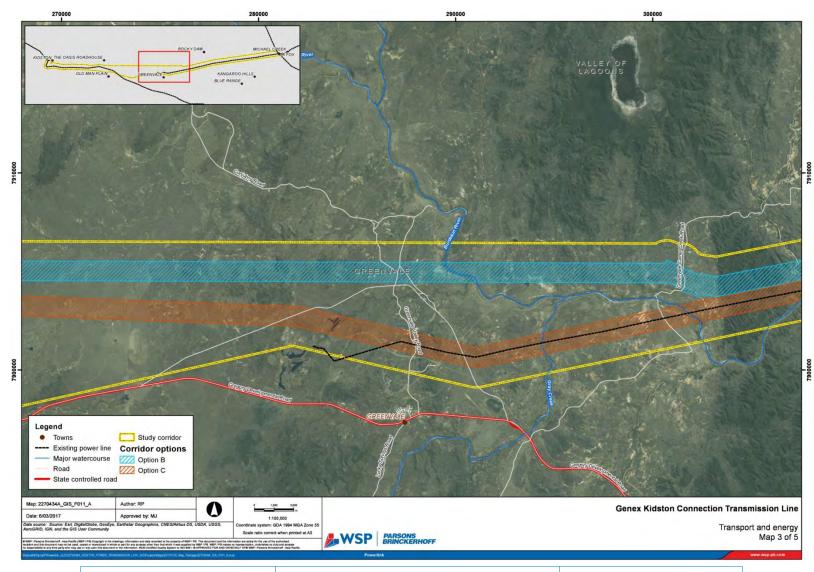
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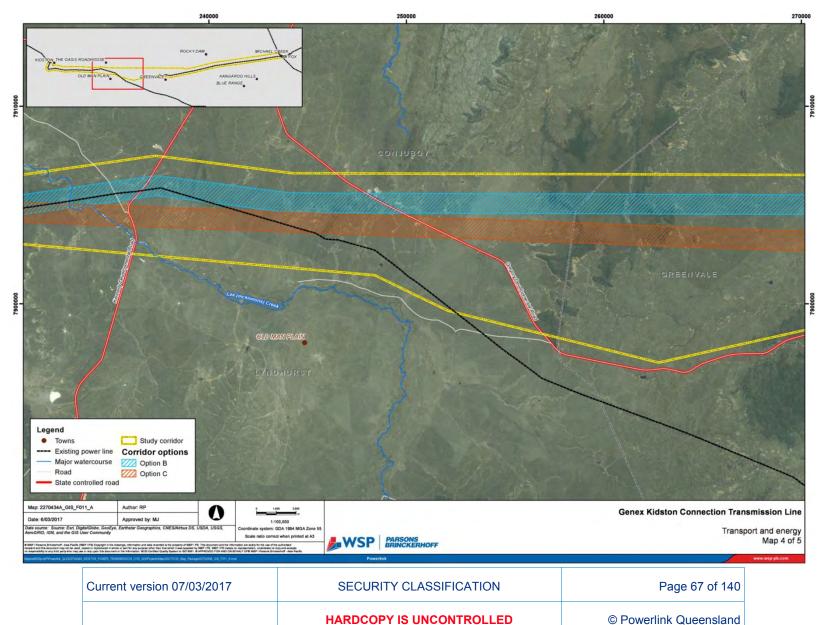
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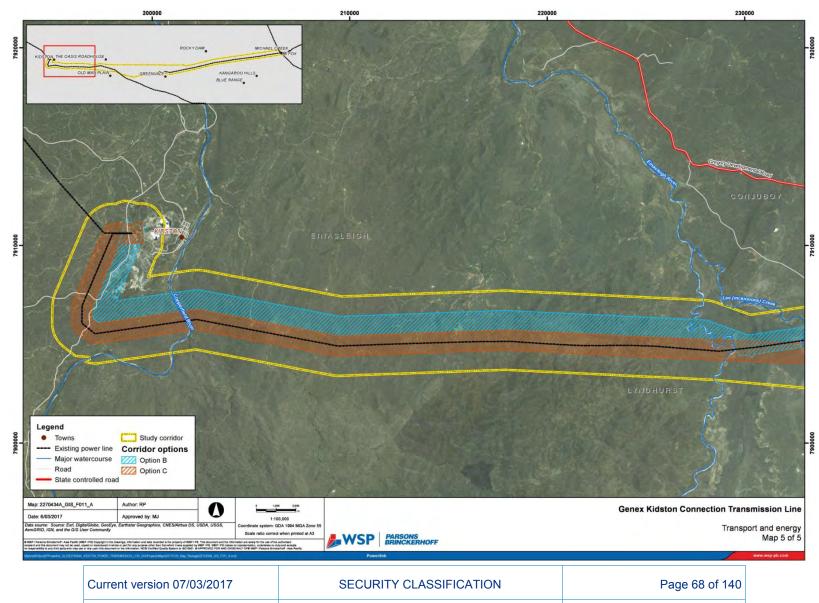


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#### 10.1.6 COMMUNITY BENEFITS

The project potentially provides a range of benefits, including:

- Supporting a major renewable energy project
- Supporting both the Commonwealth Renewable Energy Target and State planning intent
- Supporting the diversification of Queensland's electricity generation mix
- Creation of jobs during the construction period and for ongoing operation.

The benefits of renewable energy projects are widespread and supported by legislation and policy. Employment generated during the construction period is likely to be of regional and local benefit. Because the proposed transmission line is approximately 200km long, it is likely the opportunity for project-related employment will occur across a broad geographic area and flow-on effects to small business will be equally wide spread.

Consultation will occur with relevant authorities regarding strategies to maximise community benefits should the project proceed to the next phase in the assessment process. All corridors provide similar community benefits.

## 10.2 CULTURAL HERITAGE AND NATIVE TITLE

#### 10.2.1 ABORIGINAL CULTURAL HERITAGE

A search of the Department of Aboriginal and Torres Strait Islander Partnerships (DATSIP) database identifies three Aboriginal Parties whose Native Title determination areas are intersected by the study corridor.

- Gugu Badhum People #3
- Ewamian People #2
- Ewamian People #3.

Activities to be carried out during the clearing and construction phase for transmission lines are typically assessed as Category 5 of the Duty of Care guideline under the *Aboriginal Cultural Heritage Act 2003* (ACHA) i.e. ground disturbing activities within a greenfield site.

Powerlink's strategy to meet its Duty of Care (s23 ACHA) is through the negotiation of a formal agreement or Cultural Heritage Management Plan (CHMP) with each of the relevant Aboriginal Parties and conducting its land clearing and construction activities in accordance with that agreements or CHMP. This typically includes a detailed cultural heritage survey.

Powerlink has established processes for, and significant experience in the management of cultural heritage in transmission line development and working closely with Aboriginal parties. The processes of other development or constructing organisations are not discussed in this CSR.

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#### 10.2.2 EUROPEAN CULTURAL HERITAGE

A search of the Queensland Heritage Register (DEHP) found one listing within proximity to the project. The Kidston State Battery and Township is classified as a State heritage place, however is not located within any of the three corridors. Historical heritage is therefore not an influencing factor in differentiating between possible corridors at the present level of assessment.

#### 10.2.3 NATIVE TITLE

The study area discussed in this report intersects the following Native Title determinations:

- Gugu Badhum People #3
- Ewamian People #2
- Ewamian People #3.

The acquisition of an easement for the transmission line would be a 'future act' under the Native Title Act 1993 (C'th) and therefore requires that Native Title interests are addressed. To the extent that an easement acquisition and construction of infrastructure affects Native Title, Powerlink typically utilises section 24KA of the Native Title Act 1993 (C'th), which applies to facilities for services to the public, to validate the act and 'supress' Native Title over the easement for the 'lifetime' of the infrastructure.

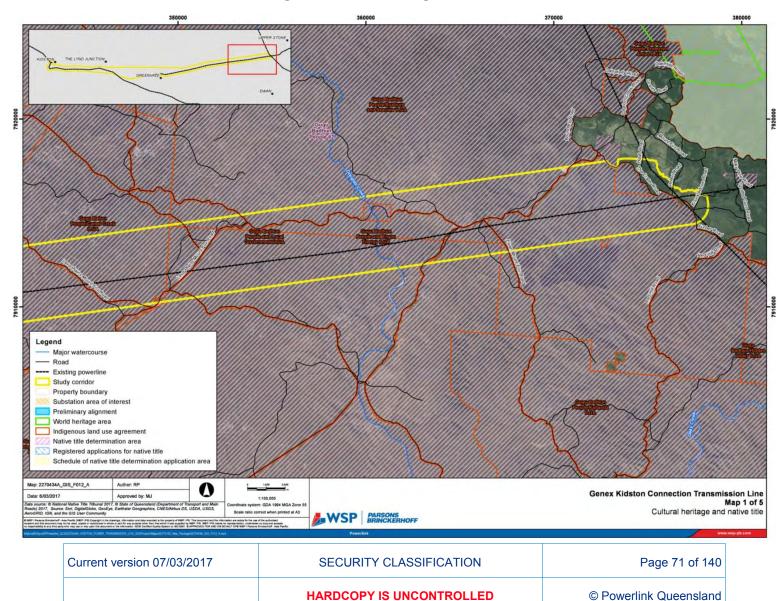
On the basis that any proposed alignment within the project area would intersect the above determination areas, Native title compliance would not significantly differentiate between corridor options at present. Powerlink has established processes for, and significant experience in the management of Native Title in transmission line development. The processes of other development or constructing organisations are not discussed in this CSR.

Figure 12 depicts the Native Title determinations that intersect the project area.

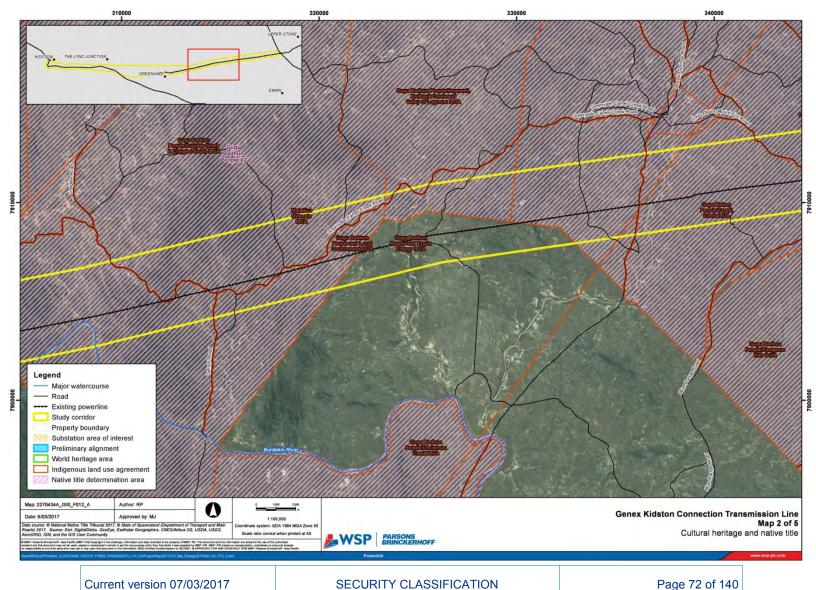
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Figure 12 – Cultural Heritage & Native Title

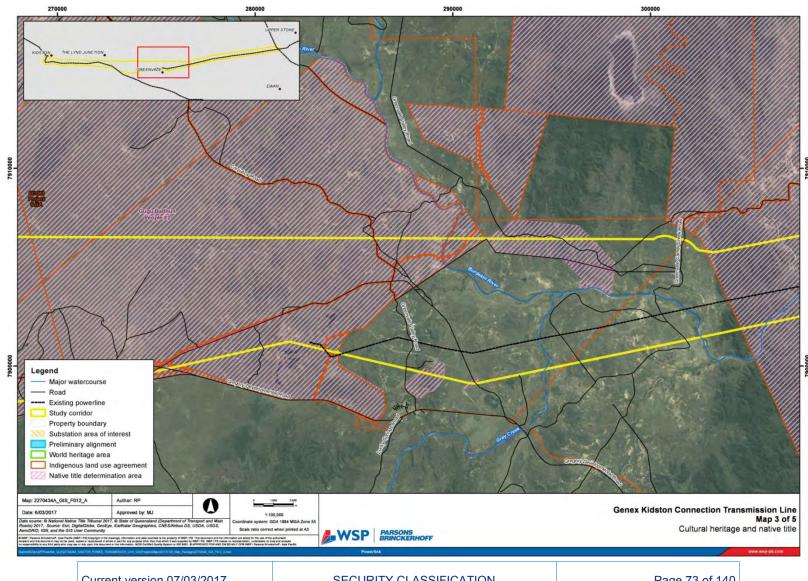






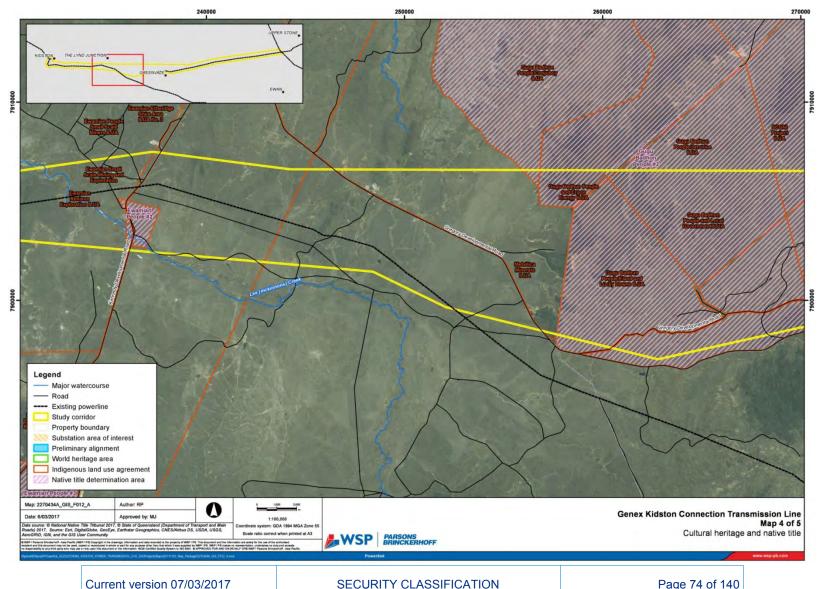
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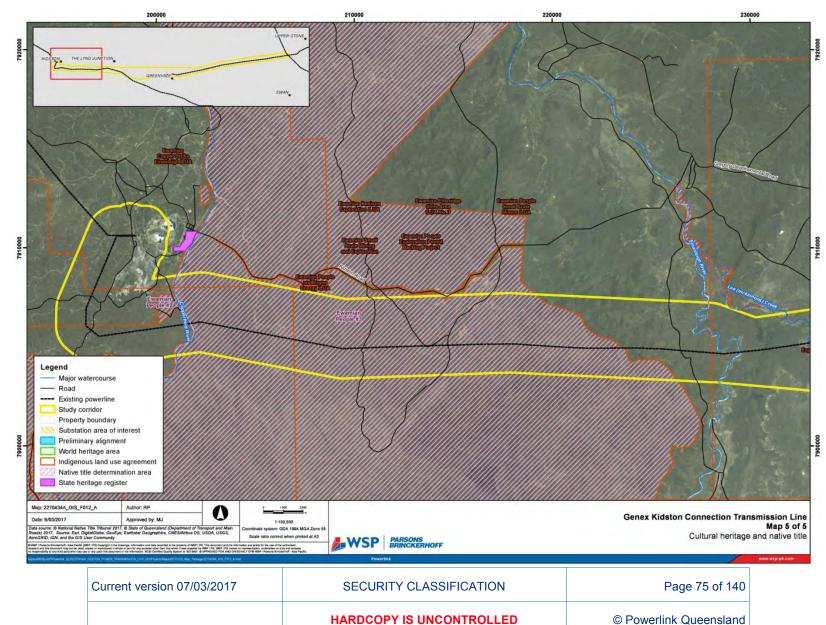
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## 10.3 VISUAL AMENITY

The transmission line has the potential to be visible from some residences, local government controlled roads, State controlled roads and Girringun National Park. The infrastructure has the potential to be visible from the following locations:

- Local residences
- Mt Fox Crater
- Girringun National Park
- The Crater Road
- Lava Plains Mt Fox Road
- Kangaroo Hills Road
- Ewan Mt Fox Road
- Valley of Lagoons Road
- Craiglee Road
- Conjuboy Road
- Runs along Gregory Development Road
- Kennedy Development Road
- Unnamed Roads.

Visual amenity can be described as comprising two elements: a near field view, comprising the immediate and permanent amenity as visible from a fixed point (e.g. residences); and a far field view, comprising a more transient visual impact from a further distance (e.g. seen from a vehicle at a road crossing point). The severity of the impact depends on the proximity to the subject (in this instance the substation structure) and the amount of time that the structures are visible.

Visual amenity impacts will be assessed in detail in subsequent approval processes. Further engagement with potentially affected stakeholders and landholders would also occur, informed by the engagement undertaken as part of this CSR.

Visual exposure to the transmission line may be either avoidable or of minimal impact given the vast and expansive nature of the surrounding environment and the ability to carefully site transmission structures. Where exposure is not avoidable and a measurable impact is likely to occur, alternative mitigations will be investigated.

Powerlink has significant experience in the assessment, mitigation and management of visual amenity associated with transmission lines.

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#### 10.4 ELECTRIC AND MAGNETIC FIELDS (EMF)

EMF readings at the edge of an easement for a 275kV transmission line are generally similar to those encountered by people in their daily activities at home or at work (e.g. a similar reading to what experienced using a typical electrical appliance or power tool). At about 100 metres from the line, the level of EMF is so small that it typically cannot be measured.

Powerlink has undertaken an EMF assessment for the proposed Genex Kidston Connection transmission line, taking into consideration the expected maximum capacity of the line based on both the hydro and solar generation facility in full operation. This technical assessment has confirmed that both the electric field and magnetic field strength for the proposed line will be below international guidelines.

## 10.5 AIR AND NOISE

The nearest town to any of the draft corridor options is Greenvale (population 150), some 2km from the nearest corridor. The nearest population centre to the project area is Ingham (approximately 40km north-east of Mt Fox, population 4,706). With regard to air and noise, minimal to negligible impacts on population centres would be expected. Any impact to individual residences will require detailed consideration in any subsequent impact assessment phases, with only a very small number of properties proximal to all corridor options.

Additionally, during construction and operation of any transmission line, the environmental duty of care and compliance with the respective Environmental Protection Policies for Air and Noise management must be demonstrated. Measures contained within these policies would be applicable to any project and do not serve to differentiate corridors.

## 10.6 TRAFFIC

Given the low-density nature of the surrounding environment, it is unlikely that major disruptions to the road network will be required. The requirement for road interruptions (or disturbance to other transport modes) and road improvements will be investigated during subsequent impact assessments and in consultation with relevant authorities.

Traffic impacts do not differentiate corridor options at the current level of assessment.

## 10.7 TOPOGRAPHY, GEOLOGY & SOILS

The topography of the project area ranges from flat low-lying land to steep crossings of multiple ranges (including the Great Dividing Range). Elevation ranges from 400-800m AHD. All corridor options will be required to cross the Great Dividing Range (100km west of Mt Fox). Range crossings have been identified as a key constraint and guide the general direction of the corridor options to minimise associated constructability issues.

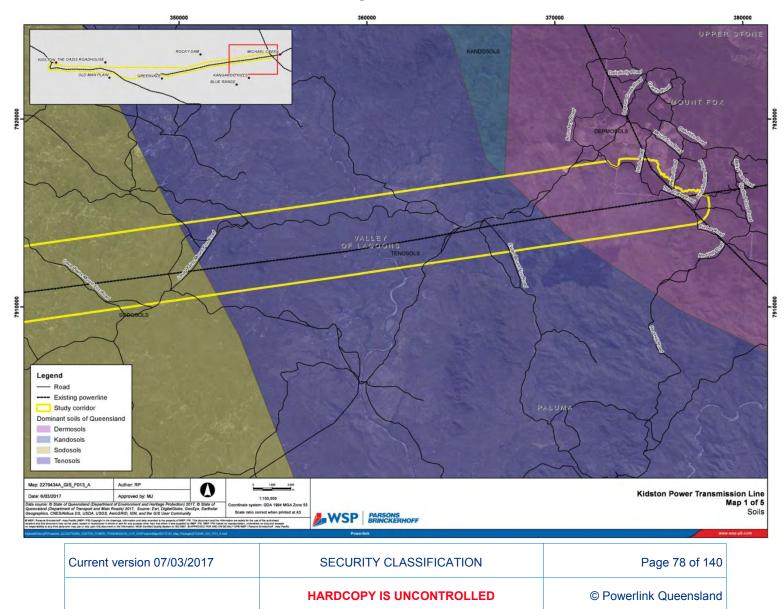
All draft corridor options traverse mainly chromosol and sodosol soil types (Figure 13). There is a very low probability of encountering acid sulphate soils (ASS) along any of the corridors as they mainly occur on coastal land that has elevations below 5m AHD. ASS are also required to be considered on land that is below 20m AHD. Both conditions are not relevant to this project.

A detailed assessment of acid sulphate soils and other highly reactive soils will be undertaken as part of future approvals processes and geotechnical investigations undertaken as part of the design phase.

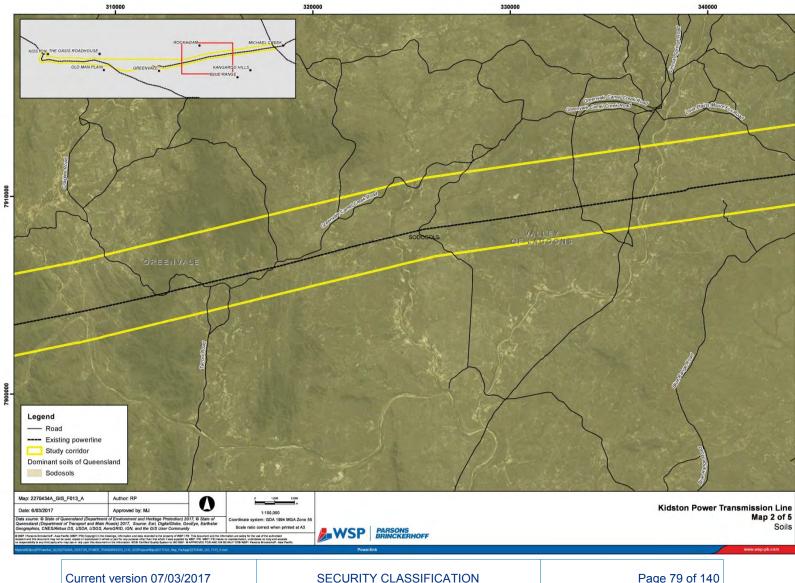
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Figure 13 – Soils

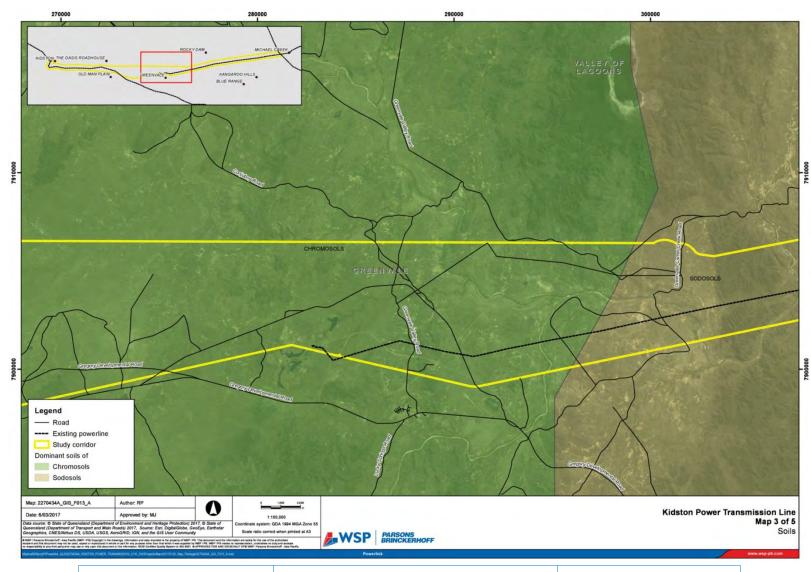






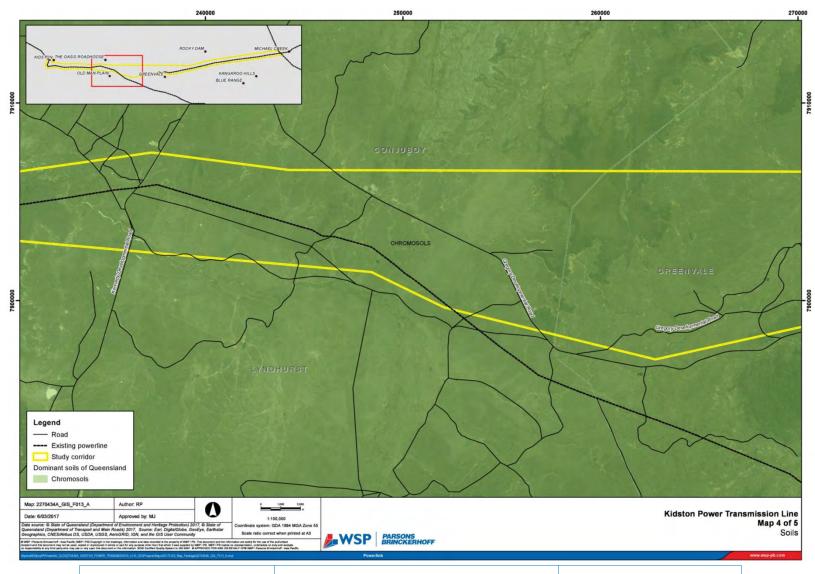
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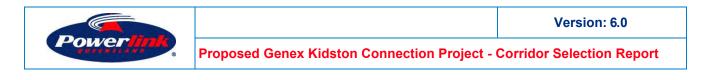


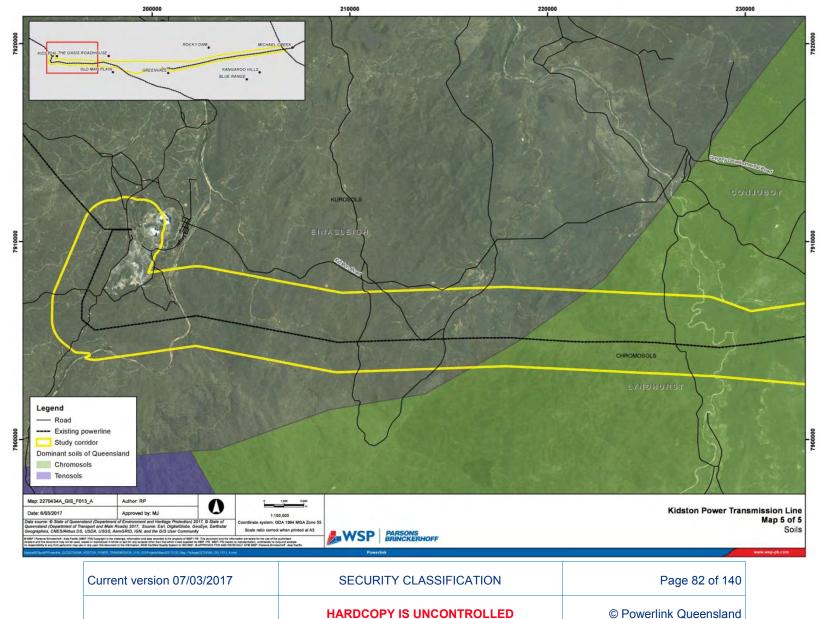
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## 10.8 CONTAMINATED LAND

A search of DEHP's Contaminated Land Register (CLR) and Environmental Management Register (EMR) was undertaken for parcels intersected by each of the corridors in early 2016 as part of the DCSR. No properties were listed on the CLR, whilst a number of properties listed are listed on the EMR for a notifiable activity which are provided in the table below.

The EMR listings are mostly for properties surrounding Kidston and activities were most likely associated with the previous mining undertaken there. It is also noted that some of these activities may no longer be occurring but may however still remain on the register. The listing is not considered a constraint, but is noted as an input to latter phases of the project.

Additionally, a notifiable activity on the EMR does not necessarily mean that the parcel (or part of) contains contaminated land, but rather that the activities have the potential to cause contamination. Furthermore, the register attaches an activity to a whole parcel, whilst specific activities may be confined to part thereof.

Further consultation and field investigations in subsequent detailed assessment processes will assist in revealing more detail about the extent of these activities. Consideration would likely be given to the footprint of operations, access and compatibility of electricity infrastructure with specific notifiable activities. At present there is negligible difference between the study corridor options in relation to potentially contaminated land.

Notifiable Activity	Number of Parcels Listed on EMR	
Notifuble Activity	Corridor B	Corridor C
Abrasive Blasting	3	3
Chemical Manufacture or Formulatio	n 3	3
Chemical Storage	3	3
Engine Reconditioning Works	3	3
Explosive Production or Storage	3	1
Gun Pistol or Rifle Range	1	1
Landfill	4	4
Livestock Dip or Spray Race	8	11
Metal Treatment or Coating	3	3
Mine Wastes	3	4
Mineral Processing	6	6
Petroleum Product or Oil Storage	5	6
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#### **Table 5 - Notifiable Activities**



Notifiable Activity	Number of Parcels Listed on EMR	
Smelting or Refining	3	3
Waste Storage Treatment or Disposal	1	1

\*Note: A single parcel may contain one or more notifiable activities.

In addition, it should be noted that due to the rural location of the project there is potential for illegal waste burial (e.g. poison drums) to be present.

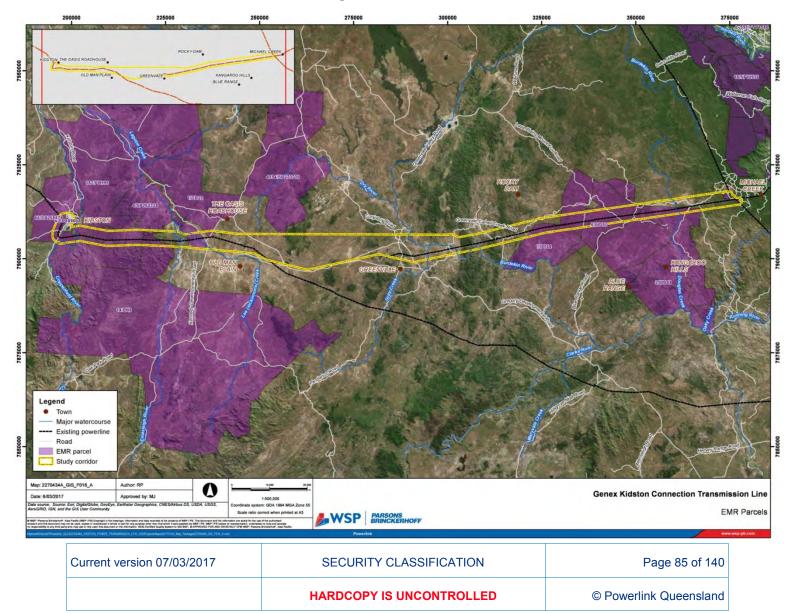
Powerlink's geotechnical engineers undertake investigations prior to construction and will therefore be able to advise on the presence of any unforeseen actual or potential contamination prior to excavation or other earthworks if Powerlink proceeds to further stages of project development. Should any risks be identified during these investigations then appropriate risk management procedures will be incorporated into an Environmental Management Plan (EMP) and implemented during construction. If contaminated material is discovered it will be managed in accordance with DEHP requirements.

Parcels registered on the EMR are shown in Figure 14.

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Figure 14 – EMR Parcels





## 10.9 UNEXPLODED ORDINANCE

Unexploded ordinance mapping is noted in the far eastern extent of the study area at Mt Fox. The mapping notes the use of the area for training purposes post World War II.

Unexploded ordinance mapping is broad and will not differentiate corridor options at the current level of assessment. The potential presence of unexploded ordinance will be assessed in more detail in subsequent phases of the project, supported by field verification studies as required.

## 10.10 HYDROLOGY AND WATER QUALITY

The topography within the project area ranges from flat low-lying land to steep crossings of multiple ranges (including the Great Dividing range) with elevation ranging from 400-800m AHD.

Each of the study corridor options cross the following major watercourses within the Burdekin River Basin:

- Douglas Creek
- Camel Creek
- Burdekin River.

Each of the study corridor options cross the following major watercourses within the Gilbert River Basin:

- Einasleigh River
- Lee (McKinnons) Creek
- Gray Creek (Option C only)
- Copperfield River.

All study corridor options cross potential flood-prone areas west of the Gregory Development Road to Kidston. These areas are associated with the Copperfield and Einasleigh Rivers.

The likelihood of impacting hydrology and water quality characteristics of waterways within the project area is considered low as the infrastructure can usually be designed to span over constraints without the need to clear or disturb beds and banks. Alluvial soils on the banks and approaches to watercourses are generally of a loamy sand nature, prone to erosion and dispersion when disturbed.

Riparian vegetation performs a vital role in times of flood, reducing flow velocities, resisting channel erosion, anchoring streamside top soil, and helping to trap and retain water borne top soil washed from upstream. Through transmission line design and careful use of topography it is possible to locate the infrastructure on high points to elevate the lines over streamside vegetation and minimise clearing of riparian vegetation.

General ground disturbance associated with the construction of transmission structures and access tracks within the corridors has limited potential to result in sediment entering waterways. Where possible, existing access roads with established and maintained erosion and control measures (culverts and spoon drains) would be used, minimising vegetation clearing and disturbance to soil structures. Option C offers the greatest apparent advantage in this regard, reducing clearing requirement through co-location and allowing shared access via existing maintenance tracks.

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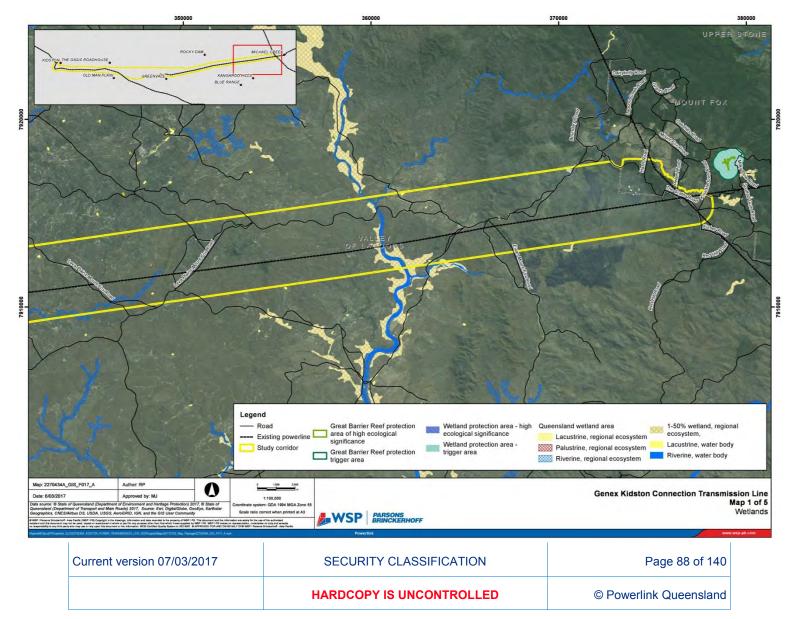
The suitability of the existing tracks to enable construction machinery to access sites will be assessed through future field reconnaissance and there may be repair and minor upgrades of tracks and crossings required.

Further assessment of impacts and mitigation measures will be undertaken as part of future detailed impact assessments and would be subject to further engagement with the relevant landholders and stakeholders.

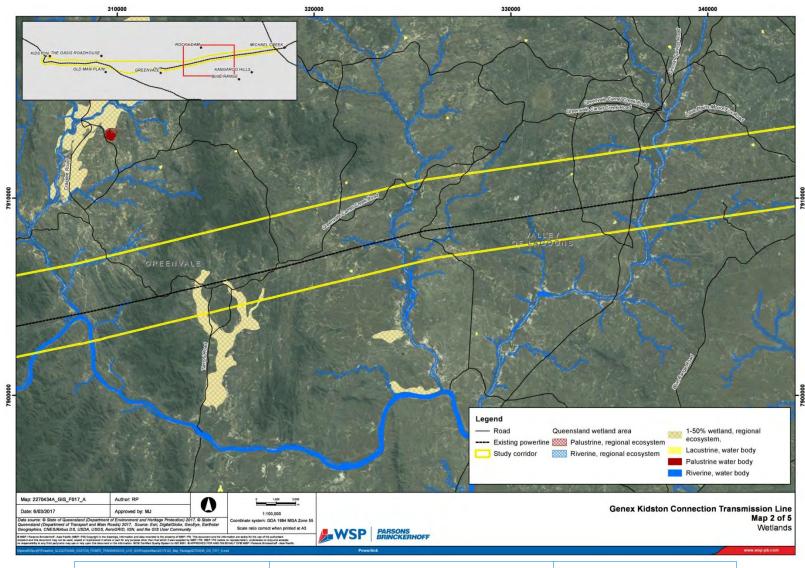
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Figure 15 – Wetlands

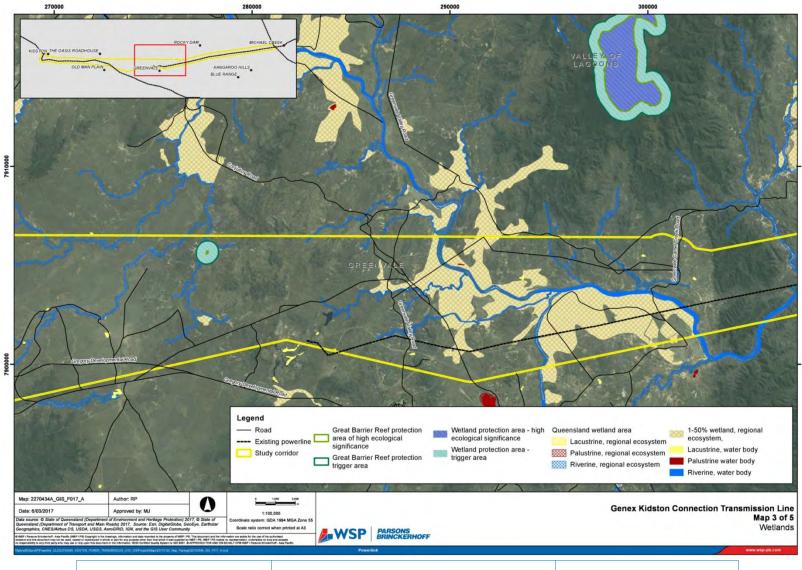






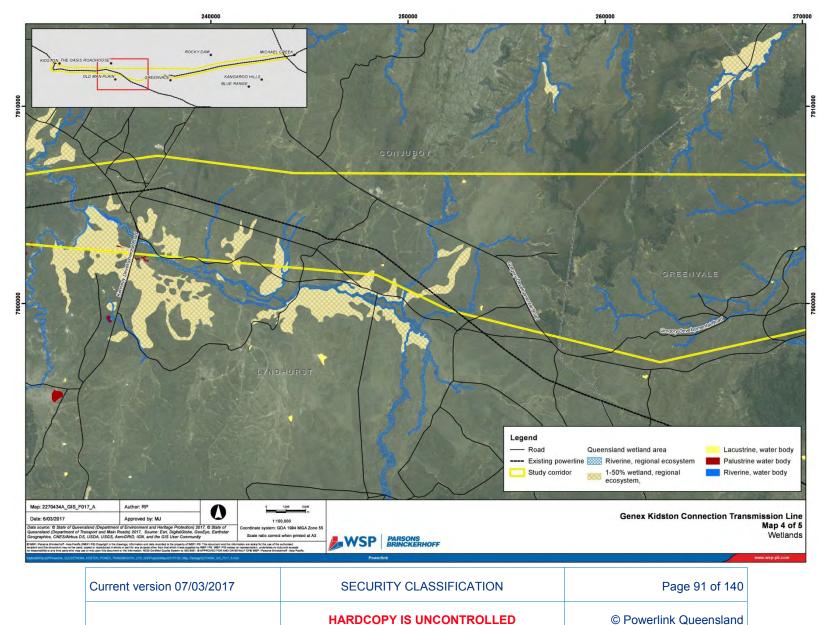
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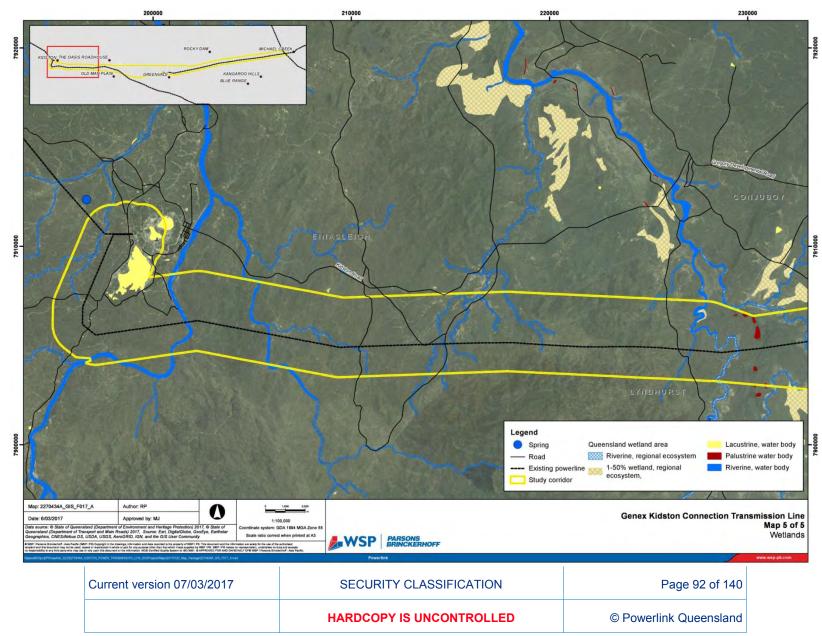


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# PROPOSED GENEX KIDSTON CONNECTION PROJECT

# **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"



## 11 ECOLOGY AND VEGETATION

Powerlink commissioned third party ecological advice in late 2016. The purpose of the advice was to assess the likely presence of MNES, Matters of State Environmental Significance (MSES) and other ecological constraints, providing a better understanding of potential environmental impacts for each study corridor option. The full ecological assessment is provided at Attachment 1.

The ecological advice included the three study corridor options from the DCSR. It was prudent to add study corridor A from the DCSR to the ecological assessment to validate the conclusions of desktop assessment in the DCSR that study corridor A had the highest relative environmental impact. It was as a result of the DCSR that study corridor A was removed from assessment in this CSR and the ecological advice provided an opportunity to confirm that removal of study corridor A had been undertaken on sound logic. The ecological advice confirmed that study corridor A had the highest potential for environmental impact (Attachment 1).

Due to the expansive nature of the study area and the early nature of these investigations, it was not feasible or justified to undertake field studies (ecological or otherwise) for the purpose of the CSR. Field studies will be part of any future detailed impact assessment process

In an endeavour to improve the ecological assessment, recognising that field studies were not feasible, Powerlink commissioned a small scope of remote sensing work over the region. The methodology of the remote sensing work is described in Section 11.1. The purpose of remote sensing was to define the physical extents of woody vegetation in the project area, which are not readily understood from other publicly available data sources.

Defining the extent of woody vegetation allows for primary quantification of potential vegetation impacts associated with the study corridor options. Understanding the physical extent of vegetation impacted also allowed the ecological advice to make more informed inferences about the type of vegetation and habitat potentially impacted.

## 11.1 REMOTE SENSING

Remote sensing was undertaken over the same area as the ecological advice, being the general extent covered by all study corridor options from the DCSR. Satellite imagery was utilised due to the vast size of the project area and limited availability of aerial imagery.

The methodology captured the extent and spatial distribution of current woody vegetation using an object based image classification process. This methodology has been previously deployed at a wide range of scales and using different remotely sensed source datasets for organisations across Australia and internationally. Landsat satellite imagery was used to derive landcover classification for the project extent from which indicative regional ecosystem mapping was generated. This was then aggregated into the mapping of woody vegetation for the project.

The final report associated with the remote sensing scope is provided at Attachment 2 and contains extensive detail about the methodology adopted. The output of greatest importance was spatial data representing the woody vegetation in the project area, which formed the basis of the ecological assessment.

#### 11.2 ECOLOCIAL ASSESSMENT METHODOLOGY

The general methodology adopted for the ecological assessment is described in Table 6.

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#### Table 6 – Ecological Assessment Methodology

Process	Comments	
Database searches	<ul> <li>EPBC Protected Matters Search Tool (by coordinates, buffer 5km)</li> <li>Wildlife Online (Figure 16)</li> <li>Atlas of Living Australia.</li> </ul>	
GIS analysis and mapping	All study corridor options buffered to 3km wide and interfeatures mapped and quantified: The number of Wildlife Online species records for thread listed under the NC Act and/or EPBC Act that have been precorridor investigation area. The number of Atlas of Living Australia's species recorrect under the NC Act and/or EPBC Act that have been precorridor investigation area. The presence or absence of 'high risk areas' for endant threatened plants (EVNT plants) listed under NC Act or survey trigger map' within each corridor investigation area. The extent (hectares) of MSES regulated vegetation in Department of Natural Resources and Mine's mapping (endangered and of concern regional concern regional ecosystems) Virtual GIS's remote sensing analysis and ma predicted and potential extent of remnant, regivegetation regulated remnant watercourse vegetation (erconcern regional ecosystems). The extent (hectares) of other MSES listed under Sche Offsets Regulation 2014 within each corridor investigation area. Wildlife habitats Strategic environmental areas High ecological significance wetlands High ecological value waters (watercourses) Environmental offsets	atened flora and fauna species en previously recorded within ds for threatened fauna listed eviously recorded within each agered, vulnerable and near in the 'protected plants flora rea. (DNRM) regulated vegetation al ecosystems) agered, of concern and least upping of woody vegetation – prowth and non-remnant indangered, of concern and least
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	Connectivity.	
Regulated remnant watercourse vegetation	1:100,000 scale regulated vegetation mapping published by the State of Queensland covers approximately two thirds of the project area, while the remaining third (the western extent) is covered by the 1:250,000 scale mapping. The 1:250,000 scale mapping does not have stream order attributes. Therefore, the Strahler method was applied to the 1:250,000 watercourse layer using GIS to appropriately assign stream orders.	
Threatened ecological communities	An assessment of the mapped regional ecosystems within each corridor was undertaken to identify the potential presence of regional ecosystems that may potentially constitute threatened ecological communities listed under the EPBC Act	
Constraint Ranking	A ranking system was applied for each environmental matter and/or potential ecological constraint, to capture the degree of ecological constraints within each study corridor option.           RED         Greatest amount of potential impact and ecological constraint	
	ORANGE         Moderate amount of potential impact and ecological constraint           GREEN         Least amount of potential impact and ecological constraint	
	The ranking system was not applied to database search results (i.e. Wildlife Online and PMST) because they have been used to inform species record searches and assessment. The ranking system was also not applied to matters that do not require significant impact assessments or trigger environmental offsets (i.e. least concern regional ecosystems and regrowth and areas of non-remnant). The 'high risk areas' for endangered, vulnerable and near threatened plants (EVNT plants) listed under NC Act on the 'protected plants flora survey trigger map', were not ranked because they were captured as flora species records.	

#### 11.3 POTENTIAL ENVIRONMENTAL IMPACT SUMMARY

Table 7 summarises the potential environmental impacts described in Sections 11.4 to 11.5.14. The summary is copied from the third party ecological assessment described in 11. The only variation is the removal of 3ha of impact to protected areas in study corridors A and C, resulting from the study area amendment described in Section 5.1.5.

The summary is based on a traffic light system and demonstrates that study corridor option C is preferred from an environmental perspective. The difference in potential impact between study corridors B and C is small for many criteria but the cumulative benefit of study corridor C is likely to be significant. Co-location options and shared access opportunities around study corridor C offer potential to further lower overall impacts during impact assessment and design phases.

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It is important to reiterate that study corridor A only appears in the ecological assessment to validate its removal from assessment at the DSCR stage. The traffic light system demonstrates that study corridor A had the potential for highest environmental impact by a reasonable margin.

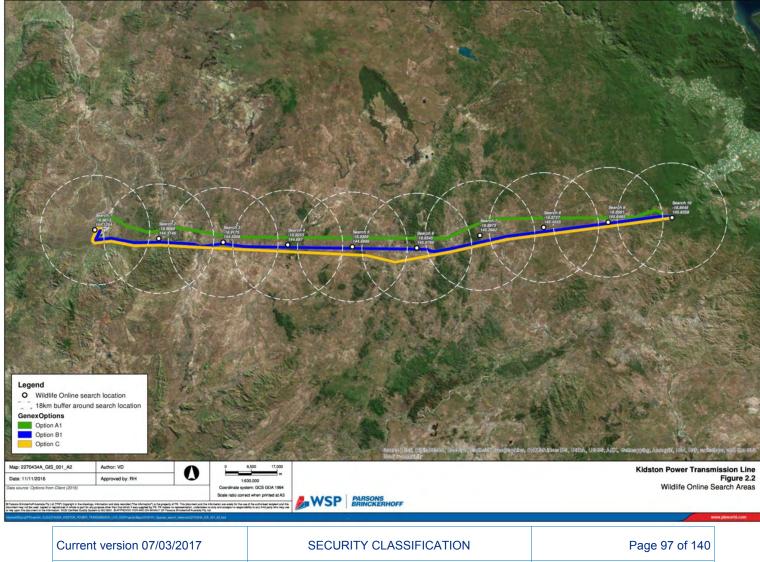
	CONSTRAINT RANKINGS		
Ecological constraint	Option A	Option B	Option C
Matters of State Environmental Significance			
NC Act threatened species records (numbers)	2	4	5
Regulated of concern dominant vegetation (ha)	2,059	1,819	1,519
Regulated of concern sub-dominant vegetation (ha)	1,008	442	334
Regulated of high value regrowth vegetation (ha)	3.6	2.4	2.4
Regulated Category R regrowth vegetation (ha)	422	296	257
Mapped of concern dominant regional ecosystems (ha)	1,008	442	334
Mapped of concern dominant regional ecosystems (ha)	2,059	1,891	1,519
Predicted of concern dominant woody vegetation (ha)	1,535	1,460	1,223
Predicted of concern sub-dominant woody vegetation (ha)	549	295	618
Potential of concern dominant woody vegetation (ha)	240	344	431
Potential of concern sub-dominant woody vegetation (ha)	364	87	37
Regulated remnant watercourse Wet Tropics bioregion (ha)	118	50	45
Regulated remnant watercourse vegetation Einasleigh Uplands bioregion (ha)	5,585	5,547	5,493
Protected areas (ha)	655	214	142
Wildlife habitats (ha)	522	411	385
High ecological significance (HES) wetlands (ha)	0	1.1	0
Matters of National Environmental Significance			
Threatened and migratory species records (numbers)	3	6	5
Proximity to Wet Tropics of Queensland heritage properties (km)	4.8	5.2	5.6

#### Table 7 – Potential Environmental Impact Summary

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Figure 16 – Wildlife Online Searches



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#### 11.4 MATTERS OF NATIONAL ENVIRONMENTAL SIGNIFICANCE

#### 11.4.1 EPBC ACT THREATENED AND MIGRATORY SPECIES

Database searches identified few records of threatened and migratory species. A low number of records was expected due to the remote and largely undeveloped and unsurveyed (ecological survey) nature of the project area. There is however the potential for additional species to be identified in subsequent impact assessment stages.

The records retrieved have been mapped for each study corridor option (Figure 3.1 - 3.3 in Attachment 1).

The potential to identify other species does not dictate that a referral under the *EPBC Act 1999* should, or should not, occur. It is rather a gap in existing database information. In an effort to address the shortfall in existing information, Powerlink commissioned a likelihood of occurrence assessment.

#### 11.4.1.1 LIKELIHOOD OF OCCURRENCE ASSESSMENT

The likelihood of occurrence assessment (Attachment 3) built on the third party ecological advice described in Section 11.2. It related mapped vegetation types to fauna and flora species known to be associated with those features in the broader region well beyond the CSR study area, targeting more developed areas where species records are more readily available.

The likelihood of occurrence assessment represents best endeavours to identify flora and fauna species potentially occurring in the study area and responds to feedback from the Department of Environment and Heritage Protection in January 2017 suggesting such an investigation.

This additional assessment confirmed the initial ecological advice procured for the CSR and identified only a relatively small number of additional EPBC species that may occur in the study area. Importantly, no new Threatened Ecological Communities or World Heritage Values were identified through this additional assessment.

It is important to note that the likelihood of occurrence assessment does not influence the preferred study corridor at the current level of assessment because vegetation communities and habitats occur in a broad and homogenous manner across the CSR study area.

Future ecological survey approaches will be designed to reflect the potential for additional species and provide for an adequate survey intensity to explore the ecological values of the project area. Appropriate management measures can then be designed on the basis of survey results. A range of management and mitigation measures are available in the development of transmission lines and terrestrial impacts can be minimised in most circumstances.

The likelihood of occurrence assessment does not dictate that a referral under the *EPBC Act 1999* should, or should not, occur. Detailed discussion of the referral process is provided in Section 9.1.1.1.

#### 11.4.2 THREATENED ECOLOGICAL COMMUNITIES (TEC)

Broad Leaf Tea-tree (Melaleuca viridiflora) woodlands in high rainfall coastal north Queensland was the only TEC revealed by the PMST as potentially occurring within the project area. The Broad Leaf Tea-tree TEC in

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the Wet Tropics and Central Mackay Coast Bioregions is associated with varieties of regional ecosystems described as 7.3.8/7.5.4 (floodplain), 8.3.2, 8.5.2 or 8.5.6 (*Melaleuca*).

The ecological assessment determined that the likelihood of The Broad Leaf Tea-tree TEC occurring in the project area is low as the associated regional ecosystems are not present. Based on the assessment undertaken to date the risk of impacting TECs is considered low for all study corridor options.

#### 11.4.3 WET TROPICS OF QUEENSLAND – WORLD, NATIONAL AND INDIGENOUS HERITAGE PROPERTIES

The ecological assessment considered proximity of the proposed line to the World Heritage-listed Wet Tropics. None of the study corridors impact the World Heritage area. Option A is the closest, located 4.8km at its closest point. At that distance, a transmission line is highly unlikely to impact the Wet Tropics in any manner (see Figure 3.65 in Attachment 1).

#### 11.5 MATTERS OF STATE ENVIRONMENTAL SIGNIFICANCE

#### 11.5.1 NC ACT THREATENED SPECIES

Wildlife Online database search results revealed that 26 threatened species listed under the NC Act and/or EPBC Act have been previously recorded within the search area, including 11 birds, six mammals and nine plants. Threatened species listed as endangered, vulnerable and special least concern under the NC Act are recognised as MSES under the *Environmental Offset Act 2014*. Near threatened species do not need to be assessed in terms of significant residual impacts under the *Environmental Offset Act 2014*, but have been included, just in case of any future changes in legislation that may occur during the approval pathway of the project.

As noted for EPBC Act threatened and migratory species (Section 11.4.1), the largely undeveloped and unsurveyed (ecological survey) nature of the project area dictates that species records are likely to be partial. There is potential for additional species to be identified in subsequent impact assessment stages.

Future ecological survey approaches will be designed to reflect the potential for additional species and provide for an adequate survey intensity to explore the ecological values of the project area. Appropriate management measures can then be designed on the basis of survey results. A range of management and mitigation measures are available in the development of transmission lines and terrestrial impacts can be minimised in most circumstances.

Appropriate management measures can then be designed on the basis of survey results. A range of management and mitigation measures are available in the development of transmission lines and terrestrial impacts can be minimised in most circumstances.

The records retrieved have been mapped for each study corridor option (Figure 3.1 - 3.3 in Attachment 1).

#### 11.5.2 PROTECTED PLANTS

High risk areas for endangered, vulnerable and near threatened plants (EVNT plants) listed under NC Act on the 'protected plants flora survey trigger map' are presented in Figure 3.4 in Attachment 1. The high risk

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areas indicate confirmed records of *Acacia tingoorensis* and represent a potential constraint to all three study corridors.

Subsequent impact assessment processes will need to target these areas and flora surveys will need to be undertaken in accordance with the *Flora Survey Guidelines – Protected Plants Nature Conservation Act 1992.* Powerlink has extensive experience in designing and developing infrastructure to minimise impacts to EVNT plants.

#### 11.5.3 REGULATED VEGETATION

#### **Category R Vegetation**

DNRM's *Category R regrowth vegetation: A self-assessable vegetation clearing code* states that Category R regrowth vegetation is native woody vegetation on freehold land, Indigenous land or leasehold land granted for agriculture or grazing purposes, located within 50 metres of a watercourse in the Burdekin, Mackay, Whitsunday and Wet Tropics Great Barrier Reef catchments.

It is prudent to consider potential impacts to Category R vegetation, irrespective of the legislative treatment of Category R under the latter approval process for the project which may exempt the project dependant on the pathway chosen. Study corridor option C impacts the least amount of Category R vegetation. The physical extents impacted are described in Section 11.3.

#### 11.5.4 DNRM REGIONAL ECOSYSTEMS

No endangered regional ecosystems are mapped in the study area. Study corridor option C impacts the least of concern regional ecosystem based on DNRM mapping. The physical extents impacted are described in Section 11.3.

Regional ecosystem mapping is undertaken by broad scale processes, with most maps prepared at a scale of 1:100,000 and based on the best available information. There are established processes for broad scale regional ecosystem mapping to be refined. Survey information collected during project development by Government and the private sector is used to amend the high level mapping, providing a clearer picture of the actual ecological makeup of the region.

The Genex Kidston Connection Project study area is largely undeveloped and therefore does not benefit from the same refinement of regional ecosystem mapping as more intensely developed regions. It is for this reason that the remote sensing scope described in Section 11.1 was undertaken. Remote sensing provides limited insight into the actual regional ecosystem communities present, but it does serve to rationalise the physical extent of vegetation in an otherwise largely unexplored region.

Higher regard is given to an assessment of potential vegetation impacts based on the intersection of regional ecosystem mapping with the results of the remote sensing exercise.

# 11.5.5 WOODY VEGETATION ASSESSMENT (REMOTELY SENSED)

The woody vegetation assessment used GIS remote sensing to predict the extent of woody remnant vegetation, and the potential extent of woody remnant vegetation and regrowth vegetation in each study corridor.

The assessment broadly involved:

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- Physical identification of woody vegetation by remote sensing
- Removal of areas mapped as regional ecosystem where no vegetation was present
- Addition of areas where vegetation was present but not mapped in regional ecosystem mapping
- Attribution of remotely sensed vegetation with DNRM mapped regional ecosystem descriptions where the two intersected.

This process provides the best possible understanding of vegetation cover and potential impacts without the intensive investment of time and resources associated with a full impact assessment process. The description above is a summary in simple language for the purpose of this report. The full methodology is available in Attachment 2.

Study corridor C is the least constrained in terms of potential impacts upon 'of concern' dominant regional ecosystems. Study corridor B is least constrained in terms of potential impacts upon 'of concern' subdominant regional ecosystems. The cumulative impact to 'of concern' regional ecosystems is similar for both options.

Greater emphasis should be applied to the predicted of concern dominant remnant vegetation, which therefore suggests that study corridor C is least constrained in terms of regional ecosystems. The physical extents impacted are described in Section 11.3.

In interpreting the table it is useful to understand that 'predicted' woody remnant vegetation is those areas where woody vegetation was verified within the regional ecosystem mapping. Potential woody remnant vegetation is those areas identified by remote sensing of similar composition to other remnant vegetation but not currently mapped as regional ecosystems. It is expected that these areas will be verified as regional ecosystems in subsequent field surveys.

All study corridor options involve clearing. Subsequent impact assessment processes will focus heavily on the extent of impacts to vegetation. Option C is considered favourable not just for lower potential physical impact to vegetation, but for the benefit of co-location and the potential to further reduce those potential impacts by refining design of a shared corridor.

Mapping of potential regional ecosystem impacts based on the woody vegetation assessment are provided in Attachment 1.

# 11.5.6 REGULATED REMNANT WATERCOURSE VEGETATION

Regulated remnant watercourse vegetation occurs throughout the study area. A distance from the defining bank of the mapped watercourse is specified in regulation that defines a distance from the defining bank of the watercourse within which remnant vegetation represents remnant watercourse vegetation. The distance varies by bioregion and stream order.

Study corridor C has the lowest potential impact to remnant watercourse vegetation. The physical extent of potential impact is described in Section 11.3.

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#### 11.5.7 PROTECTED AREAS

Several protected areas under the *Nature Conservation (Protected Areas) Regulation 1994* were identified in the study area. These areas are nature refuges (Newcastle Range – The Oaks and Liefway) and National Parks (Girringun). Study corridor C intersects the lowest total extent of protected area.

The extent of impact to protected areas associated with study corridor C was lowered by the amendment to the study area described in 5.1.5, which remove the Liefway Nature Refuge from the study area entirely, benefitting study corridors A (removed at DCSR stage) and C. The refuge fell outside of study corridor B.

Study corridor C intersects the lowest amount of the Girringun national park area based on existing published boundaries, with about 70ha less of the protected area intersected than study corridor B.

#### 11.5.8 WILDLIFE HABITAT

Wildlife habitat is mapped by the State and broadly includes:

- Threatened wildlife that is classified as endangered or vulnerable
- Special least concern animals (iconic) under the NCA.

Species mapping adopts wildlife habitat using, in order of preference:

- Essential habitat mapped under the VMA
- Modelled habitat (peer reviewed and accepted) mapped using climate, elevation, bioregion, and regional ecosystems
- Point records (buffered to 1,000m) that intersect with native vegetation (remnant or regrowth regional ecosystems) where it overlays a species record location.

Study corridor C intersects the least mapped wildlife habitat. As noted in Section 11.5.1 there is potential for additional NC Act species to be present in the project area due to a lack of historic surveys. The mapped wildlife habitat is based partially on these records and there is also logically potential for additional wildlife habitat to be present.

Notwithstanding limitations in the data, study corridor C impacts the least vegetation and benefits from colocation with existing transmission lines. It is likely that these features will also minimise potential habitat impacts. Habitat values will be explored in detail in subsequent impact assessments.

#### 11.5.9 STRATEGIC ENVIRONMENTAL AREAS

Strategic environmental areas do not exist in the study area.

#### 11.5.10 HIGH ECOLOGICAL SIGNIFICANCE (HES) WETLANDS

Only study corridor B intersects an identified HES wetland area. The physical extent of the intersection is described in Section 11.3.

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#### 11.5.11 HIGH ECOLOGICAL VALUE WATERS (WETLAND)

High ecological value waters (wetland) areas do not exist in the study area.

#### 11.5.12 HIGH ECOLOGICAL VALUE WATERS (WATERCOURSES)

High ecological value waters (watercourse) areas do not exist in the study area.

#### 11.5.13 ENVIRONMENTAL OFFSET AREAS

Environmental offset areas do not exist in the study area.

#### 11.5.14 CONNECTIVITY

Connectivity is assessed as significant residual impacts under the Queensland Government's Environmental Offsets Policy version 1.1 (2014), once a project footprint has been determined. The residual impact is assessed using GIS and the Queensland Government's Landscape Fragmentation and Connectivity tool for an individual impact area. Essentially, the assessment is based upon the level of fragmentation to remnant vegetation as a result of a project's impact area(s).

All study corridor options would result in impacts to connectivity. On this basis, once a preferred study corridor option is selected and a project impact area(s) is defined during subsequent impact assessment processes, the impact upon connectivity will need to be assessed using the Landscape Fragmentation and Connectivity tool.

As all options impact connectivity and detailed impacts are subject to significant change, connectivity does not differentiate study corridor options.

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## 12 CONSTRUCTABILITY

Overall the study area is of rural character, dominated by timbered areas, with some areas cleared for cattle grazing. The topography of the study areas is varied, with a large portion of the area being level or gently undulating plains. There are steeper sections on both corridors, including escarpments. This may present a challenging environment for construction and associated vehicle/plant access and consideration has been given to avoid where possible. Concrete batch plants will be required for the construction of the transmission line foundations.

Construction of a substation site at Mt Fox may present challenges due to topography within the area. A flat site is required for substation, which would likely require some level of cut and fill earthworks to achieve. Further detail is provided in Section 12.3. Despite potentially challenging topography, Mt Fox remains the closest viable connection point to the existing electricity transmission network (Section 5.1.3).

Constructability considerations remain similar across all draft corridor options and do not serve to differentiate options at this level of investigation. Powerlink is the foremost authority in transmission line and substation development in Queensland and will support Genex in the development of best for project design solutions.

## 12.1 ROAMES DATA REVIEW

Ergon has compiled a comprehensive virtual model of their network using aerial survey techniques. Software called ROAMES, developed by spatial consultancy Fugro, is used to display and navigate this model.

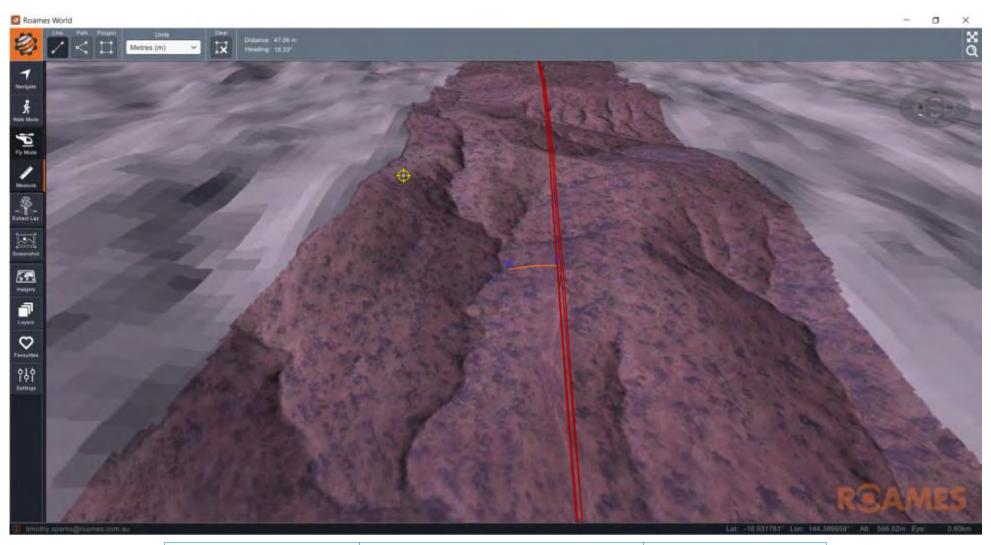
Powerlink contacted Fugro to enquire about data available in the Genex project area. The largely undeveloped nature of the study area means that spatial data available in the public realm is limited. Data was available for the existing Ergon 66kV and 132kV feeders in study corridor C and Powerlink was afforded the opportunity to view the ROAMES system on January 6, 2017 at the Fugro offices in Eight Mile Plains.

The principal benefit of viewing the ROAMES system was to gain greater insight into the variable terrain that might be encountered. Several areas of interest were identified and a screenshot for each was captured from the ROAMES system at those locations, including coordinates so the locations can be located and investigated in subsequent impact assessment phases (Figure 17).

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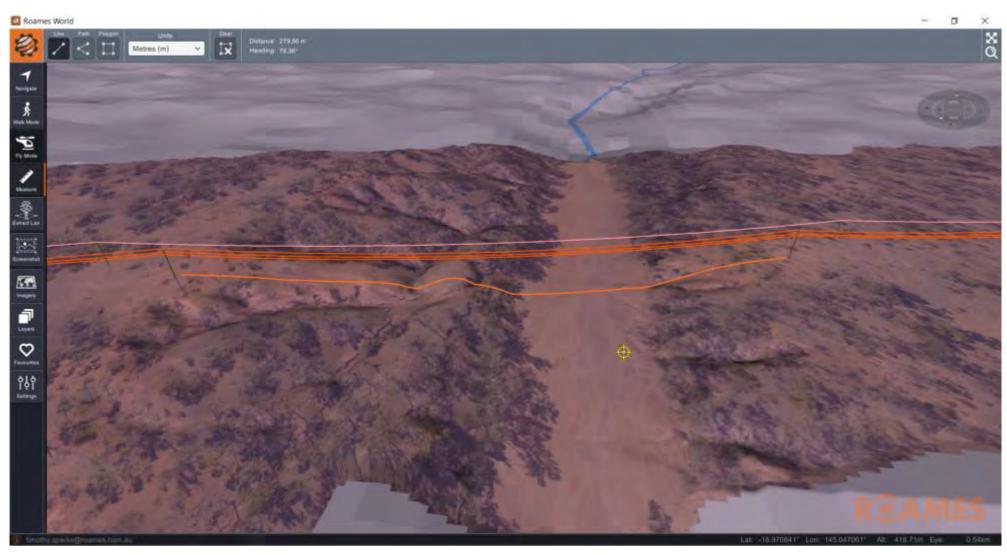
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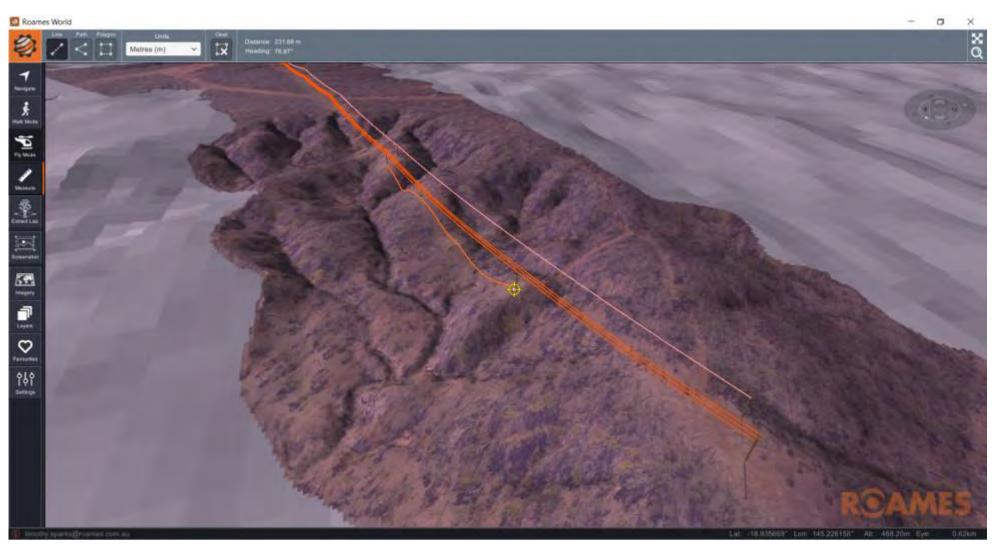
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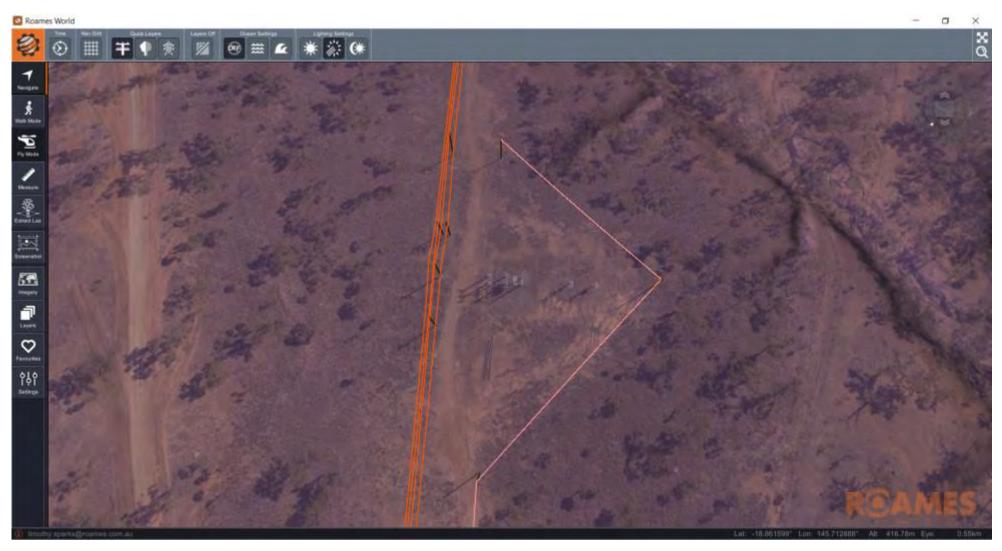
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## 12.2 HELICOPTER FLYOVER

Powerlink undertook an aerial inspection of the study area by helicopter in mid-January 2017. A helicopter was selected to minimise interference to landholders and their operations. All landholders were contacted in advance of the flight and any concerns raised were addressed by increased separation (altitude).

The fly over did not reveal any material constraints to delivery of the project. It was evident that the terrain is challenging in some locations, but through careful route selection and site specific design, terrain challenges can be overcome. Subsequent impact assessment and design phases will delve into greater detail around localised terrain and appropriate design responses. Challenging terrain is likely to occur along each study corridor and does not serve to differentiate options at the current level of assessment.

The flight supported that the study area amendment west of Greenvale discussed in Section 5.1.4 would facilitate constructability by avoiding escarpments. The preliminary alignment in this area will be outside of the initial extent of study corridor option C in response to the terrain that was not identifiable in the DCSR. No new properties were impacted as a result of the amendment and no material change to this assessment is realised by the deviation. See Section 5.1.4.

It was also evident from the fly over that recent rainfall had made much of the area difficult to traverse. Subsequent project planning will need to regard wet seasons and allow for restricted mobility through the area during those times. It was also evident that much of the soil was highly erodible, which will require further investigation during subsequent project phases. See Figure 18 and Figure 19.



Figure 18 – Swollen Creeks

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## 12.3 MT FOX SITE OPPORTUNITIES

Constructability for the substation at Mt Fox was considered as part of this CSR. The assessment included identification of high level site opportunities and constraints based on landholder engagement, affording an understanding of the likely availability of a suitable substation site. It was not in the scope of this CSR to negotiate acquisition of a site or formal option over a site. No substation site has been selected.

Terrain in the Mt Fox area is undulating. There are likely to be sites that are suitable for a substation, but some care will be required in selecting one with suitable grades and access. Despite potentially challenging topography, Mt Fox remains the closest viable connection point to the existing electricity transmission network (Section 5.1.3).

Proximity to the existing 275kV Powerlink Ross to Chalumbin transmission line is also an important consideration. A site west of the existing line is preferable from a network configuration perspective, whether adjacent or nearby. It is partially for this reason that the study area amendment discussed in Section 5.1.5 was undertaken, with property east of the existing 275kV being less desirable for the substation.

Finally proximity to the existing Ergon 66kV line is desirable so that co-location of infrastructure begins at Mt Fox and the benefits of co-location apply to as much of the proposed Genex Kidston connection as possible. An area of interest for the substation has been identified, bounded generally by the refined study area and existing Powerlink and Ergon networks. The area is depicted in Figure 21.

Subsequent project phases would seek to identify a specific site in the area of interest through a targeted and detailed assessment process, involving higher resolution terrain data (and imagery if available) than what is currently available, and further engagement with landholders. The site selection process would consider matters including, but not limited to:

Drainage

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- Cut and fill volumes
- Environmental footprint
- Availability of clean fill
- Ability to import clean fill to site (access)
- Requirements for auxiliary works
- Operational parameters.

An option over a site may be considered during the site selection process, dependant on the outcomes of technical investigations and further landholder and stakeholder engagement. Powerlink is the foremost authority in substation development in Queensland and will support Genex in the selection of a site from the area of interest.

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## 13 ENGAGEMENT FEEDBACK

Powerlink has undertaken early and targeted consultation with Federal, State and Local governments, peak bodies and landholders within the project area as part of preparing this CSR. The objective of stakeholder engagement at this point in the process is to raise awareness among relevant stakeholders and to gain meaningful input from those stakeholders to inform the CSR. In all cases, further and more extensive engagement and consultation will be undertaken in future stages of the project.

The following provides a summary of feedback received to date from stakeholders. Landholder engagement feedback is summarised in Section 13.7.

## 13.1 STAKEHOLDER WORKSHOP

Nine stakeholder organisations were invited to attend an CSR engagement workshop for the proposed Genex Kidston Connection transmission line project. The workshop on 21 November 2016 was attended by representatives of six stakeholder organisations.

The purpose of the workshop was to engage these stakeholders in identifying opportunities and constraints in the study area for the Genex Kidston Connection transmission line project Corridor Selection Report.

Representatives from the following organisations attended the workshop:

- North Queensland Conservation Council
- AgForce
- Department of Infrastructure, Local Government and Planning
- Department of Natural Resources and Mines
- Charters Towers Regional Council
- Etheridge Shire Council.

The following organisations declined the invitation to attend the workshop:

- Department of Environment and Heritage Protection
- Department of National Parks, Sport and Racing
- North Queensland Dry Tropics.

Drawing on their expert and local knowledge, workshop participants identified detailed opportunities and constraints applicable to the selection of a corridor for the Proposed Genex Power Kidston Connection transmission line. Key themes discussed are presented in Table 8. Detailed participants' input was comprehensively recorded at the workshop and informed the development of the CSR.

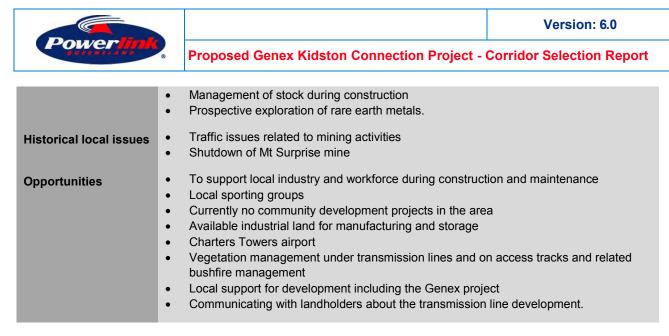
It is also noted that individual meetings were held with the Department of Environment and Heritage Protection and Department of National Parks, Sport and Racing (Section 13.2).

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#### Table 8 – Summary of workshop participant input

Key theme		Workshop participants provided input and in	nformation about:
Environment	<ul> <li>Bio</li> <li>Wa</li> <li>Bus</li> <li>Cor</li> <li>Fau</li> <li>min</li> <li>Geo</li> </ul>	getation including remnant vegetation and high value areas security including weed management and cattle tick management ter flow, erosion and sedimentation issues sh fire risk management nservation areas, wildlife refuges and National Parks una including pygmy glider, cassowary and redclaw yabbies (recreational in Kidston te ponds) ology including dormant volcano and lava tubes mate including drought impacts.	
Social	sma prov Mar Lan Fen Roa Air Visi Acc Lim Diff poo Acc at M Nat	nd use which is predominantly cattle grazing with some cropping on larger holdings; aller land holdings include chicken farm and hobby farms; includes organic aducers ny properties are run by family owned businesses ndholdings are predominantly leasehold ncing is limited on most properties ad conditions and accessibility; strips and wide use of aerial mustering vitors include fossickers at small active goldmines, 4WD enthusiasts and retirees cess tracks are used by shooters and 4WD enthusiasts nited local education options for students, schools and School of the Air ficulties with connectivity due to limited mobile phone and Internet coverage, and or radio reception commodation exists at the former Kidston mine, Junction, Greenvale and camping VIt Fox site tive title and Indigenous Access Agreements (ILUA's) in place on some properties ligenous and European Cultural Heritage sites cent defence force plans and associated land acquisition.	
Economic and technical issues	<ul> <li>Acc</li> <li>Opp</li> <li>Wo</li> <li>Cos</li> <li>Acc</li> <li>Rel</li> <li>Cor</li> <li>We</li> <li>Opp</li> <li>Opp</li> <li>Opp</li> <li>Opp</li> <li>Opp</li> <li>Opp</li> </ul>	rkforce access and retention sess to raw materials and water portunities for local industry and workers rkforce accommodation at of transport and alternative freight routes ressibility of stock routes iability and accuracy of mapping instruction traffic impacts to farmers and school bus ed wash down areas portunities for other solar and renewable development portunities downstream of the hydro dam bility of irrigation and agricultural projects that require	
Key stakeholders and communication	gov • Opt	keholders identified within the following groups: elected representatives, local vernment, interest groups, traditional owners, individuals tions for communication with landholders include mail, email, newsletters and uncil bulletins.	
Potential local issues	<ul> <li>Acc</li> <li>Tratility</li> <li>Imp</li> <li>We</li> <li>Aer</li> </ul>	<ul> <li>Acquisition of land process and recent experiences with defence force resumptions</li> <li>Traffic management including vehicle movements and driver behaviour</li> <li>Impacts on roads and road maintenance</li> <li>Weed management and wash down facilities</li> <li>Aerial mustering during project construction</li> </ul>	
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## 13.2 DEPARTMENTAL STAKEHOLDERS

# 13.2.1 DEPARTMENT OF ENERGY AND WATER SUPPLY & TREASURY

Powerlink met with representatives of the Department of Energy and Water Supply (DEWS) and the Treasury Department on 15 April 2016 as part of the DCSR. Advice was also provided to both departments on 11 November 2016 about the CSR and Powerlink's engagement intentions. Neither department raised major issues or concerns at this early stage in the process.

#### 13.2.2 DEPARTMENT OF ENVIRONMENT & HERITAGE PROTECTION

Powerlink met with representatives of the Department of Environment and Heritage Protection (DEHP) on 15 April 2016 to discuss the DCSR. A second meeting was held on 18 November 2016 to introduce the scope of this CSR.

Discussion was primarily held around DEHP's position on key areas of interest. These included protected flora and fauna, regional ecosystems and catchments/wetland/watercourses. Powerlink provided the third party ecological advice procured for this CSR to DEHP on 10 January 2017 and DEHP provided a response on 13 January 2017.

DEHP also noted that a likelihood of occurrence assessment would be a positive initiative to further identify species potentially occurring on the project area. Powerlink commissioned an assessment in response to DEHP feedback (Section 11.4.1.1). It is important to note that the likelihood of occurrence assessment will not influence the preferred study corridor at the current level of assessment because vegetation communities and habitats occur in a broad and homogenous manner across the CSR study area.

## 13.2.3 OFFICE OF THE COORDINATOR-GENERAL

Powerlink met with representatives of the Office of the Coordinator-General (OCOG) in late 2016. The OCOG is aware of the project through ongoing discussions with the proponent, Genex Power.

On 3 March 2016, the Minister for State Development declared the Genex Solar, Pumped Storage and Transmission Line project as a 'Prescribed Project' under the *State Development Public Works Organisation* 

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Act 1979. The Minister for State Development administers the SDPWO Act via the Office of the Coordinator General.

Discussion was primarily held around administrative functions of the department through the potential overarching approval process and appropriate approvals solutions. It was agreed that detailed discussion of approvals pathways was best postponed until this CSR was complete and a greater understanding of the study area was gained.

If the project was to proceed via the Coordinated Project process and Powerlink was also engaged to undertake subsequent work, we would continue to engage with the OCOG to achieve the respective project approvals and corridor acquisition. As the project progresses, a close working relationship with the OCOG will ensure delivery is done on a best-for-project basis.

#### 13.2.4 BUILDING QUEENSLAND

Powerlink met with representatives of Building Queensland (BQ) on 15 April 2016 as part of the DCSR. Advice was provided to Building Queensland on 16 November 2016 about the CSR. Building Queensland was advised there would not be any trigger for investment decisions involving Building Queensland at this point in the process.

It is noted that the Building Queensland process is likely to affect only Powerlink and the Queensland Government as potential proponents of the transmission line development process. Building Queensland may require oversight of a future business case for development of the transmission line project.

#### 13.2.5 DEPARTMENT OF TRANSPORT AND MAIN ROADS

No consultation has been undertaken to date with the Department of Transport and Main Roads given the early phase in the project development process. The proposed connection project traverses the Northern and Far North Transport and Main Roads districts.

No State roads are intersected in the Northern District. Two State roads are likely to be intersected in the Far North District, being the Kennedy and Gregory Development Roads. The roads will be spanned by conductors only where possible, with towers or other structures placed well clear of the road corridor. The potential for impact to State roads is therefore minimal. Where spanning is not possible due to significant width careful placement of structures will be required to minimise impact to the utility of the road corridor.

It is recommended that consultation with the Department of Transport and Main Roads occurs once subsequent impact assessment phases commence. Whilst State roads do not differentiate corridor options they do become influential as the project is refined and the location of access tracks, construction camps etc become clear.

# 13.2.6 DEPARTMENT OF NATIONAL PARKS, SPORT & RECREATION

Powerlink met with representatives of the Department of National Parks, Sports & Recreation (DNPSR) on 19 April 2016 as part of the DCSR. A second meeting was held for the CSR on 18 January 2017.

Girringun National Park (Mt Fox section) is an area of particular interest for DNPSR and amenity impacts on the park area will require assessment in latter project phases.

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#### 13.2.7 DEPARTMENT OF INFRASTRUCTURE, LOCAL GOVERNMENT & PLANNING

Powerlink met with representatives of the Department of Infrastructure, Local Government and Planning (DILGP) on 19 April 2016 as part of the DCSR. DILGP advised that they were not *aware of any* proposed developments or current development applications within the corridors at that time and raised only the non-statutory Gulf Regional Plan and future North Queensland Regional Plan for further consideration.

A DILGP representative attended the stakeholder workshop on 21 November 2016. No new development applications or issues around the Gulf and North Queensland Regional Plans were noted at the workshop. Other DILGP feedback is included in the summary notes from the workshop (Section 13.1).

Powerlink is aware that the North Queensland Regional Plan is currently under production and will monitor the status of the plan. Powerlink plays an active role in submitting to emerging regulatory frameworks and will monitor the North Queensland Regional Plan as part of this activity.

### 13.2.8 DEPARTMENT OF AGRICULTURE, ANDFISHERIES

Constraints mapping and project background information has been provided to DAF to undertake a desktop assessment and provide feedback on areas of interest to the department. It is recommended that a meeting be held early in future stages of the project to ensure any areas of interest to the department have can be addressed.

#### 13.2.9 DEPARTMENT OF ENVIRONMENT & ENERGY(CWTH)

Powerlink met with a representative of the Commonwealth Department of Environment and Energy (DoEE) on 22 April 2016 as part of the preparation of DCSR to obtain initial and high-level advice regarding any Commonwealth environmental considerations. Contact was re-established with the same representative in November 2016 and again in January 2017 after the third party ecological advice was completed.

Powerlink proactively undertook a likelihood of occurrence assessment for EPBC Act threatened and migratory species and NC Act EVNT flora and fauna (Attachment 3). This additional scope of work is designed specifically to supplement the low number of records returned from the Protected Matters Search Tool.

It is important to note that the likelihood of occurrence assessment will not influence the preferred study corridor at the current level of assessment because vegetation communities and habitats occur in a broad and homogenous manner across the CSR study area.

#### 13.2.10 DEPARTMENT OF NATURAL RESOURCES & MINES

Powerlink met with representatives of the Department of Natural Resources and Mines (DNRM) on 28 April 2016 as part of the preparation of the DCSR. Feedback in relation to dealing with resource interests indicated that consultation with interest holders would be the most beneficial way to address any potential issues or compatibility conflicts. Discussion focused on:

- Vegetation clearing governance relating to available project delivery frameworks
- Land tenure

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• Resource interests.

A DNRM representative also attended the stakeholder workshop on 21 November 2016. Discussion was held around similar issues at the workshop. DNRM feedback is included in the summary notes from the workshop (Section 13.1).

Further consultation with DNRM will be an important part of subsequent impact assessment processes as further detail around tenure, resource interest and vegetation impacts emerges.

#### 13.2.11 ERGON ENERGY

Powerlink made contact with regional representatives of Ergon Energy to gain further understanding of the study area. Feedback from Ergon has assisted in the development of this CSR and further engagement with a broader cross section of the organisation is recommended as early as possible in subsequent project phases.

## 13.3 ELECTED REPRESENTATIVES

Powerlink has provided initial advice about the CSR to all State and Federal Government elected representatives including Mr Shane Knuth, Mr Robbie Katter, Mr Andrew Cripps and Mr Bob Katter. Contact with these elected representatives occurred in November and December 2016. A project briefing was also provided to Mr Cripps on request, who made subsequent enquiries to Powerlink on behalf of a constituent.

## 13.4 STATE GOVERNMENT MINISTERS

Advice was provided to the following Ministerial offices about the project and Powerlink's engagement intentions:

- The Hon. Jackie Trad MP, Deputy Premier and Minister for Infrastructure, Local Government and Planning and Minister for Trade and Investment
- The Hon. Curtis Pitt MP, Treasurer, Minister for Aboriginal and Torres Strait Islander Partnerships and Minister for Sport
- The Hon. Mark Bailey MP, Minister for Main Roads, Road Safety and Ports and Minister for Energy, Biofuels and Water Supply
- The Hon. Coralee O'Rourke MP, Minister for Disability Services, Minister for Seniors and Minister Assisting the Premier on North Queensland.

## 13.5 LOCAL GOVERNMENT

#### 13.5.1 ETHERIDGE SHIRE COUNCIL

Powerlink held a teleconference with representatives of Etheridge Shire Council on 14 April 2016 and a representative of Council attended the stakeholder workshop on 21 November 2016. Powerlink also provided initial advice to Etheridge Shire Council Mayor and CEO on 18 November 2016 about the CSR and engagement activities and offered a formal briefing.

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This was followed up with a phone conversation with the Etheridge Shire CEO on 19 November 2016. A project presentation was provided via email to the Mayor and CEO on 22 December 2016, in lieu of a face-to-face briefing. Council is very familiar with the Genex project and have developed a working relationship with the proponent.

Council has approved the development application for the 50MW solar farm proposal on the Kidston tailings? Dam and continue to work with Genex on latter stages of the development.

Etheridge Shire has provided a range of feedback, much of which was included in the DCSR and has therefore contributed to this CSR. Council's feedback from late 2016 is included in the notes from the workshop (Section 13.1). No major issues or concern were raised at this early stage in the process.

#### 13.5.2 CHARTERS TOWERS REGIONAL COUNCIL

Powerlink met with representatives of Charters Towers Regional Council on 28 April 2016 as part of the DCSR. Powerlink provided initial advice to Charters Towers Shire Council Mayor and CEO on 18 November 2016 about the CSR and engagement activities and offered a formal briefing for full Council, which occurred on 13 December 2016. A representative of Council also attended the workshop on 21 November.

Relevant feedback is included within this report and in the notes from the workshop (Section 13.1). No major issues or concern were raised at this early stage in the process.

#### 13.5.3 HINCHINBROOK SHIRE COUNCIL

Powerlink met with representatives of Hinchinbrook Shire Council on 27 April 2016 as part of the DCSR. Powerlink also provided initial advice to Hinchinbrook Shire Council Mayor and CEO on 18 November 2016 about the CSR and engagement activities and offered a formal briefing for full Council, which occurred on 6 December 2016. No major issues or concern were raised at this early stage in the process.

## 13.6 PEAK BODIES

#### 13.6.1 QUEENSLAND FARMERS FEDERATION

Powerlink met with a representative of the Queensland Farmers Federation (QFF) on 17 November 2016. Powerlink subsequently provided QFF a list of potentially impacted properties for validation against the databases of its constituents and was able to verify that QFF members were not impacted by the study area.

#### 13.6.2 AGFORCE

A representative of AgForce attended the stakeholder workshop on 21 November, providing a host of valuable insights into the study area. AgForce feedback is included in the notes from the workshop (Section 13.1).

#### 13.6.3 QUEENSLAND RESOURCE COUNCIL

Powerlink contacted the Queensland Resource Council (QRC) in late 2016 to introduce the project. QRC advised that they had no feedback at the time of contact.

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#### 13.6.4 NQ CONSERVATION COUNCIL

A representative from the NQ Conservation Council attended the stakeholder workshop on 21 November, providing a host of valuable insights into the study area. NQ Conservation Council feedback is included in the notes from the workshop (Section 13.1).

#### 13.6.5 NORTHERN GULF RESOURCE MANAGEMENT GROUP

Powerlink contacted the Northern Gulf Resource Management Group (NGRMG) in late 2016 to introduce the project. NGRMG subsequently advised that they had no feedback.

#### 13.6.6 TOWNSVILLE ENTERPRISE

Powerlink provided project details to Townsville Enterprise on 13 January 2017 and sought further information in relation to the location of Hell's Gate Dam.

### 13.7 ABORIGINAL PARTIES

Input into this CSR has been sought from each of the relevant Aboriginal Parties (as defined under the *Aboriginal Cultural Heritage Act 2003*) as well as the Department of Aboriginal and Torres Strait Islander Partnerships (DATSIP) Cultural Heritage Unit.

Powerlink met with Townsville based representatives of DATSIP on 15th December 2016 to gain an understanding of any local Aboriginal cultural heritage issues and known constraints. Whilst no specific cultural heritage issues were raised, the importance of meaningful and early engagement with the relevant Aboriginal Parties was stressed along with the importance of operating under an agreement or CHMP with the Aboriginal Parties. No comments were provided to differentiate between study corridor options.

Telephone contact was made with representatives of the Gugu Badhun Aboriginal Corporation, representing the Gugu Badhun native title holders and with Tatampi Puranga Aboriginal Corporation, representing the Ewamian People#2 and Ewamian People#3 native title holders on 15 and 16 December 2016, respectively. Further details of the project were provided by email on 22 December. A subsequent teleconference was held with various members of the Gugu Badhun native title holders on 31 January 2017.

Gugu Badham representatives noted the presence of known Aboriginal cultural heritage sites in the area north of Greenvale and the 'Old Beef Road' but felt these would be unlikely to be an influencing factor the differentiation between corridor options at the current level of assessment. They also noted a general preference for co-location with the existing Ergon Energy easement.

Both Aboriginal Parties, however, stressed the need for detailed cultural heritage assessments once a corridor is determined. This will require early engagement with the parties in order to reach agreement (i.e. a signed CHMP or formal agreement) on assessment methodology and heritage protective measures. Negotiation for such agreement would commence prior to a final alignment decision.

## 13.8 LANDHOLDER ENGAGEMENT FEEDBACK

Landholders provided input to the CSR through interactions with Powerlink representatives during the period from November 2016 to mid-January 2017. Landholders were provided with a map of their property that

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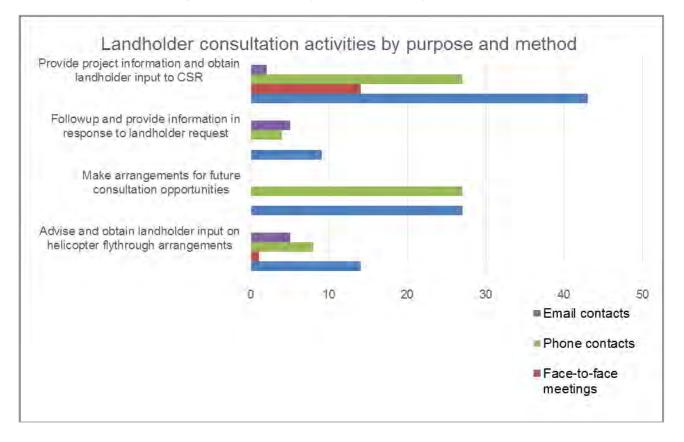
showed the extent of the study area for the CSR, constructed broadly from study corridor options B and C plus the amendments discussed in Sections 5.1.4 and 5.1.5.

Table 9 presents a summary of the input from landholders to the CSR. The input has been sorted into topic groups. Confidential, personal or private information provided by landholders is not presented in this report.

The matters most frequently raised by landholders were:

- Potential impacts on property operations including mustering activities, access and security
- Identification of constraints to the alignment including existing and proposed infrastructure and topography on their properties and in the adjoining area
- Most landholders considered co-locating with the existing ergon transmission lines as causing the least impact to their properties
- Biosecurity and weed management during construction and maintenance, and the movement of weeds throughout the life cycle of the proposed transmission line
- Line design, including conductor to ground clearance.

Figure 20 – Graph of Major Topics Raised by Landholders



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Topic raised by landholders	Detailed content of topic raised by landholders	
Potential impacts on property operations		
	Helicopter work and mustering	
	<ul><li>Helicopter maintenance of transmission lines</li><li>Loss of pasture/grass</li></ul>	
	Traffic impacts	
	Gate protocols	
	Access track maintenance/erosion	
	Disruption to telephone service	
	<ul> <li>Requirements for working on and traversing properties</li> </ul>	g registered organic status
	Potential for landholder owned generation	projects
	<ul> <li>Property value</li> <li>Biosecurity</li> <li>Proximity to residences</li> <li>Electromagnetic fields (EMF)</li> <li>Ongoing access to maintain the transmission line.</li> </ul>	
Constraints to the alignment location including existing and planned infrastructure and topography	Landholders identified the locations of existing infrastructure which may be impacted by the transmission line alignment including: dams, buildings, air strips, fences, gates, cattle yards, cattle watering points and associated infrastructure, pipelines, low voltage powerlines, houses, roads, rivers and community areas. Landholders identified the potential impacts on proposed infrastructure including	
Hell's Gate Dam, and suggested that there could be resumption greater Greenvale area by Defence.		resumption of properties in the
	Landholders identified natural features and topography which may im alignment of the proposed structure.	
	Landholders offered up solutions to many of the issues they identified.	
Co-location with Ergon transmission lines	Where relevant, most landholders considered co-locating with the existing Ergon transmission lines to have least impact to their operations, by paralleling the Ergon 132kV line all the way until it intersects the 275kV transmission line.	
Biosecurity and weed	Landholders stated their concern for the potential transmission of Parthenium weed,	
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management matters	Grader Grass and other weed seeds during construction and maintenance of the transmission line, and identified the need for wash-down stations. The need for biosecurity procedures and standards during construction and maintenance was identified.	
Line design and clearance preferences	<ul> <li>Landholders requested that during the design of the transmission line consideration be given to:</li> <li>Line clearance over roads for cattle trucks</li> <li>Line clearance from air strips</li> <li>Potential to place the line underground</li> <li>Heights, spans and tower design.</li> </ul>	
Importance of communication with landholders	Landholders stated the value of good communication with landholders during planning, construction and maintenance including access to the project decision makers.	
Potential impacts on property values	Landholders stated their concern that the proposed transmission line would impact their property value.	
Desire for local benefits from the project	<ul> <li>Landholders questioned whether the proposed transmission line would provide local benefits including:</li> <li>Improved reliability of their power supply</li> <li>Access into the electricity grid for their own solar projects</li> <li>Impacts to their power bills</li> <li>Opportunities for the location of construction base/camp</li> <li>Use of local contractors.</li> </ul>	
Compensation	Landholders raised fair compensation as a concern.	
Potential environmental impacts	Extent of vegetation clearing that would be required for the project and erosion control measures.	
Potential impacts on visual amenity	The need for screening the substation at Mt Fox.	

The vast majority of the topics raised during landholder engagement were issues to be managed in latter impact assessment phases. The top five topics included potential impacts to property operations, existing infrastructure/ topography, biosecurity, co-location and design/ clearance preferences.

The only one of these top five topics that influences the selection of a preferred study corridor from the study area discussed with landholders at the current, broad level of investigation is the desire from landholders that linear impacts are co-located. In this regard, study corridor C is preferred. All study corridor options are likely to have some level of interaction with, and potential impact on the other topics raised.

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Powerlink provided a detailed record of discussion to landholders, with a duplicate copy retained by Powerlink for its records. The records of discussion are not included in this CSR because they represent confidential discussions between Powerlink and the landholder.

Powerlink's commitment to landholders was that feedback would be included in the CSR. The CSR also serves as a record of the issues raised, which informs latter impact assessment processes. A copy of the finalised CSR will be made available to all landholders consulted on the project.

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## 14 STUDY FINDINGS

## 14.1 PREFERRED STUDY CORRIDOR

Study Corridor option C is the preferred study corridor based on the assessment in this CSR. Study Corridor option C is broadly preferred because it:

- Offers the lowest potential for environmental impact based on systematic and independent ecological assessment
- Offers the highest potential to further reduce overall impacts through refinement of design and route selection, maximising the benefits of co-location.

This CSR and the recommended corridor were developed based on a range of intensive desktop investigations, stakeholder and landholder engagement, along with virtual and aerial reconnaissance. No material constraint or opportunity was revealed during any of these investigations that would alter the preference for study corridor option C.

It is noted that a deviation from the original extent of study corridor option C is required between approximately Greenvale and Conjuboy to manage large escarpments identified during the development of this CSR. No new additional properties are impacted by the amendment (Section 5.1.4) and the preliminary alignment will reflect this deviation (Section 14.2).

## 14.2 PRELIMINARY ALIGNMENT

This CSR seeks to select both a preferred study corridor from the broader study area and a preliminary alignment to serve as the basis of further investigation during subsequent assessment processes. The preliminary alignment is an area of nominal width that can contain the final 60m easement required for the proposed 275kV transmission line.

The preliminary alignment is depicted in Figure 21 and is mapped at a nominal 120m wide around the centreline of the existing Ergon lines. The width of 120m represents sufficient area to place the proposed 275kV line either north or south of the existing Ergon lines depending on terrain and other local constraints. This width facilitates a notional clearance of 30m between both the centreline of the two transmission lines and the edge of a new easement for the 275kV line and is retained across the entire preliminary alignment for consistency.

The nominal 120m may require variation in specific areas where more than one Ergon line exists (local distribution lines were noted in parts of the Ergon transmission corridors). It may also require local variation where Ergon distribution transformers are present, where existing Ergon lines are heavily stayed or where topography dictates amendment. These areas cannot be defined with any certainty at the current level of investigation but are noted for completeness.

The preliminary alignment within study corridor option C is generally anchored by the existing Ergon transmission lines, which offer opportunity to reduce overall impacts through co-location. There are two areas where alternative alignments are required.

#### 14.2.1 MT FOX

The final substation location at Mt Fox will be identified during latter stages of project development. For the purposes of the preliminary alignment in the CSR, a general area of interest is identified for the substation.

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As noted in Section 12.3 it is likely that a substation with suitable grade and proximity to the existing Powerlink and Ergon network can be procured within the area of interest. The area of interest is depicted in Figure 21.

#### 14.2.2 GREENVALE TO CONJUBOY

There is an area of approximately 62 km between Greenvale and Conjuboy where the landholder engagement and aerial survey process provided an enhanced understanding of the local environment. In this region an alternative alignment is required to avoid terrain not identified in the DCSR, to minimise potential conflicts with the SCONI Phase 1 Project (Section 10.1.2.1) and to maximise colocation with the existing Ergon network. This area is depicted in Figure 4

#### 14.2.2.1 SCONI PHASE 1 PROJECT

Figure 9 shows graphically the minor deviation to avoid conflict with the SCONI Phase 1 Project. The preliminary alignment is currently located between the existing Stenhouse Dam in and the identified resource areas.

As noted in Section 10.1.2.1 direct contact with the tenement holder could not be established during the production of this CSR. There appears from desktop and aerial investigations however to be ample room for the mining use and transmission line to co-exist.

#### 14.2.2.2 WEST OF GREENVALE

As noted in Section 5.1.4 a study area amendment was adopted west of Greenvale to avoid escarpments not identified in the DCSR phase. The preliminary alignment in this area necessarily deviates from study corridor C as identified in the DCSR.

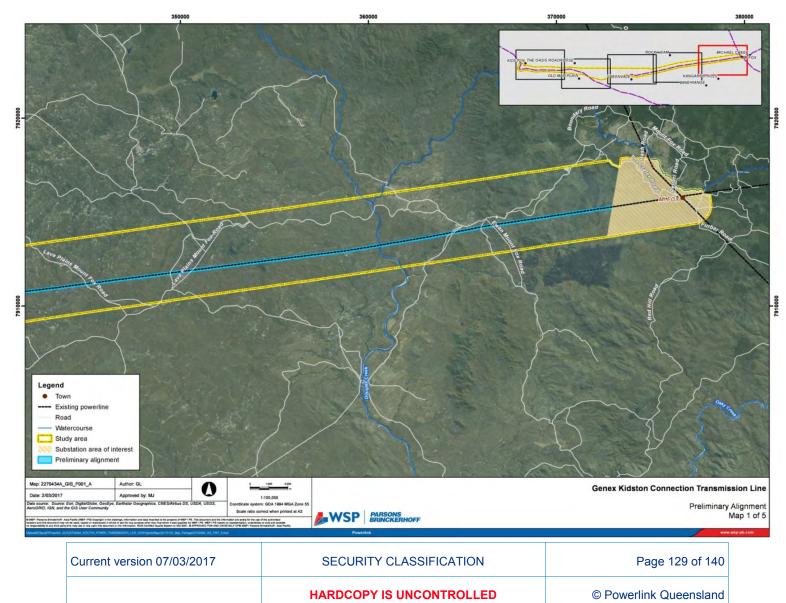
Figure 4 shows the preliminary alignment in this area. It is noted that no new properties were affected by the amendment. It is also noted that this area was not reflected in the third party ecological advice due to late identification of the need for alteration to the study area. Advice from the third party undertaking the ecological assessment is that the amendment would not materially affect the results of the investigation, with MNES and MSES occurring across the study area in a largely homogenous manner. The third party ecological advice reflects this in its Section 1.

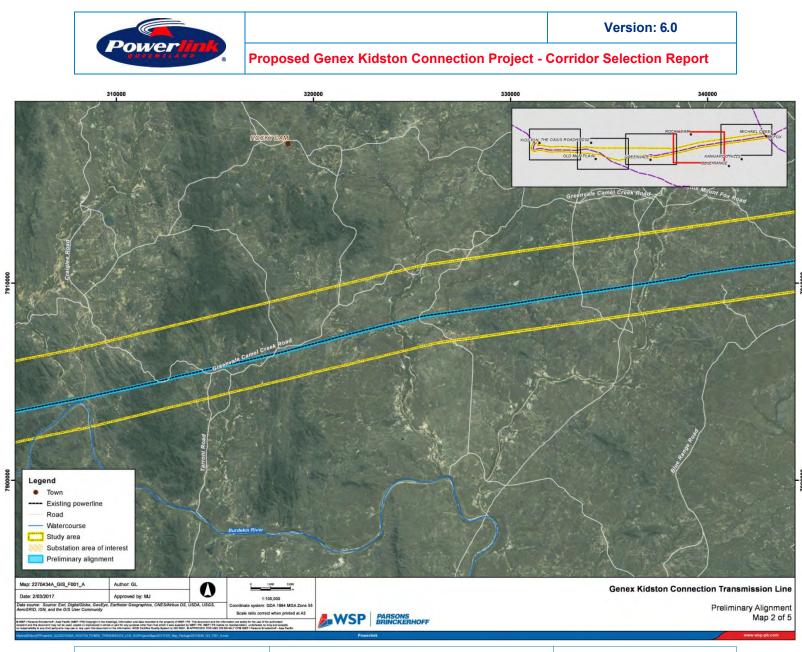
The preliminary alignment also departs from study corridor C immediately west of the amended study area discussed above. The departure of approximately 8 km is designed to maximise co-location with the existing network, which was contained in study corridor B from the DCSR for this localise region. The deviation from study corridor C, like the larger deviation west of Greenvale has no effect on the results of this CSR. It is likely instead to reduce overall impacts by maximising co-location opportunities.

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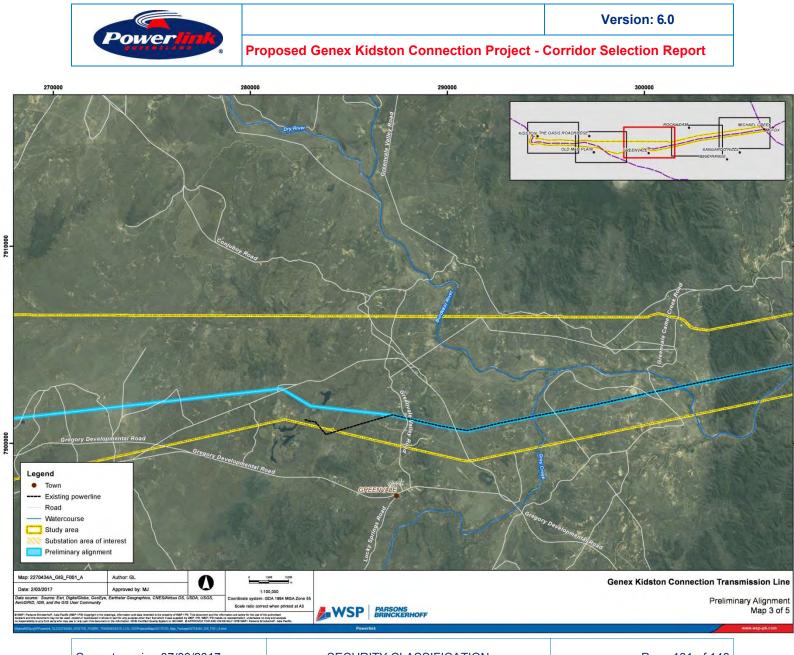


Figure 21 – Preliminary Alignment

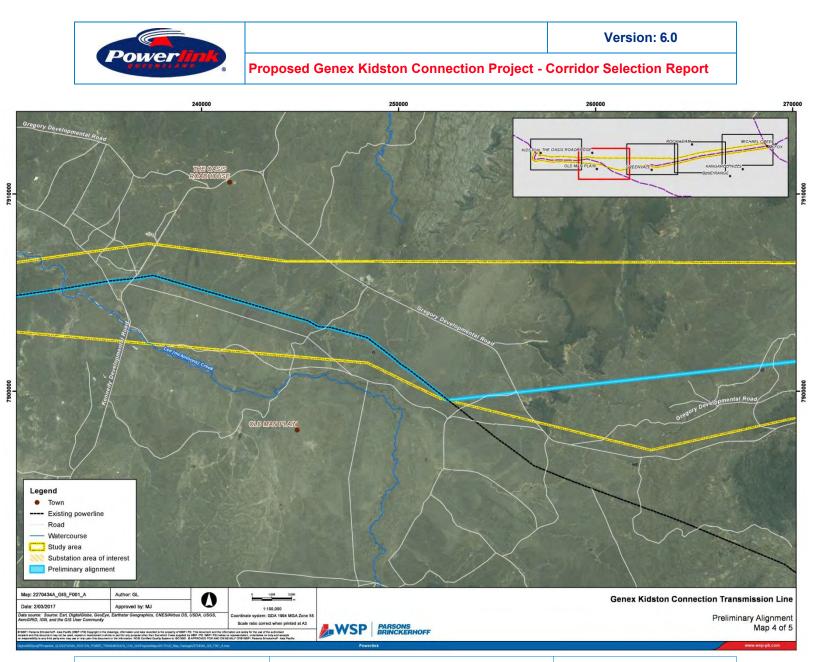




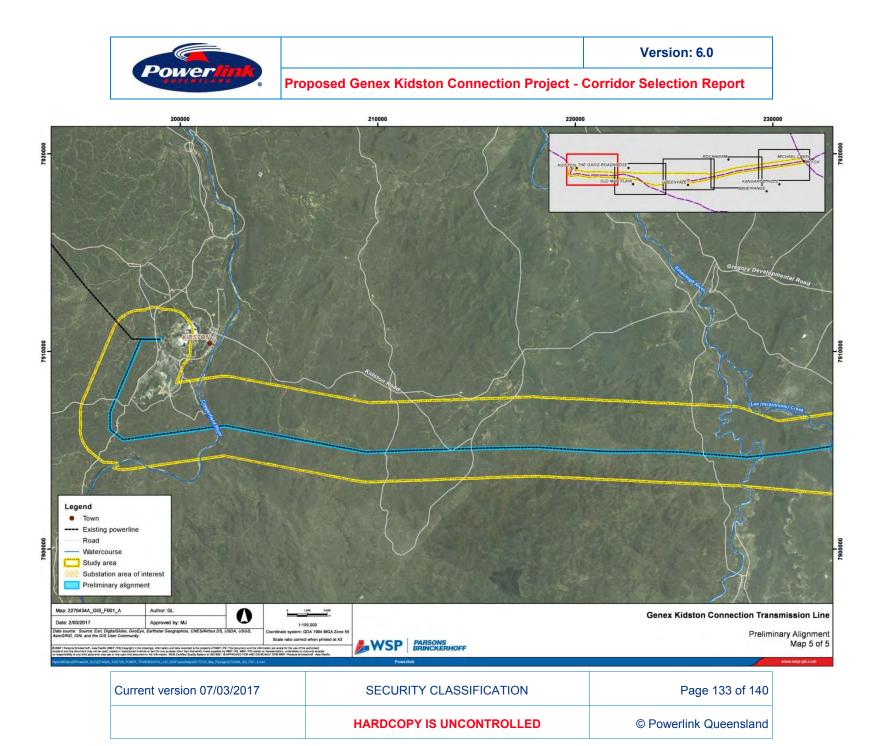
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## PROPOSED GENEX KIDSTON CONNECTION PROJECT

## **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"



## 15 FURTHER ASSESSMENTS

## 15.1 APPROVALS PROCESS

A range of approval mechanisms are available to the project. Irrespective of the major approvals process, a rigorous environmental impact assessment would be required for a project of this scale. This assessment will be targeted toward the specific natural and social values of the project area that may be impacted by an electrical transmission line, many of which were identified during development of this CSR.

It is important to record key commitments from this CSR that require specific treatment in subsequent assessment processes. Key commitments are described in Section 15.4.

Powerlink has a wealth of experience in transmission line development, impact assessment, land acquisition, stakeholder and landholder engagement to support Genex in subsequent phases, whichever process is deemed preferable.

## 15.2 STAKEHOLDER AND LANDHOLDER ENGAGEMENT

Ongoing engagement in subsequent project phases is recommended to refine the preliminary alignment and continue to develop the relationships established during development of this CSR. Ongoing engagement will ensure that potential impacts are thoroughly understood and can be effectively managed. Powerlink notes and values the contributions of stakeholders and landholders and their time and information in helping develop this CSR.

## 15.3 EPBC ACT REFERRAL

The third party ecological advice demonstrated negligible risk to Threatened Ecological Communities and the World Heritage Wet Tropics Area, which provides significant comfort around some MNES. Unfortunately database searches returned few results for MNES flora and fauna and follow-up work is required to address the gaps in available information.

Powerlink proactively undertook a likelihood of occurrence assessment for EPBC Act threatened and migratory species and NC Act EVNT flora and fauna. See Section 11.4.1.1. The likelihood of occurrence assessment related mapped vegetation types to fauna and flora species known to be associated with those features in the broader region well beyond the CSR study area, targeting more developed areas where species records are more readily available.

This additional assessment confirmed the initial ecological advice procured for the CSR and identified only a relatively small number of additional EPBC species that may occur in the study area. Importantly, no new Threatened Ecological Communities or World Heritage Values were identified through this additional assessment.

Powerlink has significant experience and relationships in managing the EPBC referral process and designing solutions to minimise potential impacts to MNES. Powerlink projects have generally not been declared controlled actions due to our ability to optimise infrastructure siting to minimise terrestrial impacts.

Based on Powerlink's experience, it is recommended that referral of the project to the DoEE for further approvals in relation to the EPBC Act should occur early in subsequent stages of the project (e.g. Draft EIS phase).

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This approach is recommended as field surveys will be necessary to provide adequate detail for DoEE assessment managers to develop sufficient understanding of any potential issues and, if necessary, develop and apply risk-appropriate controls.

Whilst every reasonable effort has been undertaken to assess potential impacts to MNES flora and fauna at the desktop level for this CSR, the broad scale nature of mapping over the largely undeveloped project area means field surveys will be important in future stages of the project to confirm any potential impacts and allow appropriate mitigation methods to be developed.

Preliminary scheduling by Powerlink indicates that sufficient time is available in Genex's schedule to refer the project and receive a decision under the EPBC Act, even if the project is declared a Controlled Action.

Importantly, field surveys and a better understanding of the presence of threatened habitats and species, may also support an application for a lower level of assessment, such as a declaration of Not a Controlled Action - Particular Manner.

The optimal approach for the project is therefore to undertake field surveys prior to referring the project to DoEE, ensuring that potential impacts to MNES flora and fauna are understood and a proposal for an appropriate level of assessment under the EPBC Act can be formulated. Powerlink will work with closely Genex to manage the assessment of MNES flora and fauna, advising on an appropriate course of action once the likelihood of occurrence assessment is available.

## 15.4 KEY COMMITMENTS & ACTIONS

A number of key commitments and matters of importance to the project have arisen from the development of this CSR. These matters are summarised in Table 10 and serve as a useful reference for subsequent impact assessment and engagement phases of the project. The matters summarised in Table 10 are project specific and are not intended to be exhaustive.

Topic/ Issue	Comments	
Hells Gate Dam	The location of the proposed Hells Gate and Mt Fullstop Dams is an important input to latter assessment phases and further investigation should be undertaken	
Planning Act 2016	The emerging Planning Act 2016 should be actively monitored for potential changes to major approvals frameworks	
Seek Detailed Landholder Input	Work closely with landholders to fully understand their properties and refine an alignment for the proposed transmission line for further more detailed investigation as part of the EIS process should the project proceed	
Resource Tenements	Consultation with mining tenement holders should occur as early as possible in subsequent assessment phases	
Mt Fox Site Selection	Undertake a targeted site selection process at Mt Fox based on higher resolution terrain information (and imagery if possible) than that currently available	
Biosecurity	Ensure a rigorous biosecurity management plan is prepared before field surveys or field activities commence	

#### Table 10 – Key Commitments and Actions

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Terrain Information	Procure higher resolution terrain information and imagery to inform project development. The ROAMES information held by Ergon may represent an opportunity to procure recent, off the shelf information for a lower capital cost for much of the preferred study corridor	
Local Providers	Commence discussions around use of local providers early in the project development process to ensure amendments to existing procurement processes can occur if required	
Defence Acquisition	Continue enquiries into the potential Defence Force acquisition of land in the Hinchinbrook/ Charters Towers Council areas (Section 13.7)	
Additional engagement	Expand engagement to include broader community groups and associations identified by stakeholders during the CSR phase (e.g. local rural fire brigade, schools etc.). These stakeholders are listed in Section 13.1.	

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## 16 REFERENCES & INFORMATION SOURCES

Bureau of Meteorology (BOM)

http://www.bom.gov.au

**Charters Towers Regional Council** 

http://www.charterstowers.qld.gov.au/

Commonwealth of Australia, Australian Bureau of Statistics

http://www.abs.gov.au

Department of Environment and Energy

http://www.environment.gov.au

Heritage Protection

https://www.ehp.qld.gov.au

Department of Natural Resources and Mines (DNRM)

http://www.nrm.qld.gov.au

Etheridge Regional Council

http://www.etheridge.qld.gov.au/

Hinchinbrook Shire Council

http://www.hinchinbrook.qld.gov.au/

Genex Power - The Kidston Solar Project

http://www.genexpower.com.au/projects/The\_Kidston\_Project

Queensland Government, Office of Economic and Statistical Research

http://statistics.oesr.qld.gov.au/qld-regional-profiles

**Queensland Spatial Catalogue - QSpatial** 

qldspatial.information.qld.gov.au

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Attachment 1 – Third Party Ecological Advice

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POWERLINK

# Initial Desktop Assessment and Preliminary Ecological Constraints Advice

PROPOSED GENEX KIDSTON CONNECTION PROJECT

CONFIDENTIAL

### DECEMBER 2016

WSP PARSONS BRINCKERHOFF

# Initial Desktop Assessment and Preliminary Ecological Constraints Advice

PROPOSED GENEX KIDSTON CONNECTION PROJECT

Powerlink

#### Confidential

REV	DATE	DETAILS
А	29/11/2016	Draft report
В	16/12/2016	Final report

#### AUTHOR, REVIEWER AND APPROVER DETAILS

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# 1 INTRODUCTION

Powerlink is investigating three potential corridor options for powerline to connect a renewable energy generator (Genex) at Kidston in Far North Queensland, to the electricity transmission grid. The site at Kidston will contain a mix of solar and pumped storage (hydro) technology with a total generation capacity of approximately 400 MW. The pumped storage component of the facility can be rapidly ramped up and down to produce energy at times of peak demand.

The closest connection point to Kidston is at Mount Fox, approximately 200 km east of Kidston. It is most likely that a 275 kV transmission line will be required to provide adequate capacity for the connection over that distance. On this basis, Powerlink has been commissioned by Genex to produce a Corridor Selection Report, to identify a preferred corridor and a preliminary alignment for the connection from a broad study area.

The assessment for the Corridor Selection Report will include engagement with Government agencies, peak bodies and landowners in the study area, as well as high level environmental, land use planning and technical investigations. It builds on a Draft Corridor Selection Report compiled for Genex in mid-2016, which identified three high level corridor options for further assessment.

The Corridor Selection Report will assess a study area based on the two southern most options from the Draft Corridor Selection Report (Options B and C), the northern most option (Option A) having been removed from assessment based on feedback during the development of the report. Part of the rationale for removal of Option A was its potential for greater environmental impacts, which was evident at the Draft Corridor Selection stage.

To validate the higher potential for environmental impact this Initial Desktop Assessment and Preliminary Ecological Constraints Advice considers all options from the Draft Corridor Selection Report. The intention is to validate the removal of the Option A, providing surety that the Corridor Selection Report is focused on the right study area and double checking the corridor refinement process undertaken to date.

This Initial Desktop Assessment and Preliminary Ecological Constraints Advice identifies the Matters of National Environmental Significance (MNES) and Matters of State Environmental Significance (MSES) that are of relevance to each corridor option. The findings herein will inform the preferred option selection process of Powerlink's Corridor Selection Report.

Since completion of this assessment, an additional area outside of Options A, B and C, is being considered. The additional area was identified during desktop investigations and allows the opportunity to avoid difficult topography and potentially seek to co-locate with the Gregory Development Road.

This additional area was identified after this report was substantially complete and has therefore not been assessed as part of this assessment. However, based upon the similarities of the MNES and MSES assessed for each alignment, it is unlikely the additional area would include anything that has not already been assessed, nor is it likely to influence the calculations for the MNES and MSES that have been assessed as occurring broadly across all three corridor options. Furthermore, the extent of ecological constraints for each Corridor option would be unlikely to increase or decrease substantially by including the additional area, and the corridor option identified as the option of least ecological constraint would be unlikely to change.

#### 1.1 Purpose

This initial desktop assessment and preliminary ecological advice identifies the matters of national and state environmental significance (MNES and MSES) that may be of potential relevance to each corridor option (A, B and C. It presents the potential ecological constraints and risks associated with each option and provides advice on a preferred option from an environmental perspective, which contains the least amount of MNES and MSES ecological constraints.

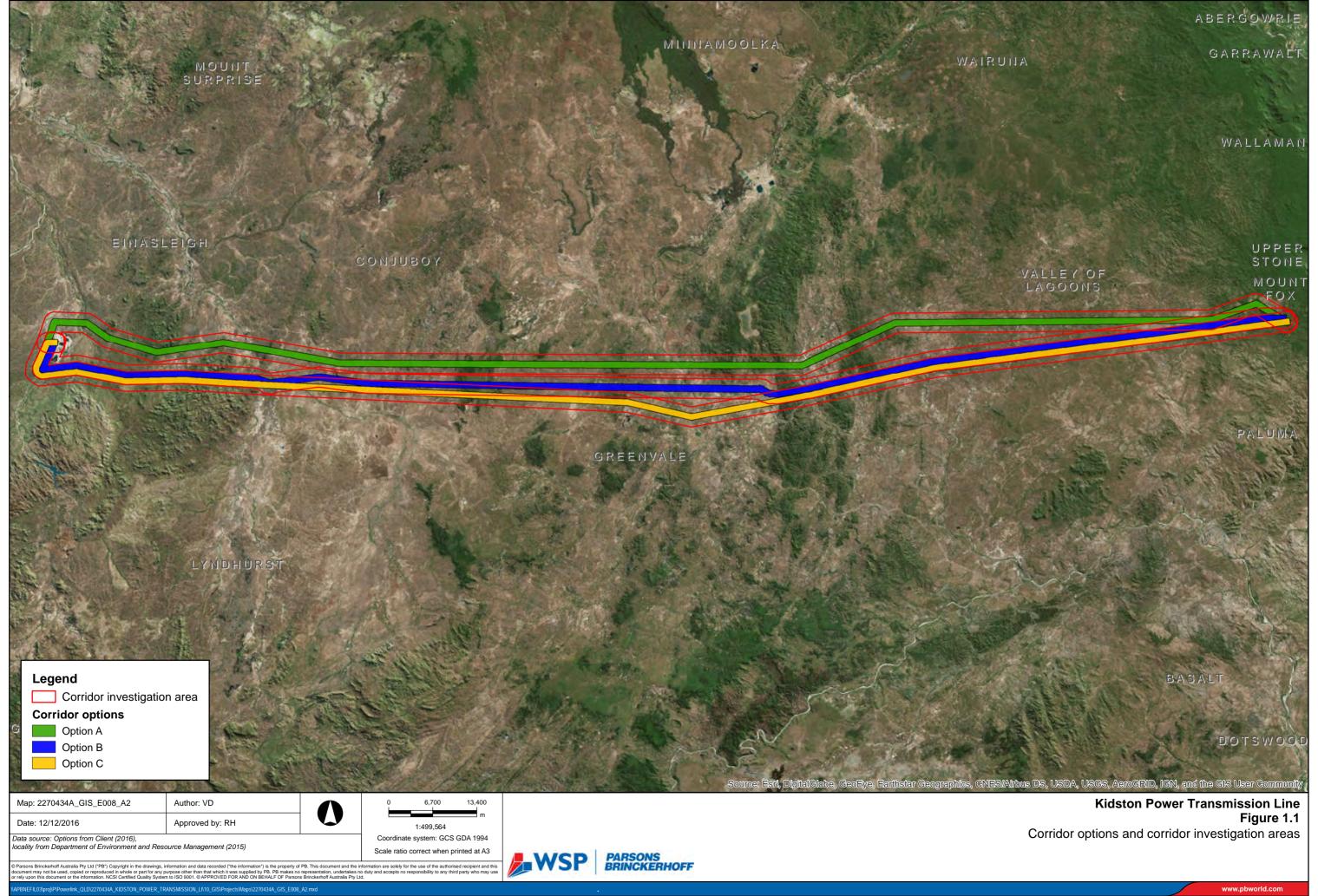
It also provides preliminary advice on the possible requirement of significant impact assessments for potential impacts to relevant MNES and MSES within each corridor, and whether the Project should be referred to the Department of the Environment and Energy (DoEE).

A brief discussion on potential Commonwealth and State environmental offset requirements has also been provided.

#### 1.1.1 Terminology

The desktop assessment discusses the corridor options and corridor investigation areas, which are defined as:

- → Corridor options three 1 km wide corridor options as defined in the Draft Corridor Selection Report, and shown on Figure 1.1.
- Corridor investigation areas a 3 km wide corridor investigation area has been applied to the centreline of each 1 km wide corridor option, to inform the Corridor Selection Report of the potential ecological constraints of relevance to each option, as shown on Figure 1.1.



# 2 DESKTOP ASSESSMENT METHODS

The initial desktop assessment has been designed to inform Powerlink of the MNES and MSES ecological constraints that are of relevance to each of the three corridor options. Powerlink has provided the three corridor options (A, B and C). Each option is 1 km wide. This initial desktop assessment applies database searches and GIS analysis to 3 km wide corridor investigation areas for each corridor option. The main purpose of the assessment is to determine the option of least ecological constraint for the Project. The methods used for the initial desktop assessment have been discussed in more detail in the following Sections 2.1–2.5.

#### 2.1 Database searches

The initial desktop assessment included searches of the following databases:

- → Commonwealth Department of the Environment and Energy's (DoEE) Protected Matters Search Tool (PMST), with a 5 km buffer applied to each 1 km wide corridor option, as indicated on Figure 2.1.
- State Department of Science, Information Technology and Innovation's (DSITI) Wildlife Online, by applying 10 point searches (18 km radius) along a shared centreline of the three options, as indicated in Figure 2.2.

#### 2.2 Data collation and record searches

The data collation process and threatened species records searches for determining the threatened species and/or ecological communities, listed under the EPBC Act and/or NC Act, that are of relevance to each corridor option are discussed in Sections 2.2.1–2.2.2.

#### 2.2.1 Data collation

The data collation process identifies the threatened and migratory species listed under the EPBC Act and/or NC Act for which species record searches are undertaken.

#### 2.2.1.1 PMST

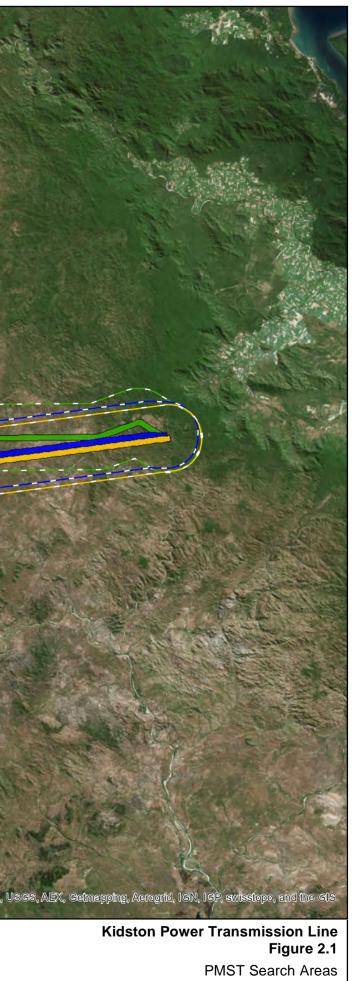
The PMST database results reveals the MNES of potential relevance to each corridor option. The PMST searches are presented in Appendix A. The PMST search results for each corridor option have been collated into a spreadsheet to present the results of relevance to each corridor option. This spreadsheet is available in Appendix B.

#### 2.2.1.2 Wildlife Online

The 10 Wildlife Online database search results were obtained as Excel files so they could be amended to draw attention to the threatened, special least concern and migratory species listed under the NC Act and/or EPBC Act. These amended spreadsheets are presented in Appendix C.

The Wildlife Online database results were taken from 10 centre point searches with an 18 km radius along a shared centreline of the three options, as shown on Figure 2.2. Wildlife Online database is based upon species records. The results of the 10 point searches (18 km radius) were collated into a spreadsheet to identify the endangered, vulnerable, near threatened and special least concern species that have records within the search area. This spreadsheet is available in Appendix D.

Legend Skm buffer around options		
Option C         GenexOptions         Option A1         Option B1         Option C         Map: 2270434A_GIS_002_A2         Author: VD         Date: 11/11/2016         Approved by: RH         Data source: Options from Client (2016)	IN BEHALP OF Paisons Britckenton Australia Pty Lto.	eographics, CNES/Airbus DS, USD



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#### 2.2.2 Threatened species records searches

The next step of the process involved undertaking species records searches for each threatened species returned from the combined PMST and Wildlife Online search results. The collated PMST and Wildlife Online search results spreadsheet that was developed to inform species records searches, as presented in Appendix E. The following two databases have been searched for species records:

- 1. State DSITI Species profile search to obtain state wide records for the threatened flora and fauna species listed under the NC Act and/or EPBC Act.
- 2. Commonwealth Government administered Atlas of Living Australia to obtain national records for the threatened fauna species listed under the NC Act and/or EPBC Act.

It is important to note that for some NC Act listed threatened flora and fauna species (e.g. orchids and frogs) are not made available by DSITI, due to the risk of them being illegally captured or removed from the environment.

Once all of the available records were obtained, they were put through the GIS analysis and mapping process.

#### 2.3 GIS analysis and mapping

Using ESRI ArcGIS, a 1 km buffer was applied to the centreline of each 1 km wide corridor option (A, B and C) to create a 3 km wide investigation area. Each 3 km wide investigation area (corridor investigation area) is the basis for quantifying and mapping the MNES and MSES ecological constraints therein. This was done to draw comparisons between each corridor option to inform the option selection process.

#### 2.3.1 Quantification of ecological constraints

The number and/ or extent (hectares) of ecological constraints, within each 3 km wide corridor investigation area (Options A, B and C), has been quantified and mapped using the GIS intersect tool for the following mapping layers:

- → The **number** of DSITI's species records for threatened flora and fauna species listed under the NC Act and/or EPBC Act that have been previously recorded within each corridor investigation area.
- → The **number** of Atlas of Living Australia's species records for threatened fauna listed under the NC Act and/or EPBC Act that have been previously recorded within each corridor investigation area.
- The presence or absence of 'high risk areas' for endangered, vulnerable and near threatened plants (EVNT plants) listed under NC Act on the 'protected plants flora survey trigger map' within each corridor investigation area.
- → The extent (hectares) of MSES regulated vegetation in reference to:
  - Department of Natural Resources and Mine's (DNRM) regulated vegetation mapping (endangered and of concern regional ecosystems)
  - DNRM's regional ecosystem mapping (endangered, of concern and least concern regional ecosystems)
  - Virtual GIS's remote sensing analysis (refer Section 3.1.4) and mapping of woody vegetation predicted and potential extent of remnant, regrowth and non-remnant vegetation
  - regulated remnant watercourse vegetation (endangered, of concern and least concern regional ecosystems), methodology provided below in Section 2.3.1.1
  - regulated remnant wetland vegetation (endangered, of concern and least concern regional ecosystems), methodology provided below in Section 2.3.1.2.
- → The extent (hectares) of other MSES listed under Schedule 2 of the Environmental Offsets Regulation 2014 within each corridor investigation area, including:
  - protected areas (e.g. National Parks and State reserves)
  - wildlife habitats
  - strategic environmental areas

- high ecological significance wetlands
- high ecological value waters (wetland)
- high ecological value waters (watercourses)
- environmental offsets
- connectivity.

#### 2.3.1.1 Regulated remnant watercourse vegetation

The 1:100,000 scale regulated vegetation mapping covers approximately two thirds of each corridor investigation area, while the remaining thirds, toward the western extent of each corridor investigation area, are covered by the 1:250,000 scale mapping.

The 1:250,000 scale mapping does not have stream order attributes. Therefore, the Strahler method has been applied to the 1:250,000 watercourse layer using GIS to appropriately assign the stream orders.

The defined distance for a remnant regional ecosystem along a watercourse has been determined on whether the area of investigation is located in a coastal bioregion and sub-regions or non-coastal bioregion and sub-regions, which are prescribed under Schedule 2 clause 2, subsections (4) and (6) of the Environmental Offsets Regulation 2014, including:

- → Coastal bioregions and sub-regions:
  - Townsville Plains (11.1), Bogie River Hills (11.2) and Marlborough Plains (11.14) Subregions, Brigalow Belt (SBRB) Bioregion
  - Central Queensland Coast (CQC) Bioregion
  - Starke Coastal Lowlands (3.2) Subregion, Cape York Peninsula (CYP) Bioregion
  - Hodgkinson Basin (9.3) Subregion, Einasleigh Uplands (EIU) Bioregion
  - Wet Tropics (WET) Bioregion
  - South East Queensland (SEQ) Bioregion.
- → Non-coastal bioregions and sub-regions:
  - Brigalow Belt (SBRB) Bioregion (excluding Subregions 11.1, 11.2 and 11.14)
  - New England Tableland (NET) Bioregion
  - Northwest Highlands (NWH) Bioregion
  - Gulf Plains (GUP) Bioregion
  - Cape York Peninsula (CYP) Bioregion (excluding Subregion 3.2)
  - Mitchell Grass Downs (MGD) Bioregion
  - Channel Country (CHC) Bioregion
  - Mulga Lands (MUL) Bioregion
  - Einasleigh Uplands (EIU) Bioregion (excluding Subregion 9.3)
  - Desert Uplands (DEU) Bioregion.

The distances for regulated remnant watercourse vegetation from the defining bank of watercourse stream orders in coastal bioregions and sub-regions, is presented in Table 2.1.

## Table 2.1Distances for regulated remnant watercourse vegetation from the defining bank of watercourse<br/>stream orders in coastal bioregions

WATERCOURSE STREAM ORDER	DISTANCE FROM THE DEFINING BANK (metres)
1 or 2	10
3 or 4	25
5 or greater	50

The distances for regulated remnant watercourse vegetation from the defining bank of watercourse stream orders in non-coastal bioregions and sub-regions, is presented in Table 2.2.

Table 2.2Distances for regulated remnant watercourse vegetation from the defining bank of watercourse<br/>stream orders in non-coastal bioregions

WATERCOURSE STREAM ORDER	DISTANCE FROM THE DEFINING BANK (metres)
1 or 2	25
3 or 4	50
5 or greater	100

#### 2.3.1.2 Regulated remnant wetland vegetation

The extent (ha) of remnant vegetation (endangered, of concern and least concern regional ecosystems) of relevance to areas of mapped regulated wetland vegetation, is determined by applying a 50 m buffer to the mapped regulated vegetation wetlands.

#### 2.4 Threatened ecological communities assessment

An assessment of the mapped regional ecosystems within each corridor was undertaken to identify the potential presence of regional ecosystems that may potentially constitute threatened ecological communities listed under the EPBC Act.

#### 2.5 Constraint ranking

A ranking system has been applied to each corridor investigation area, for each environmental matter and/or potential ecological constraint, to capture the degree of ecological constraints within each corridor option and to assist with determining the least constrained or preferred corridor.

RED	Greatest amount of potential impact and ecological constraint
ORANGE	Moderate amount of potential impact and ecological constraint
GREEN	Least amount of potential impact and ecological constraint

The ranking system has not been applied to database search results (i.e. Wildlife Online and PMST) because they have been used to inform species record searches and assessment. The ranking system has also not been applied to matters that do not require significant impact assessments or trigger environmental offsets (i.e. least concern regional ecosystems and regrowth and areas of non-remnant).

The 'high risk areas' for endangered, vulnerable and near threatened plants (EVNT plants) listed under NC Act on the 'protected plants flora survey trigger map', have also not been ranked as they are captured as flora species records.



# PROPOSED GENEX KIDSTON CONNECTION PROJECT

# **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"

# **3** ECOLOGICAL CONSTRAINTS

#### 3.1 Matters of State Environmental Significance

The following MSES have been assessed in the following sections:

- → NC Act threatened flora and fauna species
- → regulated vegetation, including:
  - DNRM's regulated vegetation mapping
  - DNRM's regional ecosystem mapping
  - Virtual GIS's remote sensing analysis and mapping of woody vegetation
  - regulated remnant watercourse vegetation
  - regulated remnant wetland vegetation
- → protected areas (e.g. National Parks and State reserves)
- wildlife habitats
- → strategic environmental areas
- → high ecological significance (HES) wetlands
- → high ecological value (HEV) waters (wetland)
- → high ecological value (HEV) waters (watercourses)
- → environmental offsets
- connectivity.

#### 3.1.1 NC Act threatened species

#### 3.1.1.1 Wildlife Online results

The Wildlife Online database search results revealed that 26 threatened species listed under the NC Act and/or EPBC Act have been previously recorded within the search area that was applied to all three corridor options, including 11 birds, six mammals and nine plants.

The results of the Wildlife Online database search is presented in Table 3.1. The complete Wildlife Online database searches are presented in Appendix C.

Threatened species listed as endangered, vulnerable and special least concern under the NC Act are recognised as MSES under the *Environmental Offset Act 2014*. Near threatened species do not need to be assessed in terms of significant residual impacts under the *Environmental Offset Act 2014*, but have been included, just in case of any future changes in legislation that may occur during the approval pathway of the Project.

### Table 3.1 Wildlife Online database search results for threatened flora and fauna species previously recorded within the search area

SCIENTIFIC NAME	COMMON NAME	NC ACT	EPBC ACT
Birds			
Calidris ruficollis	Red-necked Stint	SL	М
Calyptorhynchus lathami erebus	Glossy Black-cockatoo (northern)	V	-
<i>Casuarius casuarius johnsonii</i> (southern population)	Southern Cassowary (southern population)	E	E
Cuculus optatus	Oriental Cuckoo	SL	М
Erythrura gouldiae	Gouldian Finch	E	E
Hirundapus caudacutus	White-throated Needletail	SL	М

SCIENTIFIC NAME	COMMON NAME	NC ACT	EPBC ACT
Hydroprogne caspia	Caspian Tern	SL	М
Monarcha melanopsis	Black-faced Monarch	SL	М
Poephila cincta cincta	Black-throated Finch (white-rumped subspecies)	E	E
Rhipidura rufifrons	Rufous Fantail	SL	М
Symposiachrus trivirgatus	Spectacled Monarch	SL	М
Mammals			
Mesembriomys gouldii	Black-footed Tree-rat	_	V
Petauroides volans minor	Northern Greater Glider	_	V
Petaurus gracilis	Mahogany Glider	E	E
Petrogale sharmani	Sharman's Rock-wallaby	V	V
Phascolarctos cinereus	Koala	V	V
Tachyglossus aculeatus	Short-beaked Echidna	SL	-
Plants			
Acacia tingoorensis	-	V	-
Arytera dictyoneura	-	NT	-
Cycas cairnsiana	-	V	V
Corybas cerasinus	-	NT	-
Corymbia leptoloma	-	V	V
Glossocardia orthochaeta	Yellow jacket	E	_
Leptospermum pallidum	-	NT	-
Lepturus minutus	-	V	-
Oenanthe javanica	-	NT	-

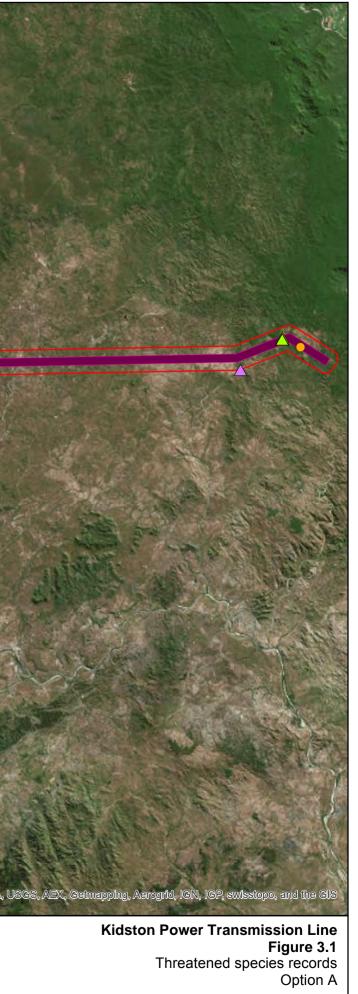
Key: E = endangered, V = vulnerable, NT = near threatened, SL = special least concern, M = migratory

#### 3.1.1.2 Threatened species records

The NC Act threatened species records, obtained from DSITI's species profile search and Atlas of Living Australia that are of relevance to each corridor option have been quantified and mapped using GIS. The number of threatened species listed under the NC Act that have been previously recorded within each corridor investigation area is presented in Table 3.2 and mapped on Figure 3.1–Figure 3.3.

The spreadsheet that contains the analysis of threatened species records is presented in Appendix F.

Legend         ● Greater Glider (EPBC Act - vulnerable)         ● Southern Black-throated Finch (EPBC Act - endangered) (NC Act - endangered)         ▲ Acacia tingoorensis (NC Act - vulnerable)         ▲ Dichanthium setosum (EPBC Act - vulnerable)         ■ Option A	Source: Esrí, DigitalGlobe, GeoEye, Earthster Geographias, CNES/Althous DS, USDA
Map: 2270434A_GIS_E003_A3a       Author: VD       0       6,300       12,600         Date: 12/12/2016       Approved by: RH       0       6,300       12,600         Data source: Species records from DSITI © The State of Queensland (2016) and Atlas of Living Australia (2016)       0       6,300       12,600         © Parsons Brinckerhoff Australia Pty Ltd (PB*) Copyright in the drawings, information and data recorded ("the information") is the property of PB. This document and the information are solely for the use of the authorised recipient and document may not be used, copied or reproduced in whole or part for any purpose other than that which it was supplied by PB. PB makes no representation, undertakes no duty and accepts no responsibility to any third party who may or rely upon this document or the information. NCSI Certified Quality System to ISO 9001. © APPROVED FOR AND ON BEHALF OF Parsons Brinckerhoff Australia Pty Ltd.         WAPBNEFIL03/projIPPowerlink_OLDI2270434A_KIDSTON_POWER_TRANSMISSION_L110.GISIProjects/Mapsi/2270434A_GIS_E003_A3.mxd	BRINCKERHOFF



#### Legend

- Curlew Sandpiper (EPBC Act critically endangered) (NC Act special least concern)
- Greater Glider (EPBC Act vulnerable)
- Koala (EPBC Act vulnerable) (NC Act vulnerable)
- Southern Black-throated Finch (EPBC Act endangered) (NC Act endangered)
- *Dichanthium setosum* (EPBC Act vulnerable)
- White-throated Needletail (EPBC Act migratory) (NC Act special least concern)
   Corridor investigation area
- Option B

		C. I DO HE CORE		1	A CONSTRUCT
Map: 2270434A_GIS_E003_A3b	Author: VD		0	6,300	12,600
Date: 12/12/2016	Approved by: RH	U		1:470,000	m
Data source: Species records from DSITI © The S	tate of Queensland (2016) and Atlas of Livi	ng Australia (2016)	Coordinate system	em: GDA 1994	MGA Zone 55
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#### Legend

- Curlew Sandpiper (EPBC Act critically endangered) (NC Act special least concern)
- Koala (EPBC Act vulnerable) (NC Act vulnerable)
- Sharman's Rock-wallaby (EPBC Act vulnerable) (NC Act vulnerable)
- Southern Black-throated Finch (EPBC Act endangered) (NC Act endangered)
- *Dichanthium setosum* (EPBC Act vulnerable)
- Corridor investigation area
- Option C

		C 10 10 10 10 10	100 M	1	
Map: 2270434A_GIS_E003_A3c	Author: VD		0	6,300	12,600
Date: 12/12/2016	Approved by: RH	U		1:470,000	m
Data source: Species records from DSITI $old c$ The S	State of Queensland (2016) and Atlas of Livi	ng Australia (2016)	Coordinate syst	em: GDA 1994	MGA Zone 55
			Scale ratio o	correct when pr	inted at A3
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### Table 3.2 The number of NC Act listed species that have been previously recorded within each corridor investigation area

NC ACT STATUS	OPTION A	OPTION B	<b>OPTION C</b>
Threatened flora species			
Endangered	-	-	-
Vulnerable	1	-	-
Near threatened	-	-	-
Threatened fauna species			
Endangered	1	1	1
Vulnerable	-	1	3
Near threatened	-	-	-
Special least concern	-	2	1
Total records	2	4	5

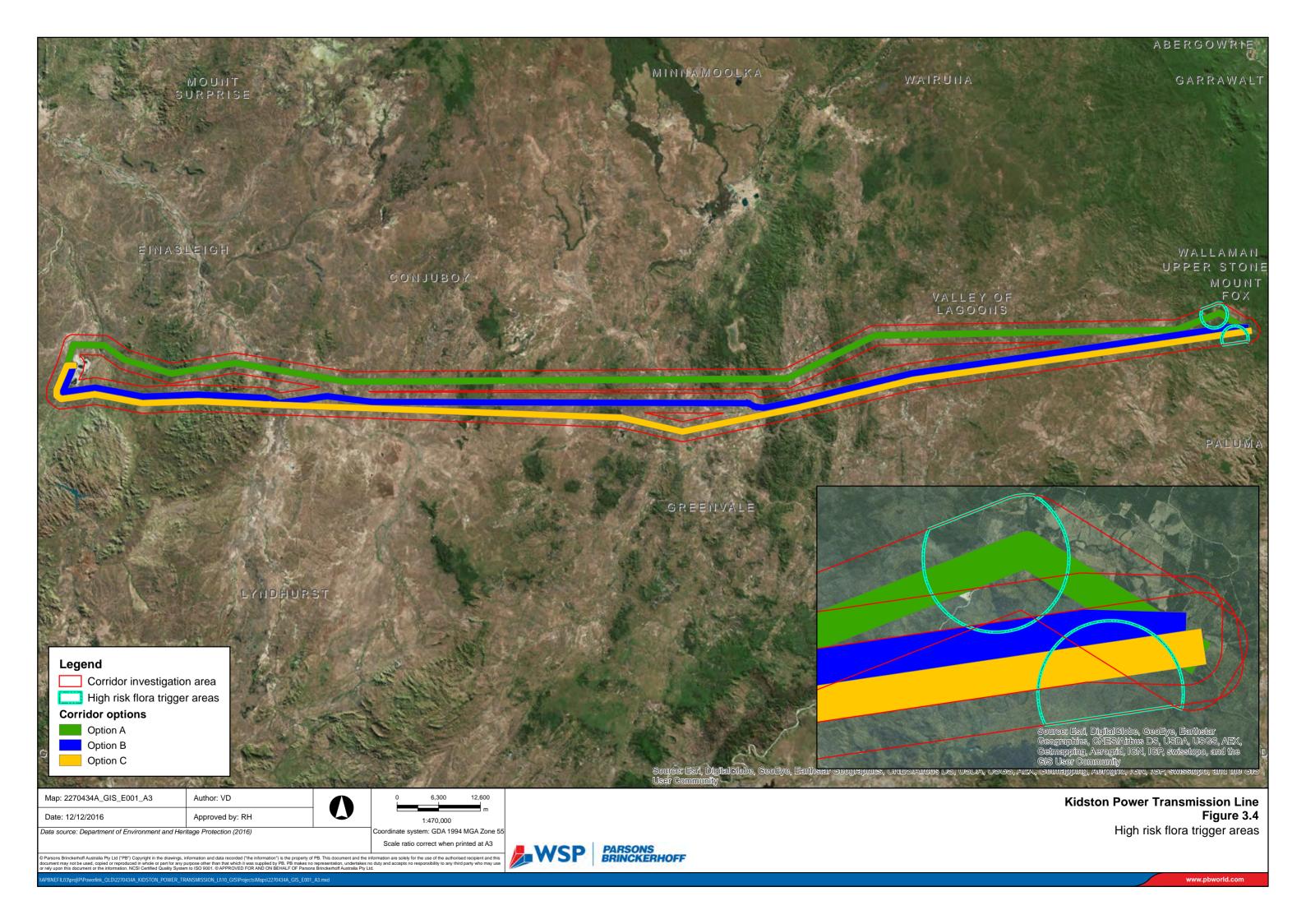
In reference to Table 3.2, Option A has the least number of NC Act threatened species records and is therefore the least constrained.

It is important to note that the number of threatened species records are representative of the number of flora and fauna surveys that are undertaken in specific areas for developments or in association with protected areas. In respect to the remoteness of the corridor investigation areas, it is possible that flora and fauna surveys for the Project may identify threatened species that may have not yet been recorded within the corridor investigation areas, and as such formerly captured on the DSITI and ALA species records databases.

#### 3.1.1.3 Protected plants flora survey trigger map high risk areas

The 'high risk areas' for endangered, vulnerable and near threatened plants (EVNT plants) listed under NC Act on the 'protected plants flora survey trigger map' within each corridor investigation area is presented on Figure 3.4. This indicates the presence of threatened flora species within the high risk areas and indicates a potential ecological constraint to all three corridor options. The high risk areas have been mapped for confirmed records of *Acacia tingoorensis*.

Once a final project footprint is obtained a protected plants flora survey in accordance with the *Flora Survey Guidelines – Protected Plants Nature Conservation Act 1992* (DEHP, 2014), would need to be conducted within the high risk areas that intersect the project footprint. This survey would need to be conducted within 12 months of construction within the high risk areas.



#### 3.1.2 Mapped regulated vegetation

The extent of DNRM's mapped regulated vegetation that is of potential of relevance to each corridor investigation area, is presented on Figure 3.5–Figure 3.19.

The area of mapped regulated vegetation that is of potential relevance to each corridor option has been further calculated using GIS, and is listed in Table 3.3.

The area (ha) of mapped regulated vegetation associated with dominant and sub-dominant of concern regional ecosystems that is of potential relevance to each corridor investigation area is presented in Table 3.3. The analysis of regulated vegetation for each corridor option is presented in Appendix G.

Based on the results of this assessment, Option C is the least constrained in terms of potential impacts upon mapped regulated vegetation, as presented in Table 3.3.

Table 3.3	Extent of regulated vegetation of relevance to each corridor option
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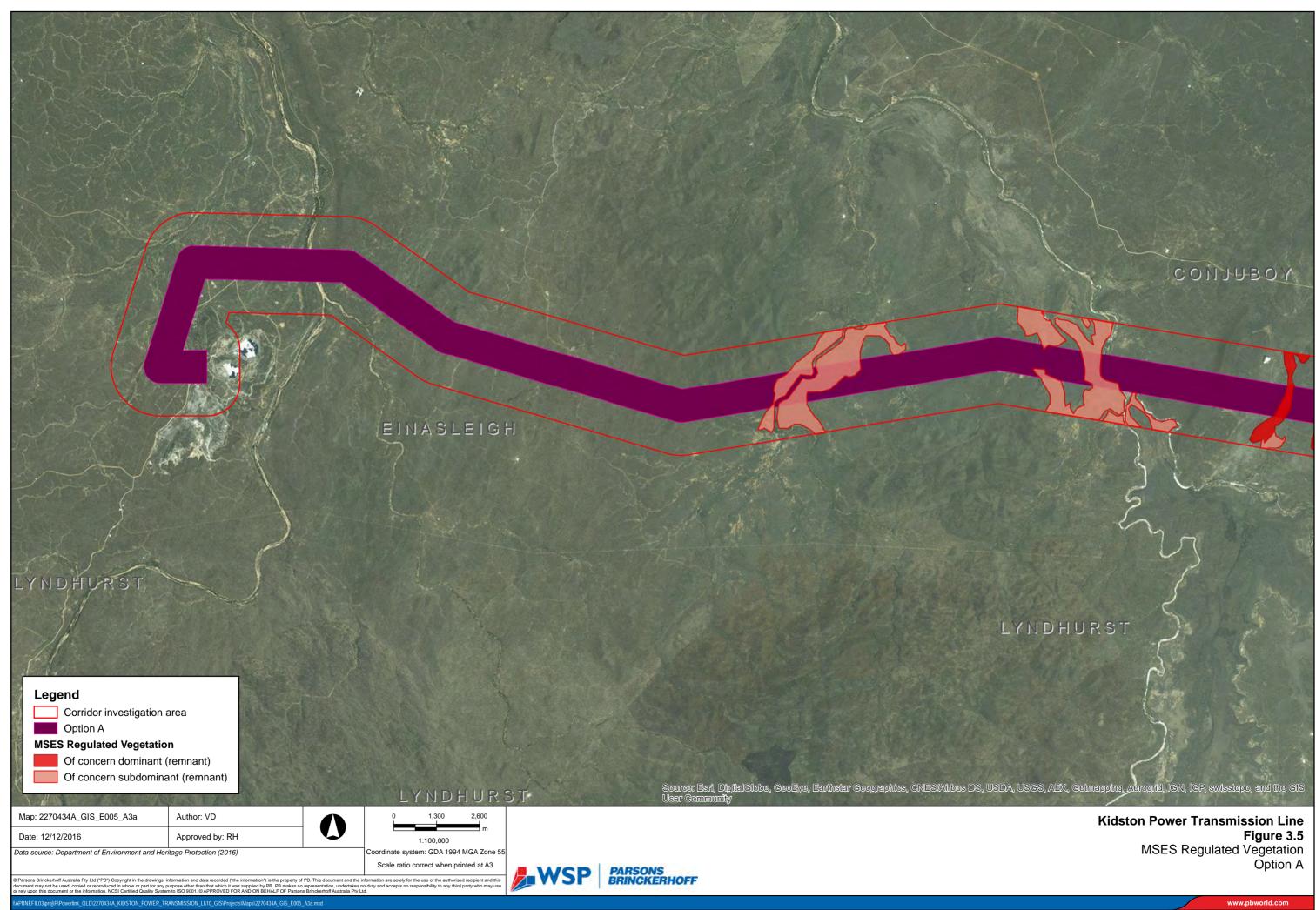
CORRIDOR	MAPPED REGULATED VEGETATION (HA)			
	OF CONCERN DOMINANT	OF CONCERN SUB- DOMINANT	OF CONCERN HVR	CATEGORY R REGROWTH
Option A	2,059	1,008	3.6	422
Option B	1,819	442	2.4	296
Option C	1,519	334	2.4	257

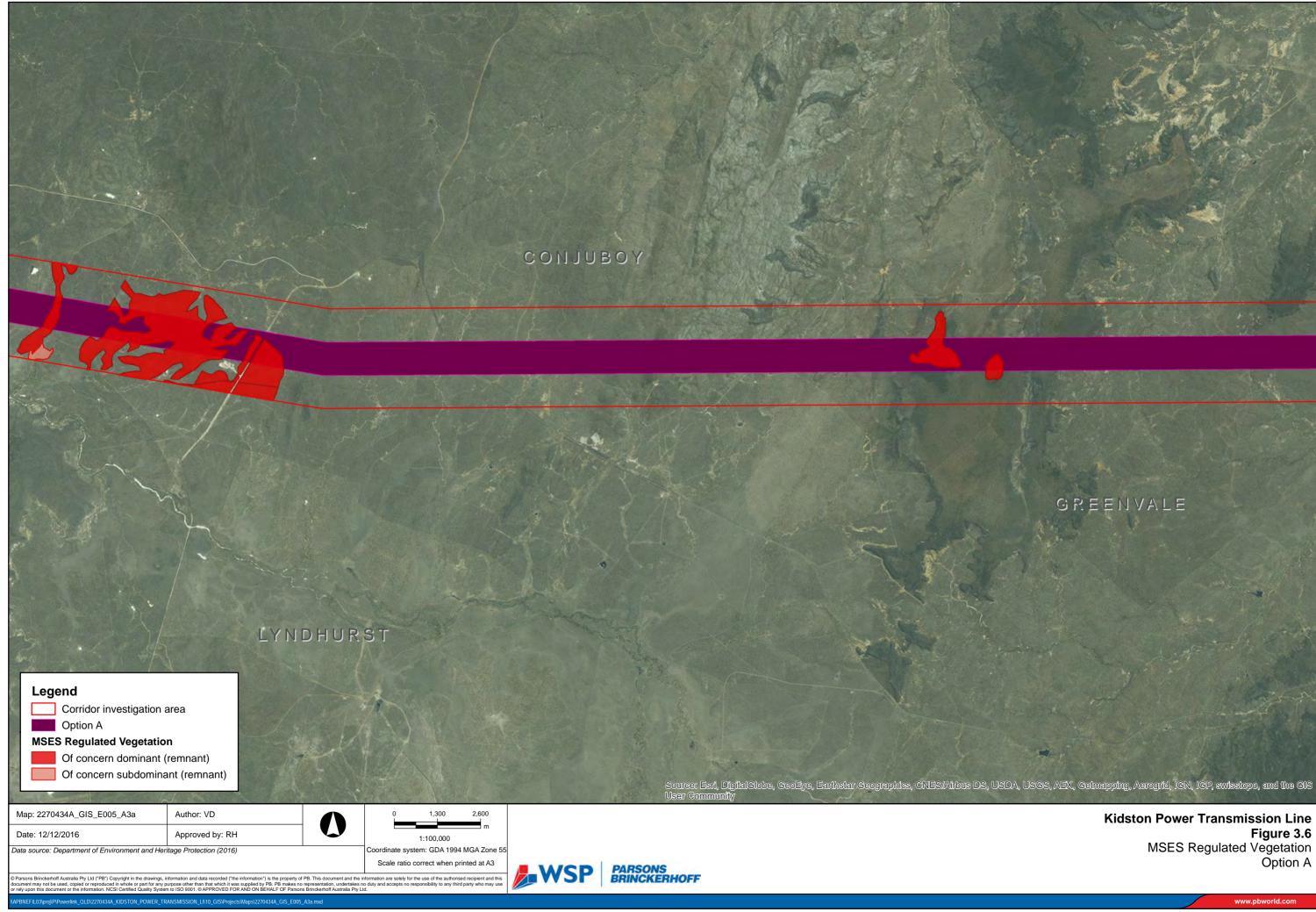
#### 3.1.2.1 Category R vegetation

The DNRM's mapped Category R regrowth vegetation: A self-assessable vegetation clearing code (DNRM, 2013), specifies that Category R regrowth vegetation is native woody vegetation on freehold land, Indigenous land or leasehold land granted for agriculture or grazing purposes, located within 50 metres of a watercourse in the Burdekin, Mackay, Whitsunday and Wet Tropics Great Barrier Reef catchments (DNRM, 2013).

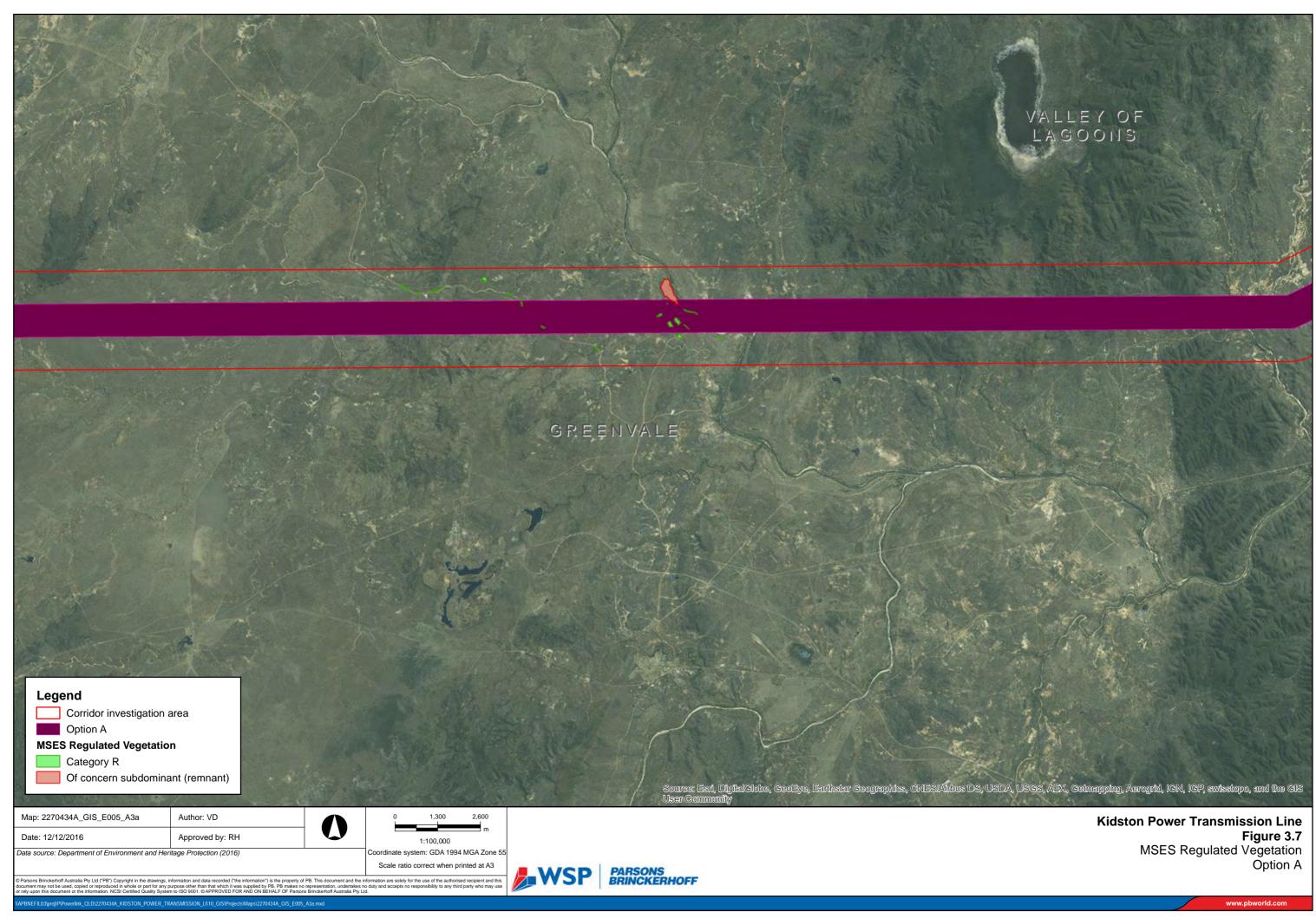
The need to consider Category R regrowth vegetation will be dependent upon the approval pathway for the Project. Assessment of the approvals pathway for the Project is excluded from the scope of this report, however a review of Community Infrastructure Designations (CIDs) database (Department of Infrastructure, Local Government and Planning, 2015) indicates that most Powerlink transmission lines are assessed and approved by the State under a CID under Chapter 5 of the *Sustainable Planning Act 2009*. Schedule 24 of the Sustainable Planning Regulation 2009 (SP Regulation) enables clearing of vegetation for community infrastructure mentioned in schedule 2 (of the SP Regulation), if the clearing is carried out on designated land.

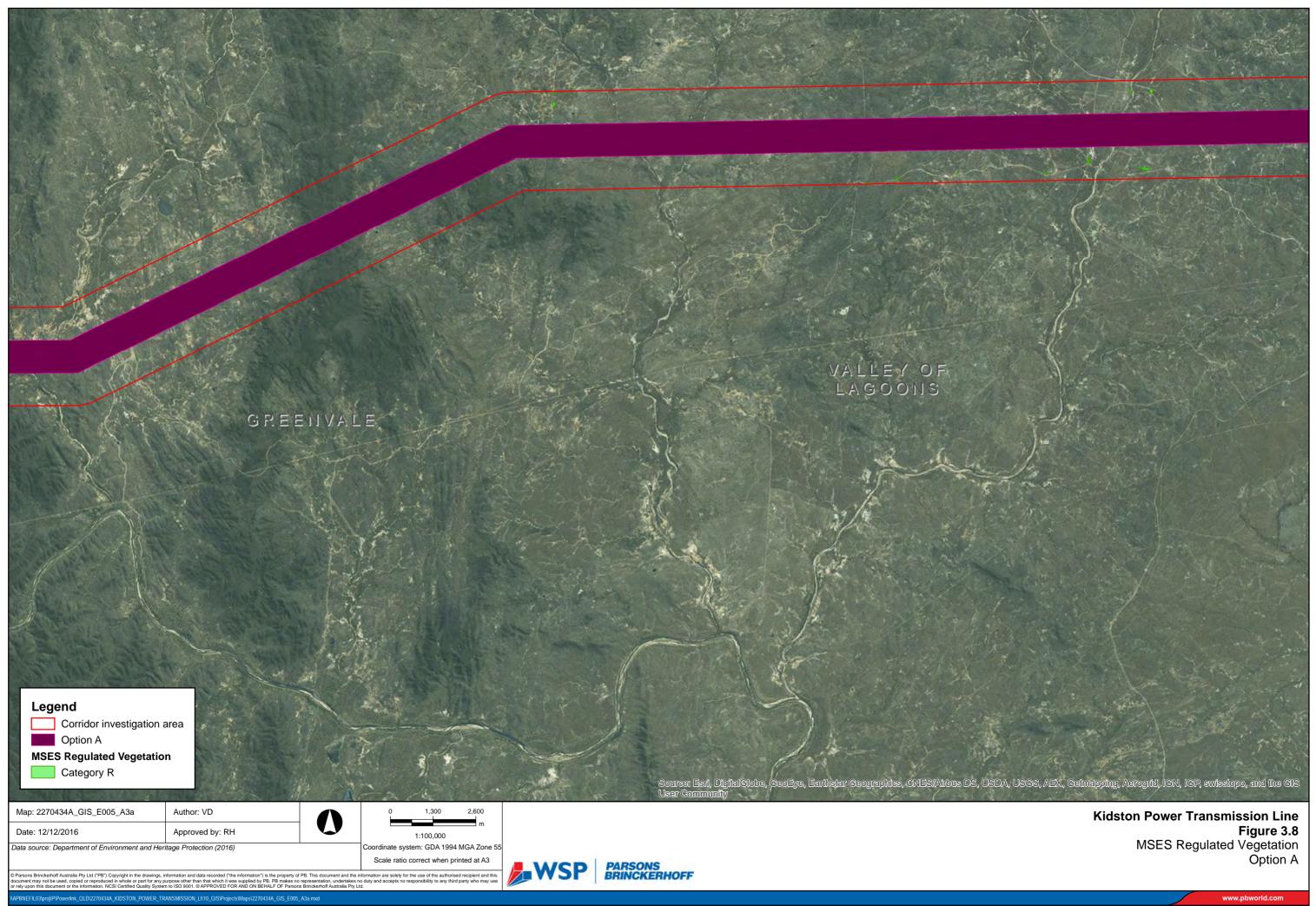
Given the likelihood that approval for the transmission line is required by the State under the CID process, it is relevant to consider the potential ecological constraints associated with Category R regrowth vegetation until the approval pathway has been determined. In reference to Table 3.3, Option C is of least constraint in terms of potential impacts upon Category R regrowth vegetation.



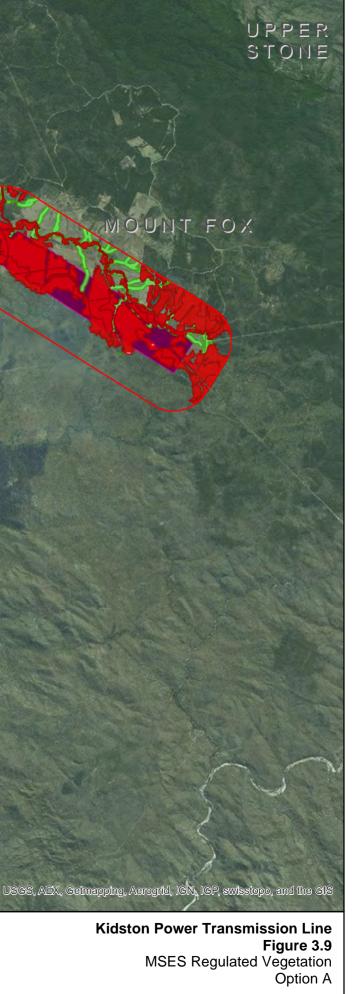


Kidston Power Transmission Line Figure 3.6 MSES Regulated Vegetation Option A

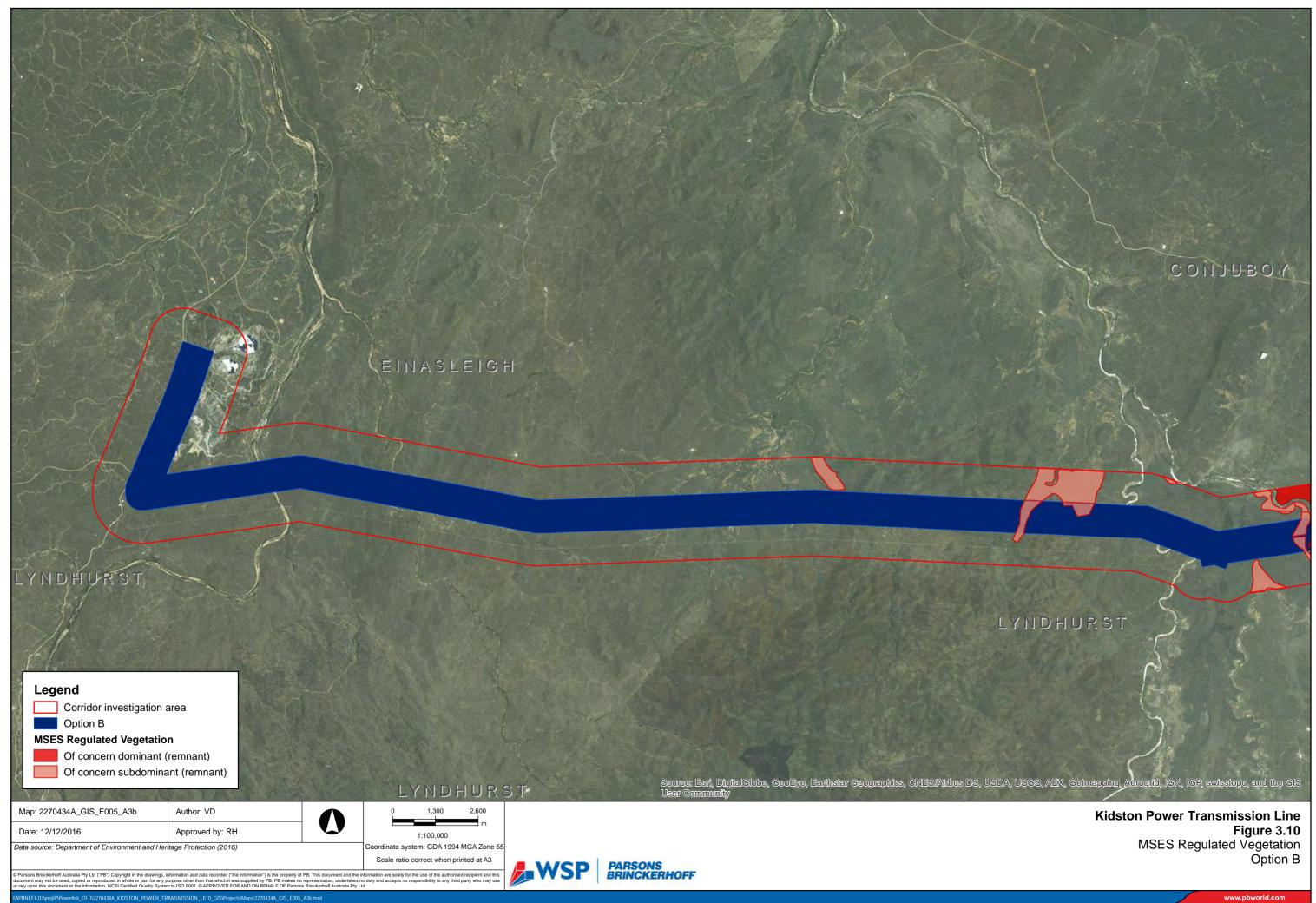


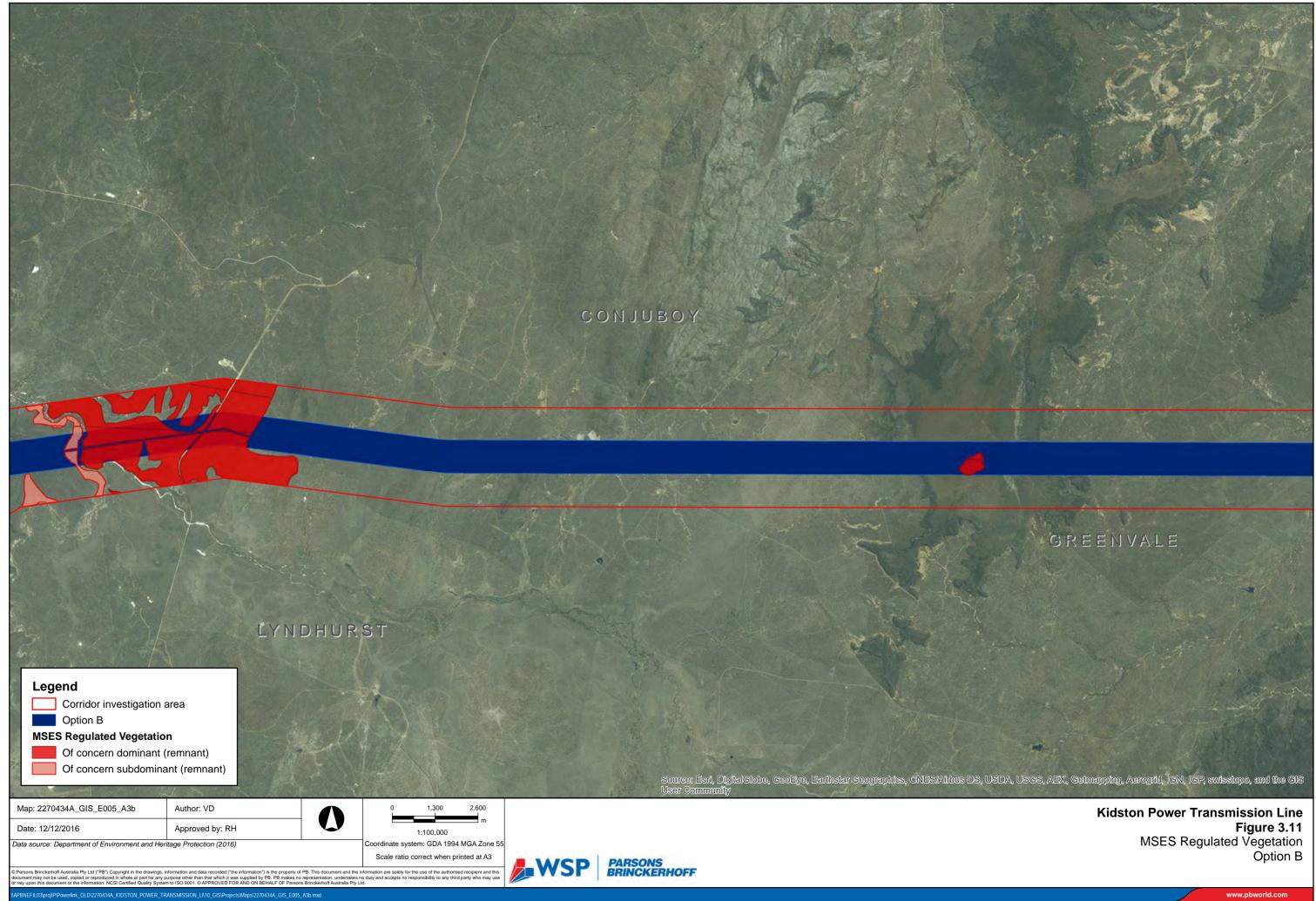


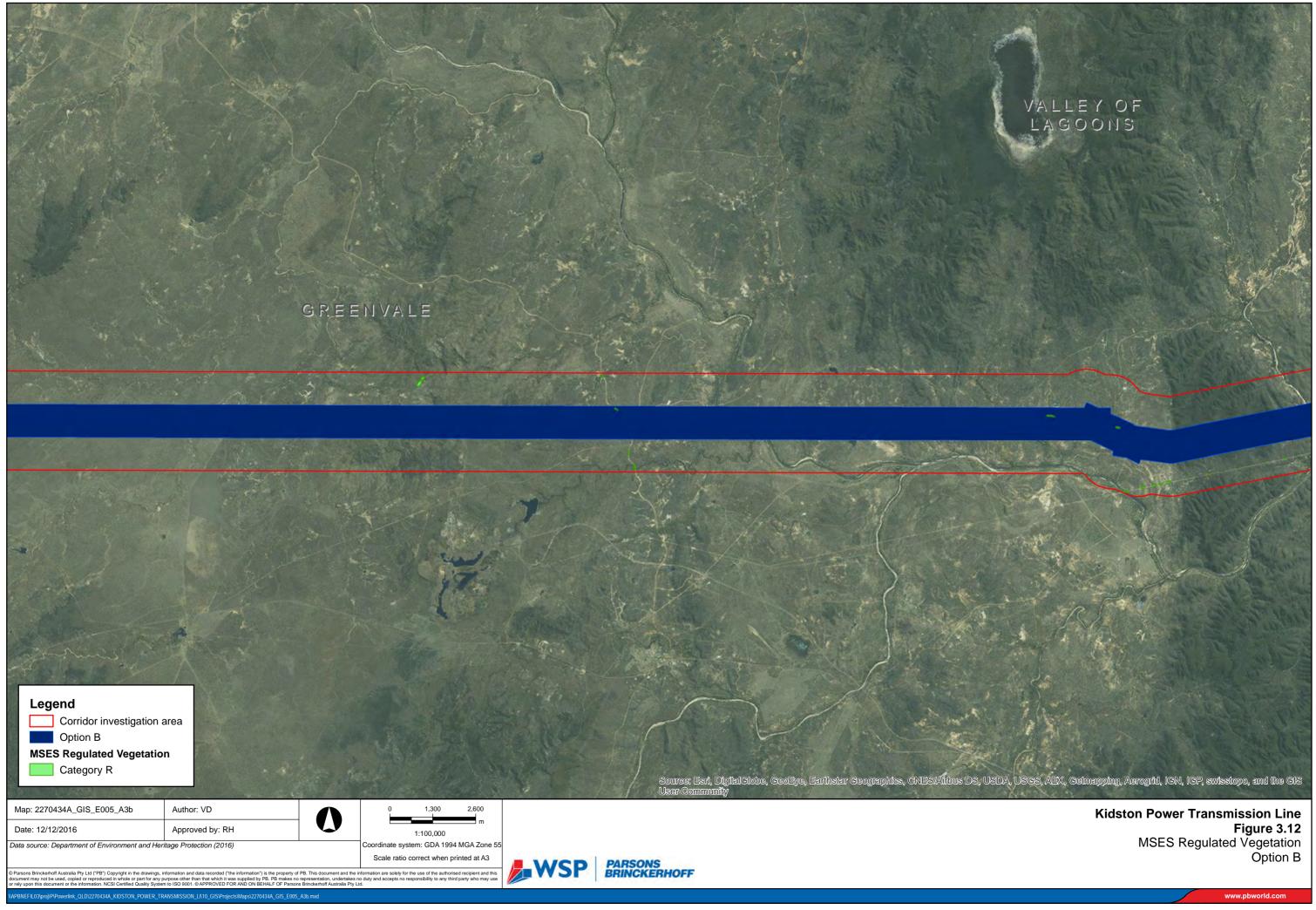
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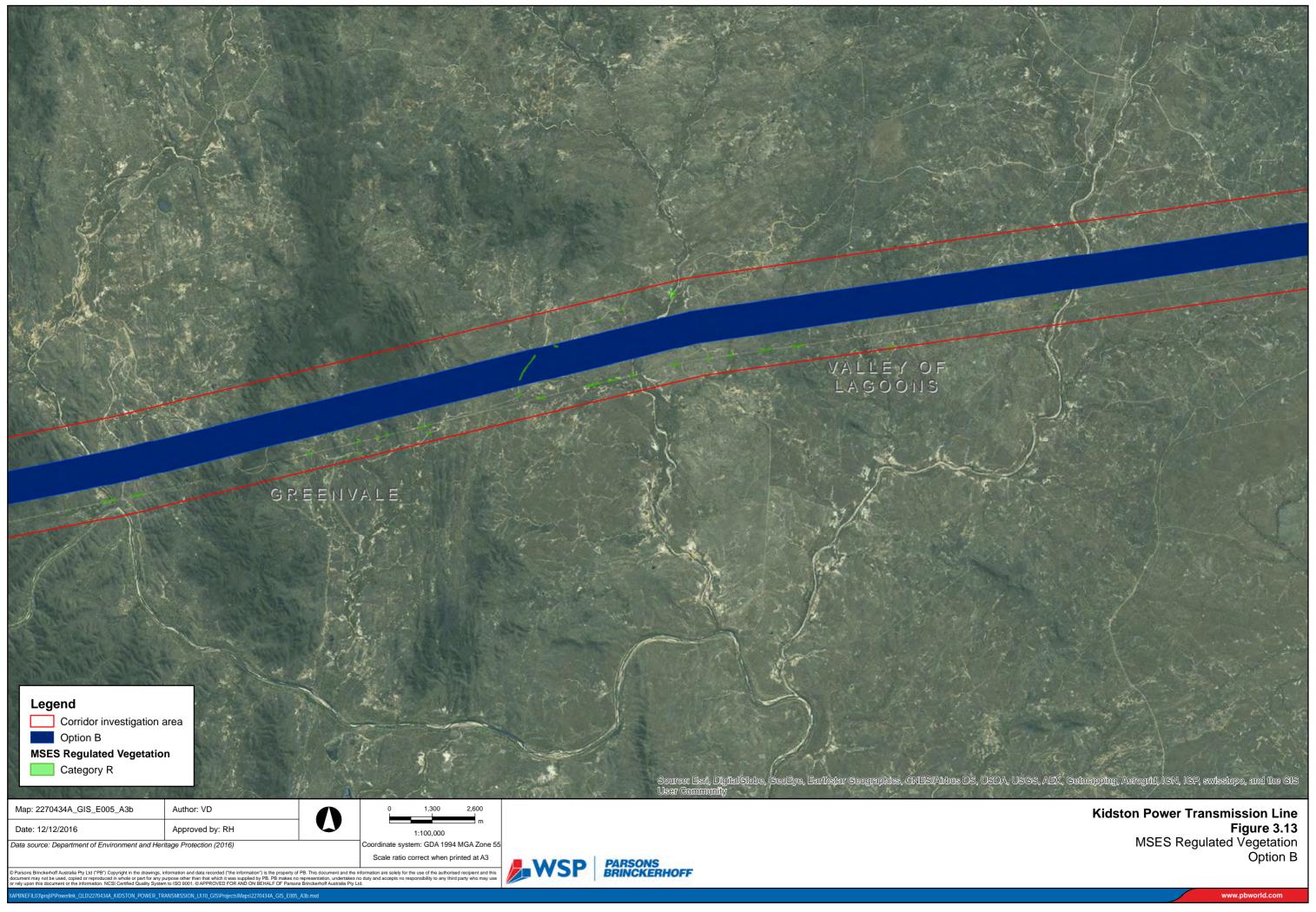


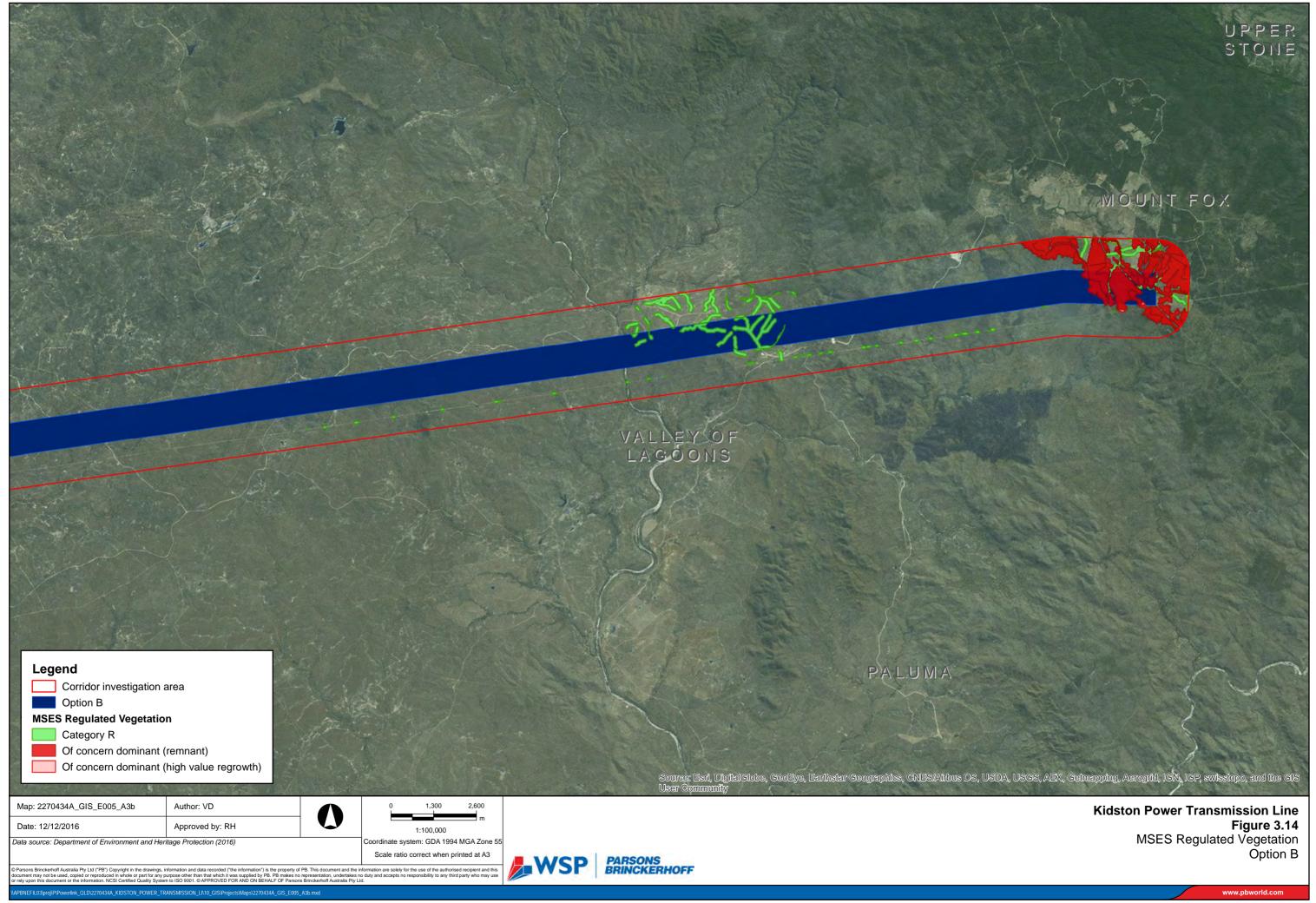
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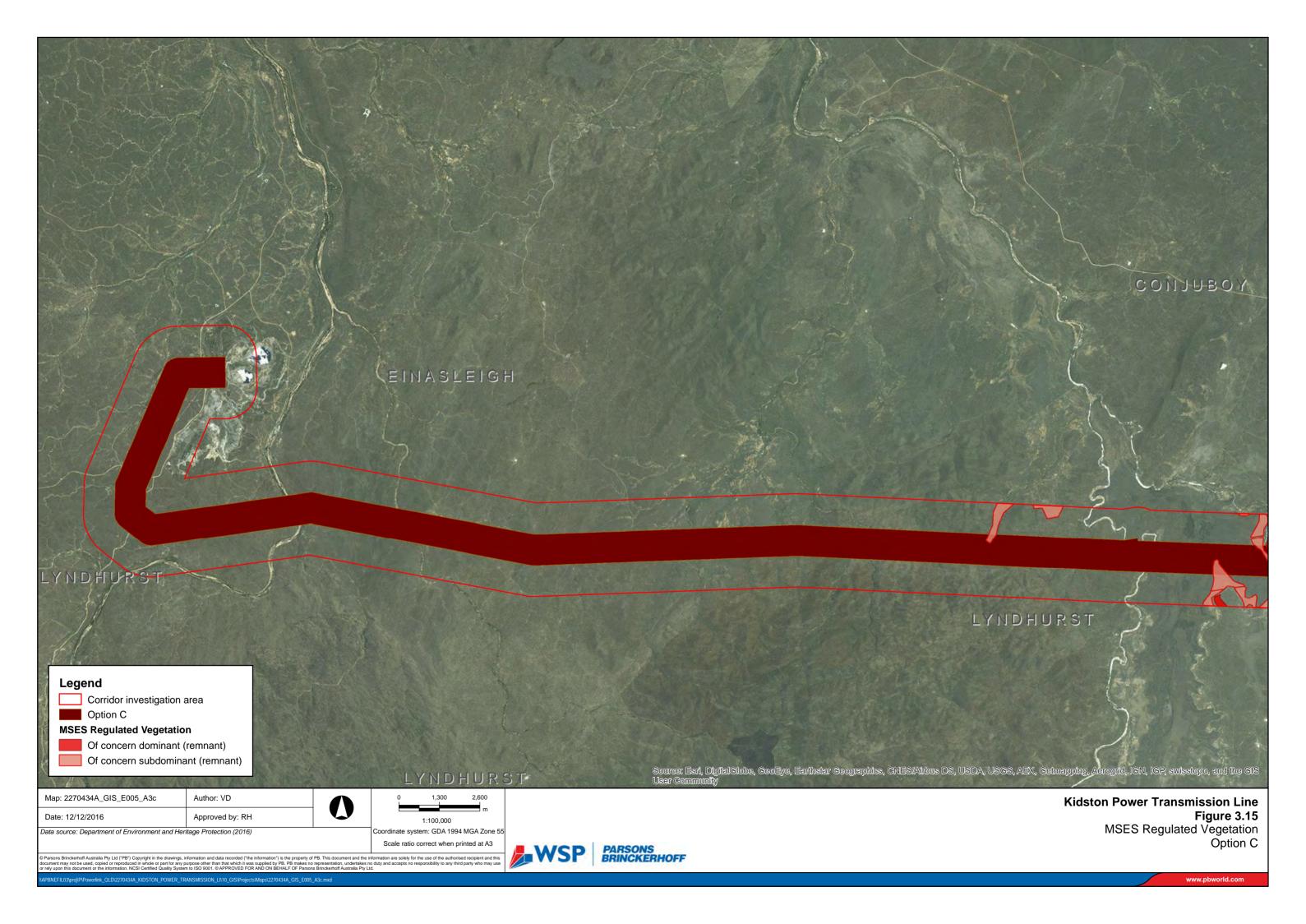


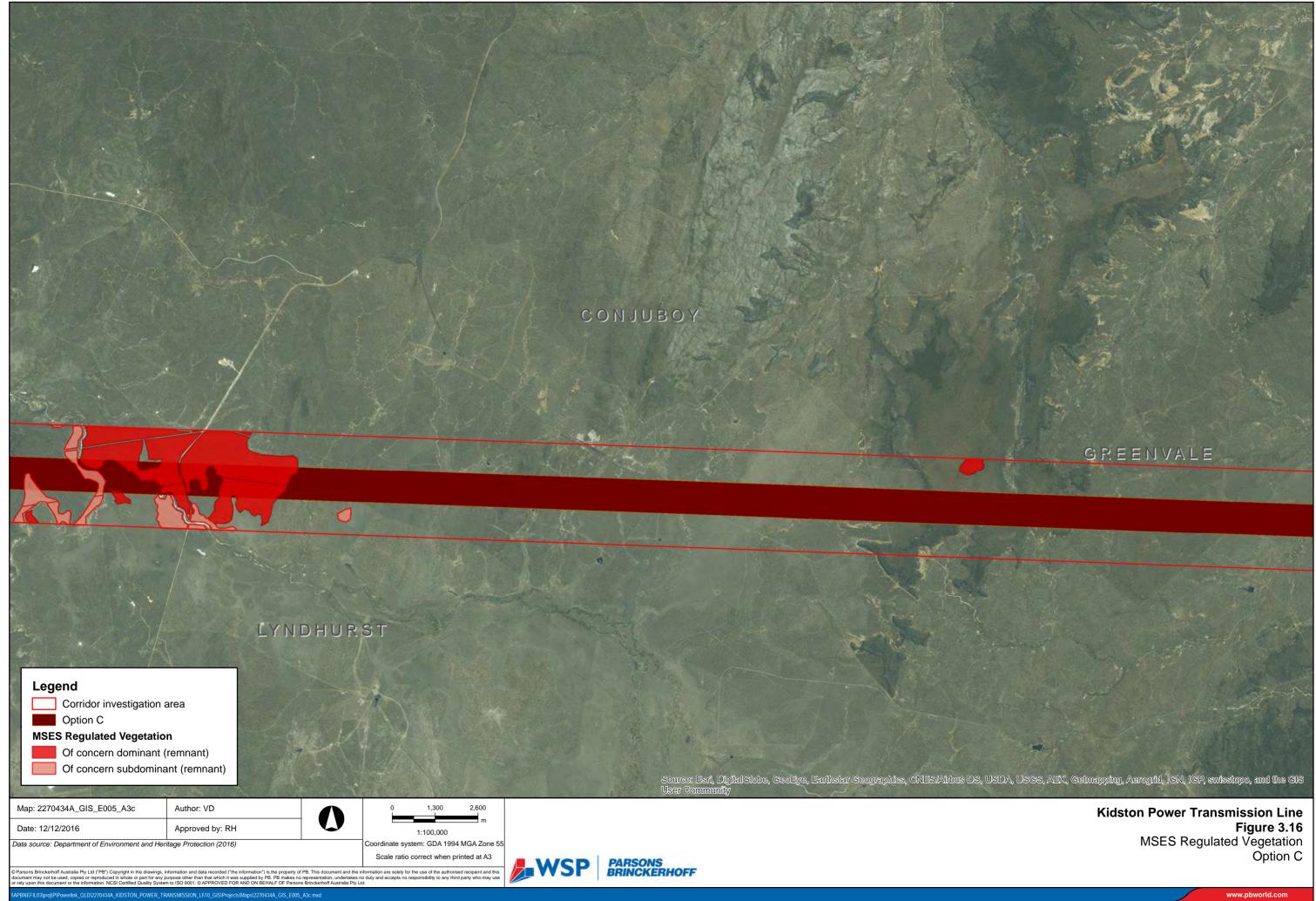


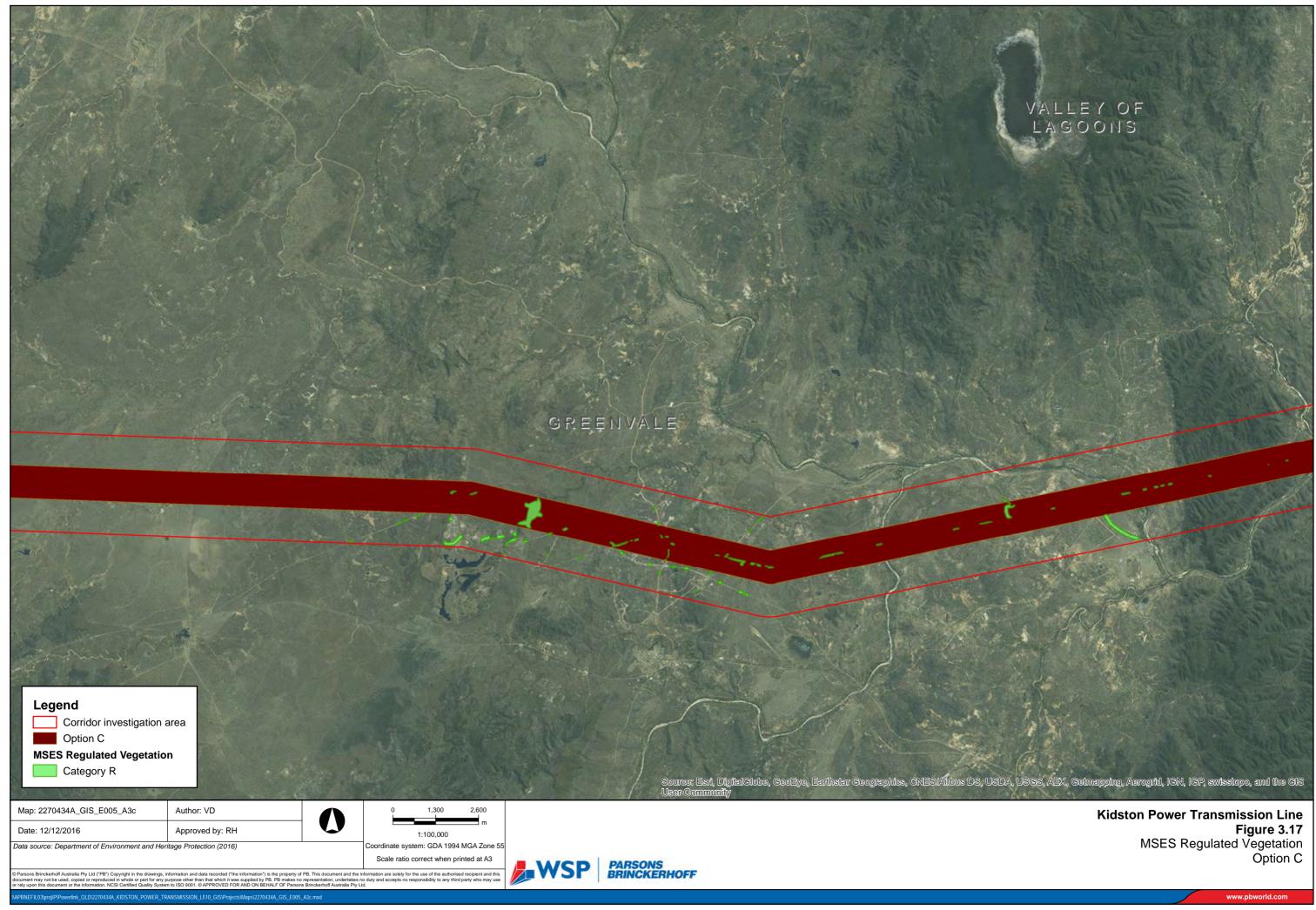


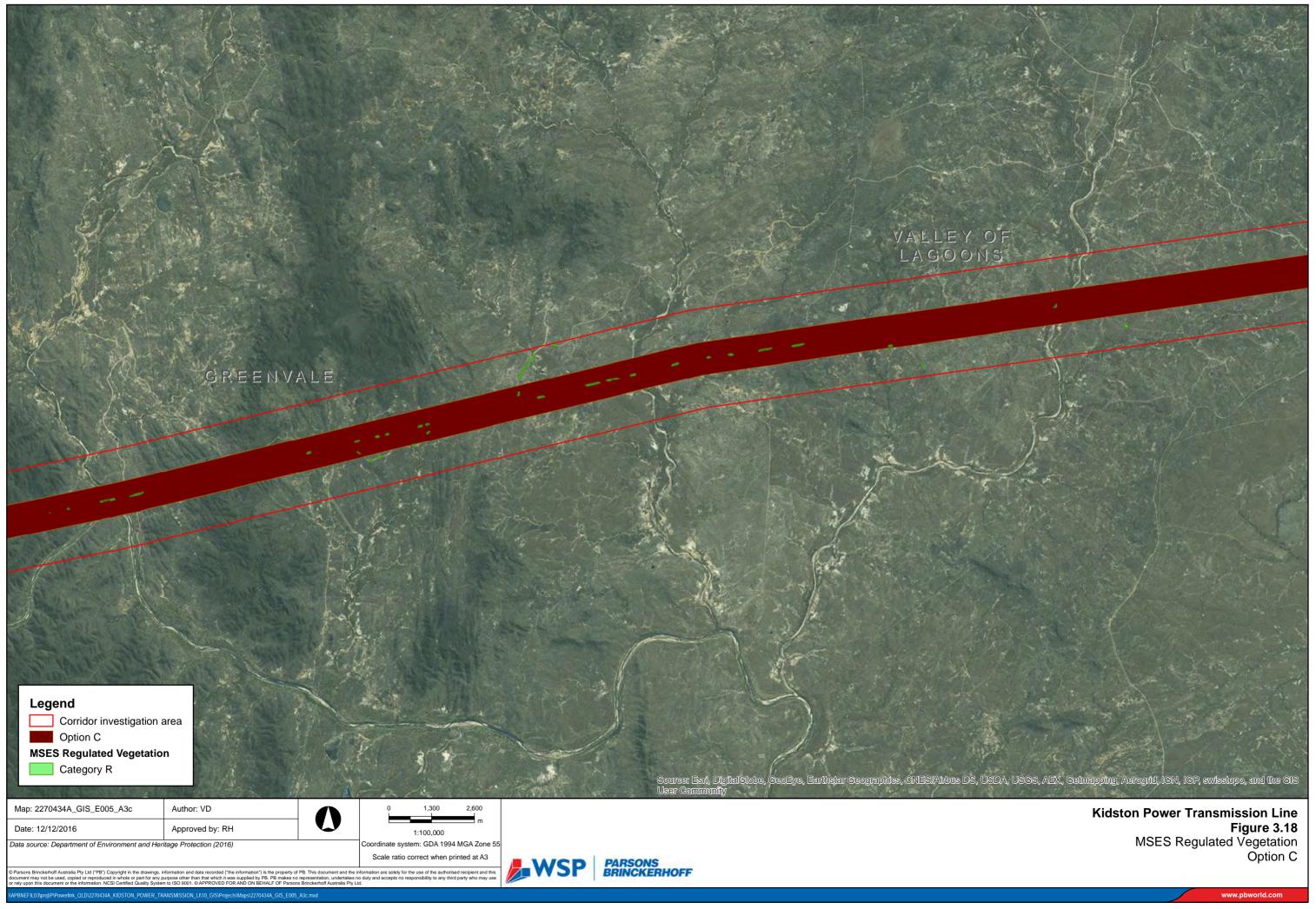


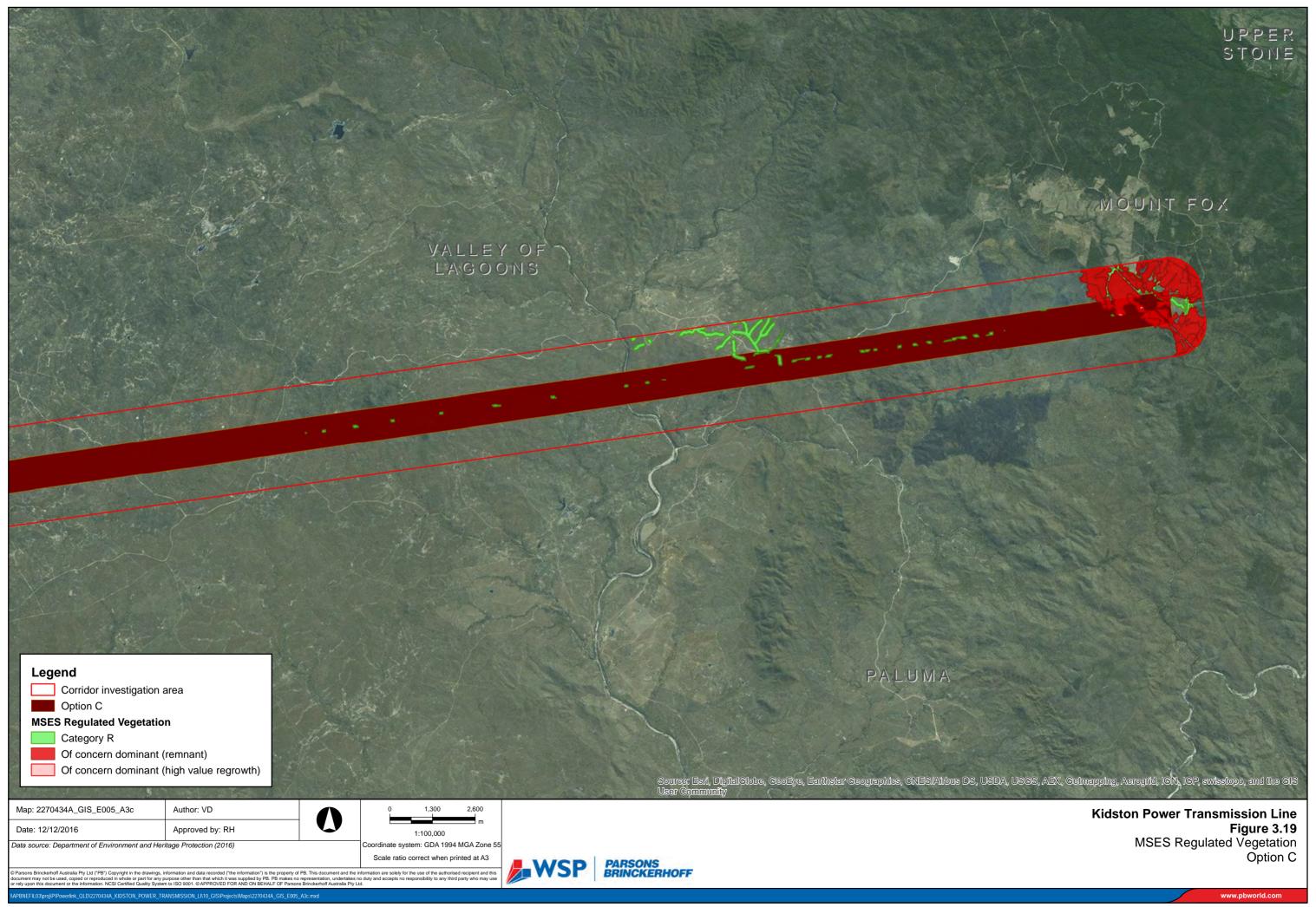












### 3.1.3 DNRM regional ecosystem mapping

The results of the GIS analysis of DNRM's regional ecosystems mapping determined the extent of mapped of concern dominant and sub-dominant, and least concern regional ecosystems that is of potential relevance to each corridor option.

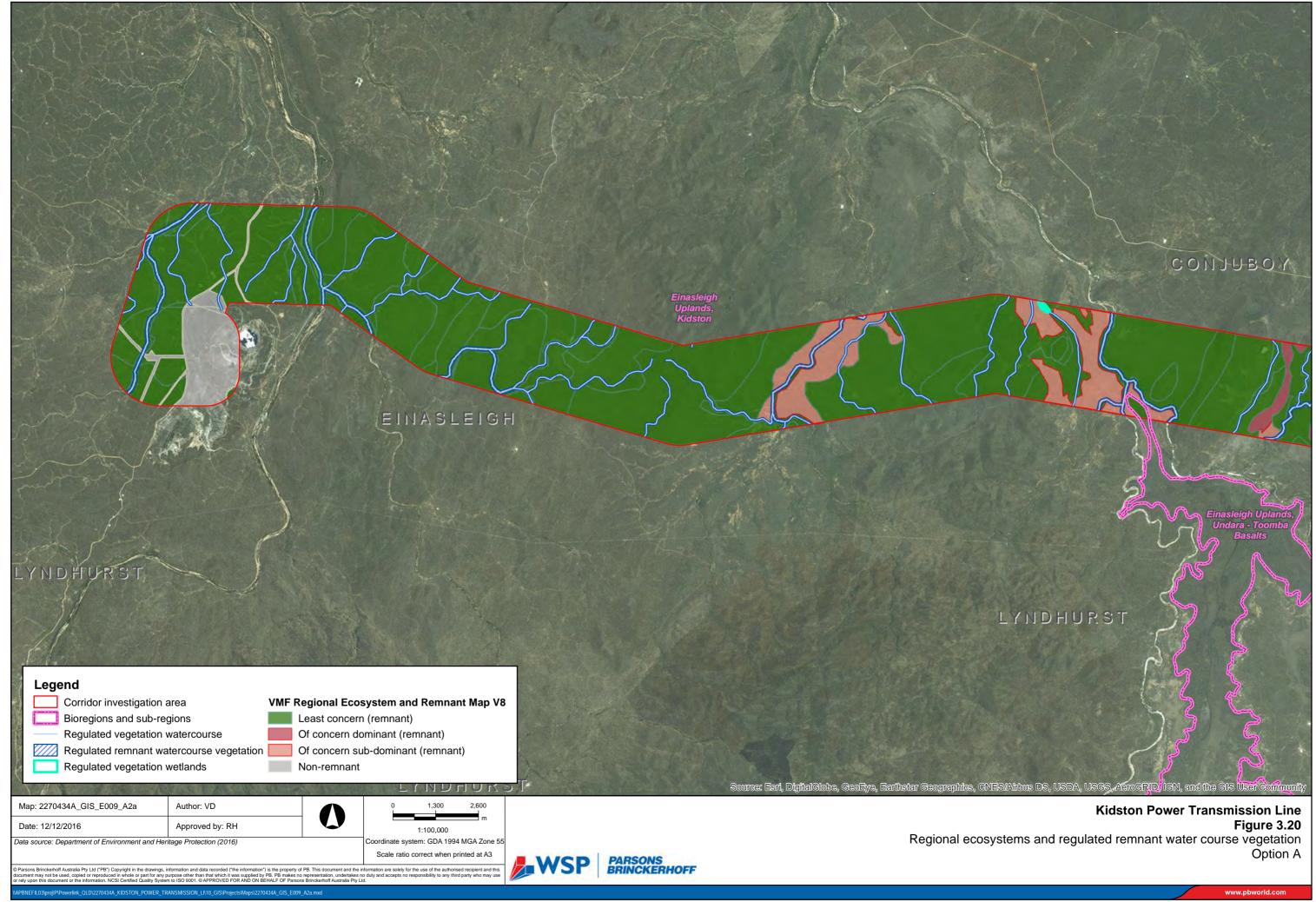
The extent of DNRM's mapped regional ecosystems that is of potential of relevance to each corridor investigation area, is presented on Figure 3.20–Figure 3.34.

The area (ha) of dominant and sub-dominant of concern, and least concern regional ecosystems that is of potential relevance to each corridor investigation area is presented in Table 3.4. The analysis of regional ecosystems for each corridor option is presented in Appendix G.

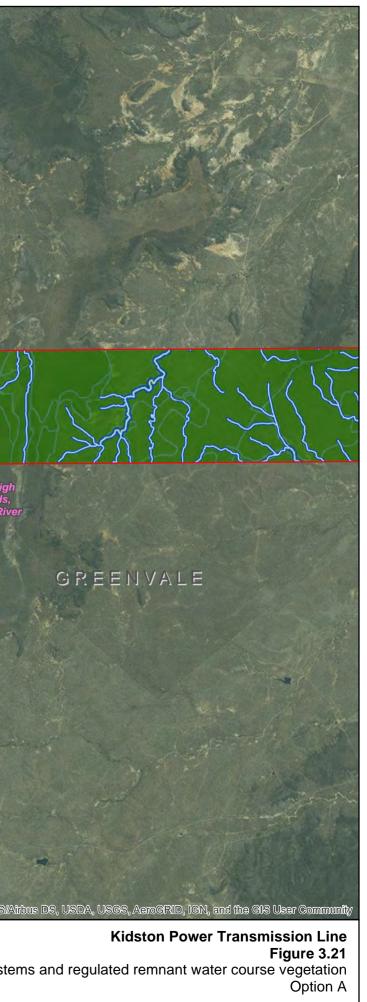
Option C is of least constraint in terms of potential impacts upon of concern dominant and sub-dominant regional ecosystems (refer Table 3.4).

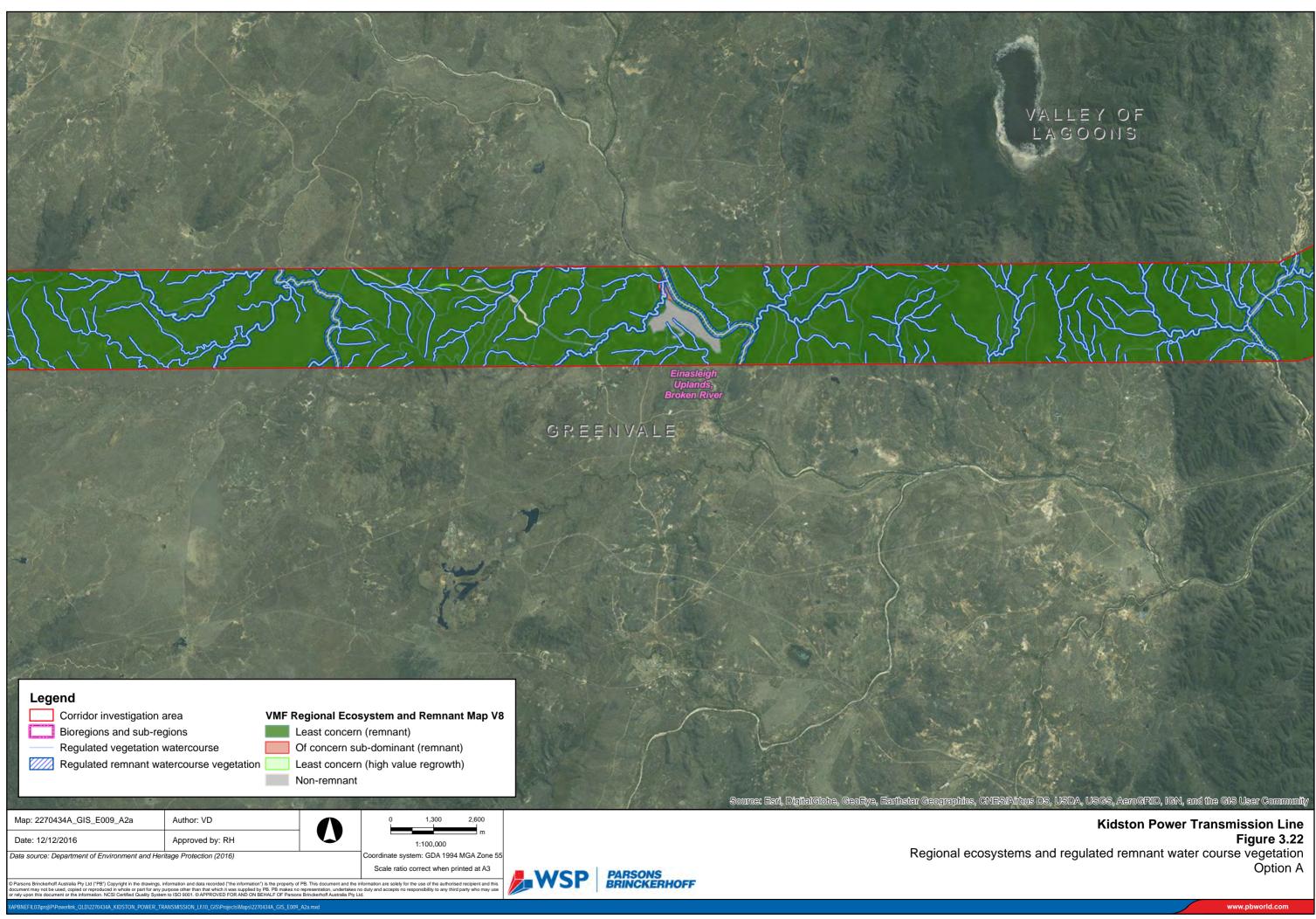
CORRIDOR	R	EMNANT VEGETATION (I	HA)	REGROWTH VEGI	ETATION (HVR) (HA)	NON-REMNANT (HA)	
	OF CONCERN DOMINANT	OF CONCERN SUB- DOMINANT	LEAST CONCERN	OF CONCERN DOMINANT	LEAST CONCERN	VEGETATION	WATER
Option A	2,059	1,008	50,675	3.6	62	2,675	20
Option B	1,891	442	51,121	2.4	202	2,379	117
Option C	1,519	334	52,609	2.4	266	2,458	29

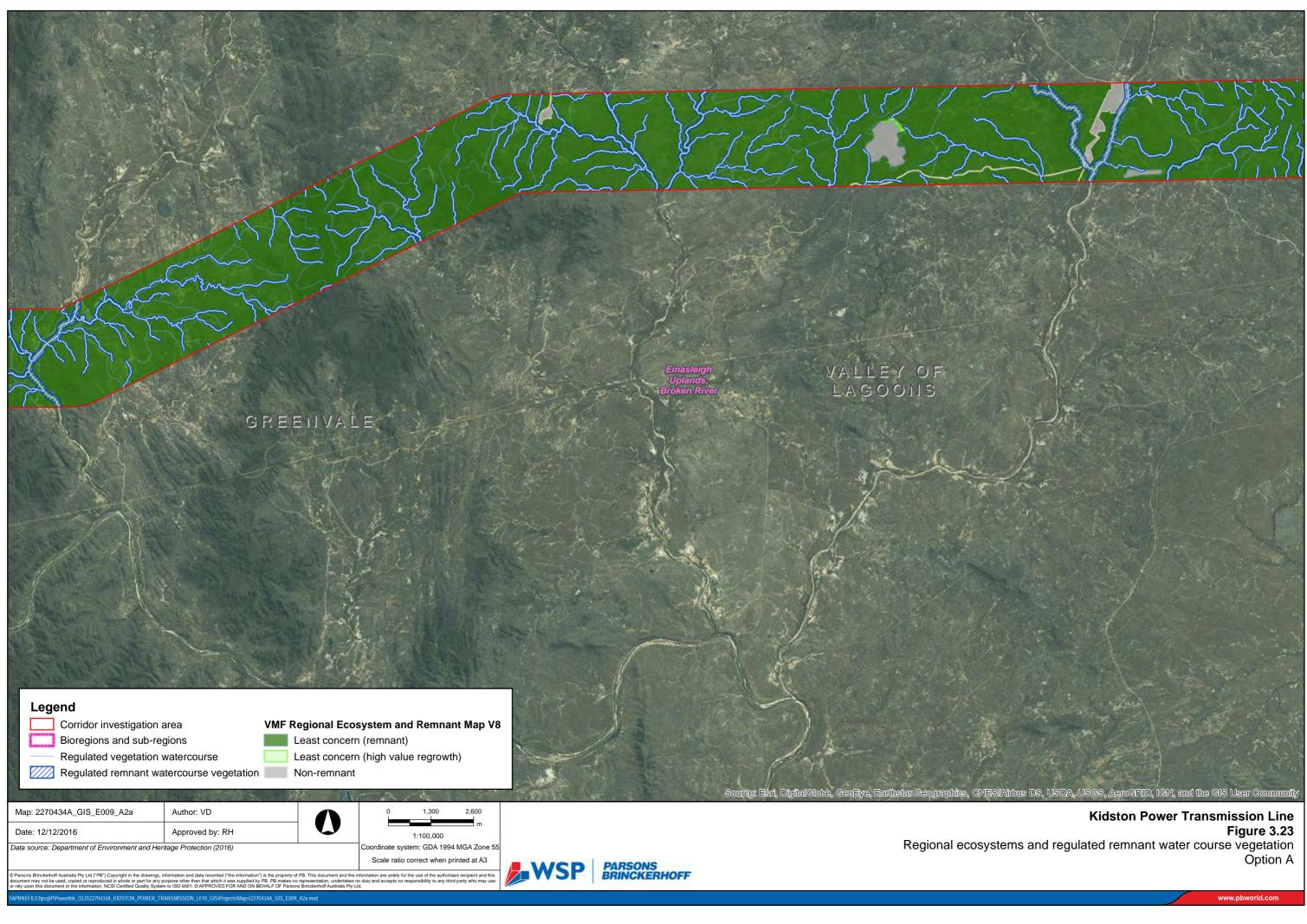
#### Table 3.4 Extent of mapped regional ecosystems of relevance to each corridor option



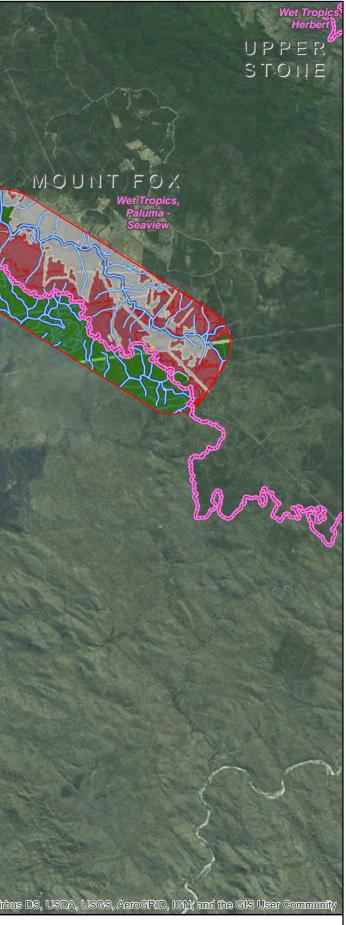
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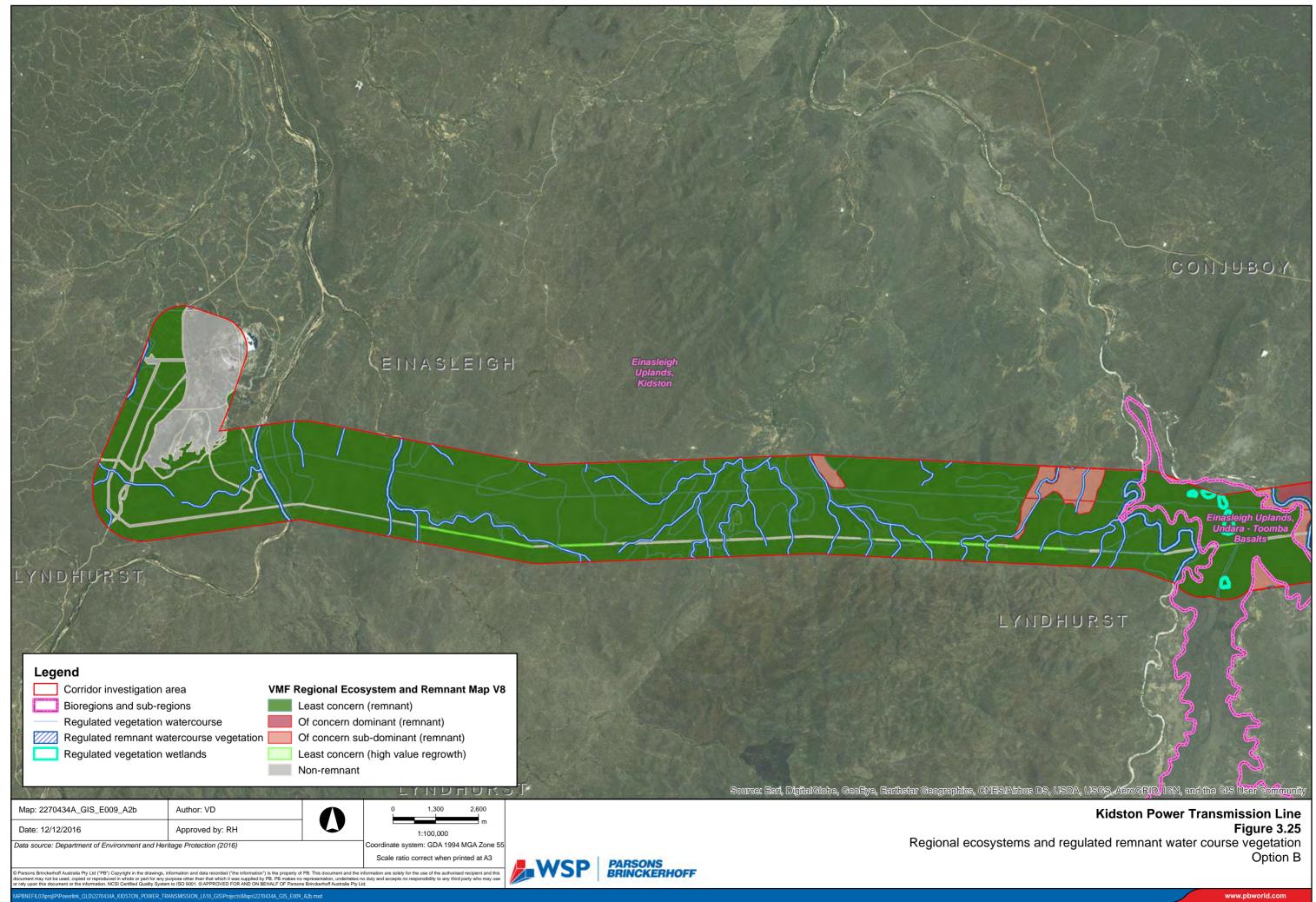
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Map: 2270434A_GIS_E009_A2a Date: 12/12/2016 Data source: Department of Environment and Parsons Brinckerhoff Australia Pty Ltd ('PB') Copyright in the drawin cument may not be used, copied or reproduced in whole or part for ref y upon this document or the information. NCSI Certified Quality Certified			0 1,300 2,600 1:100,000 Coordinate system: GDA 1994 MGA Zone 55 Scale ratio correct when printed at A3 symmation are solely for the use of the authorised recipient and this day and accepts no responsibility to any third party who may use	WSP	PARSONS BRINCKERHOFF	ource: Esri, DigitalGlobe, GeoEye, Earthstar Geographios, GNE Regional ecosy

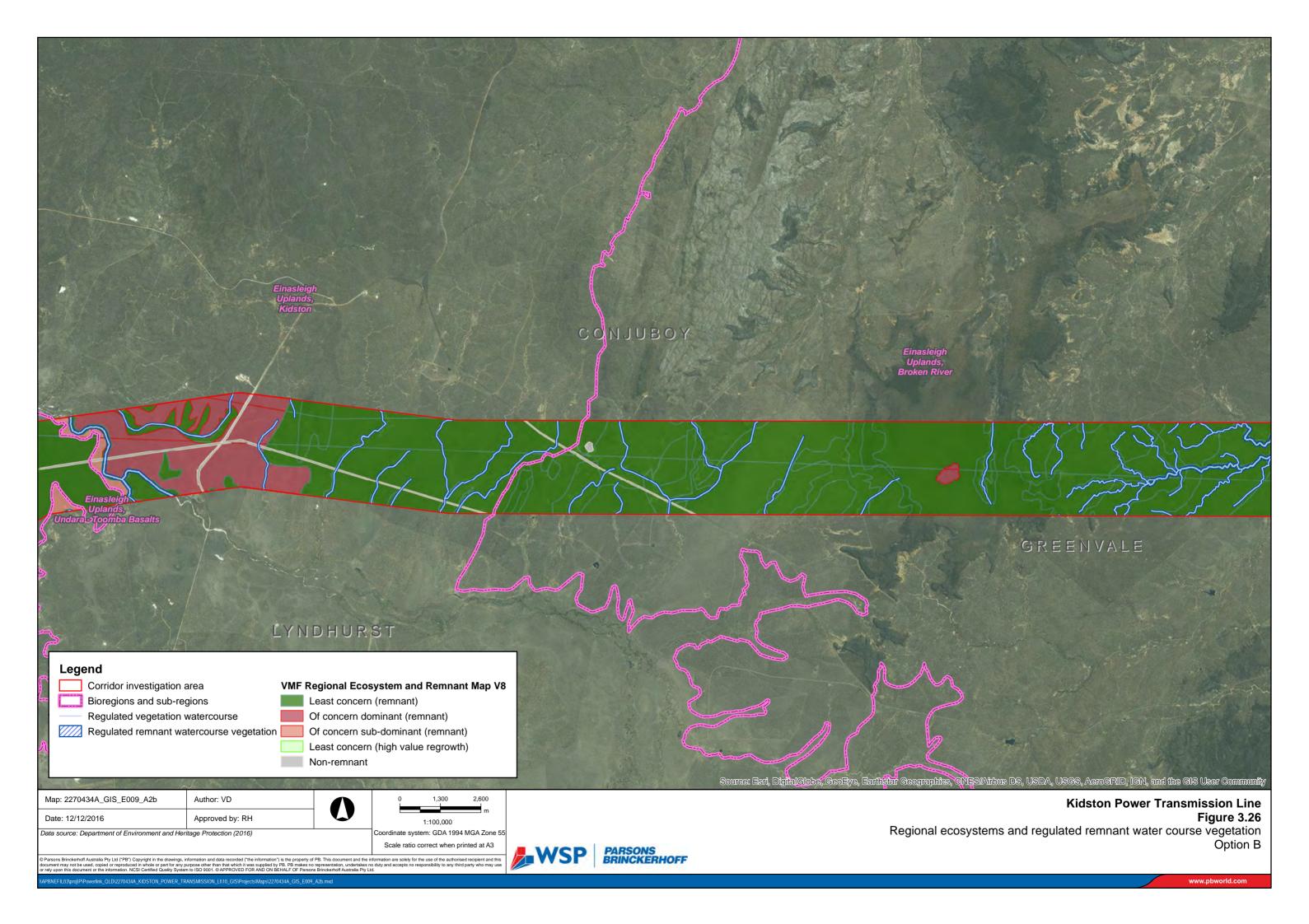


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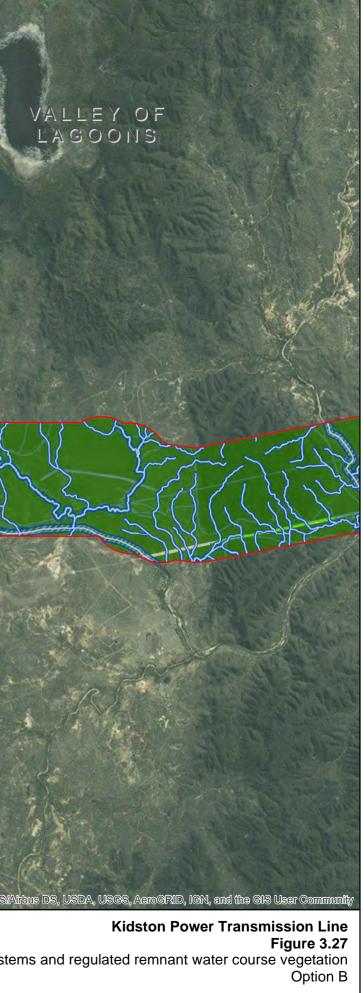
Kidston Power Transmission Line Figure 3.24 stems and regulated remnant water course vegetation Option A

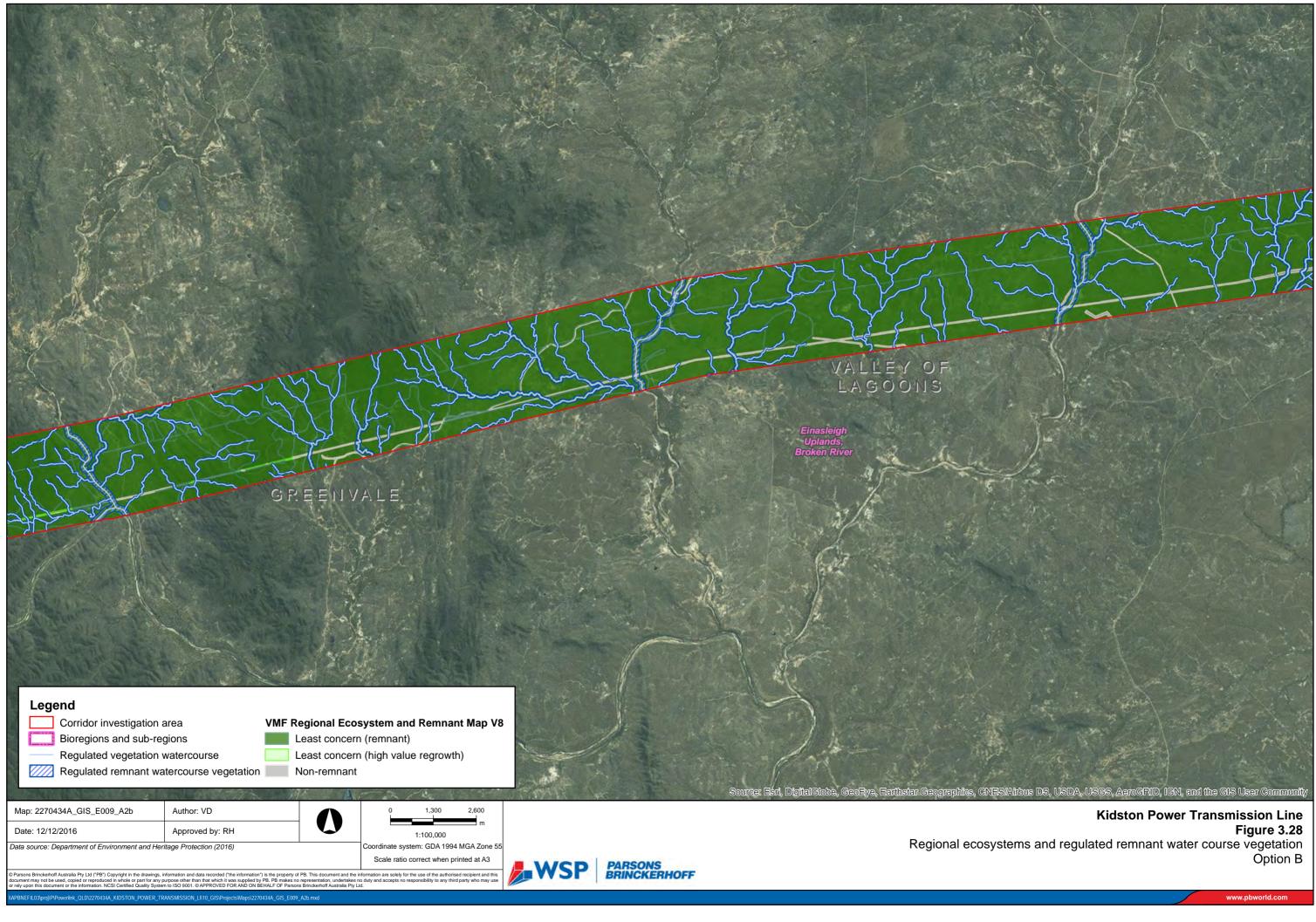
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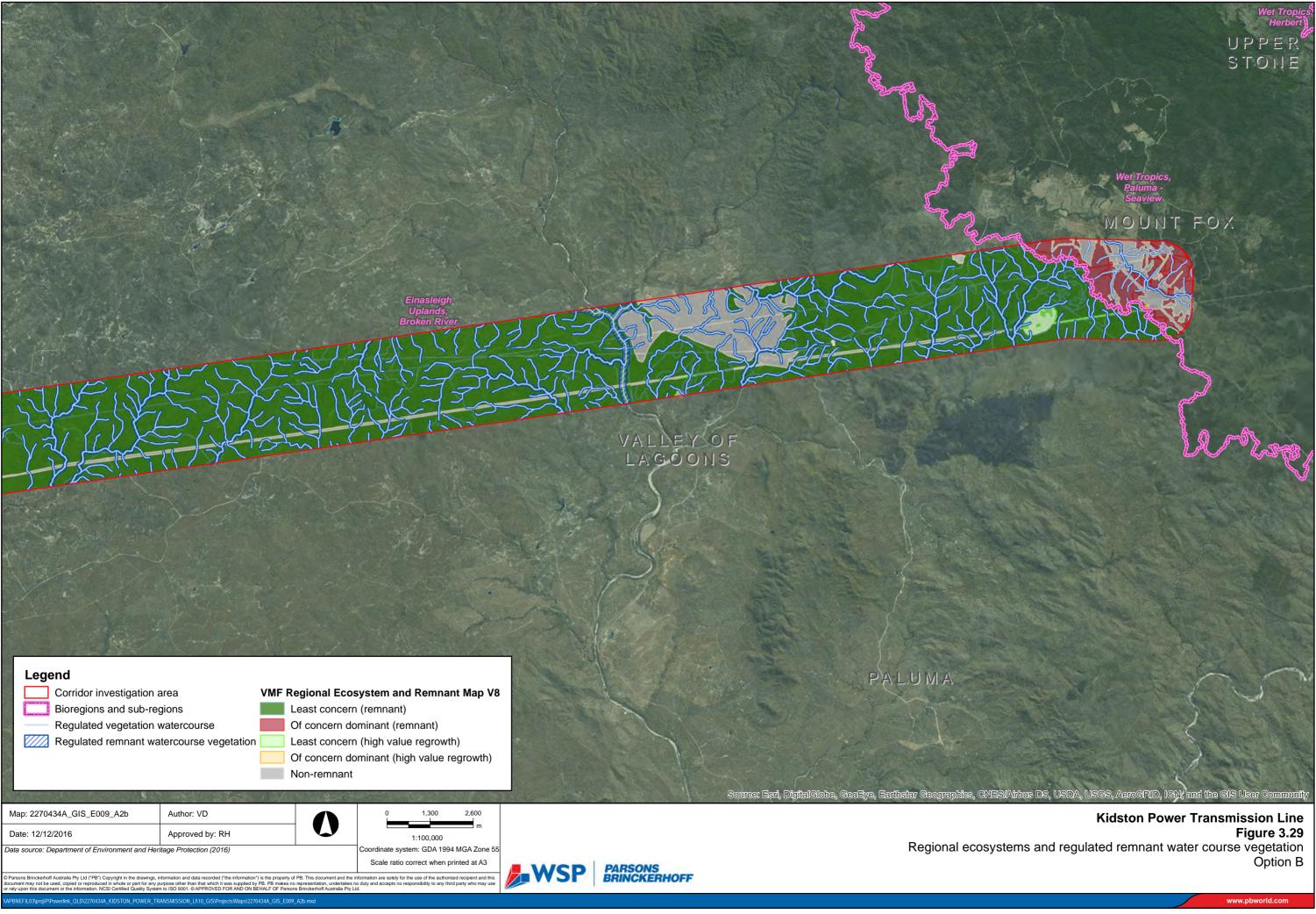


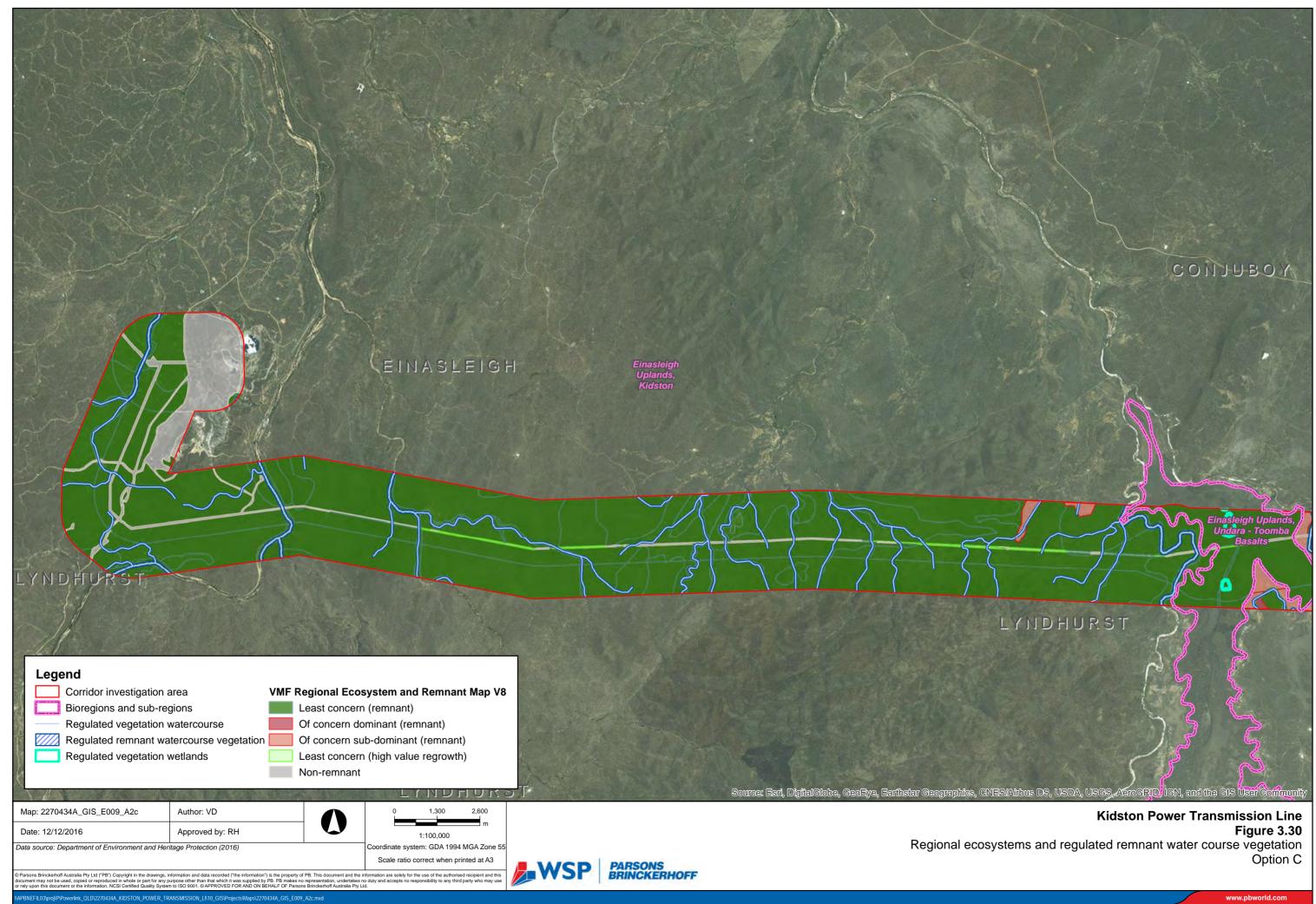


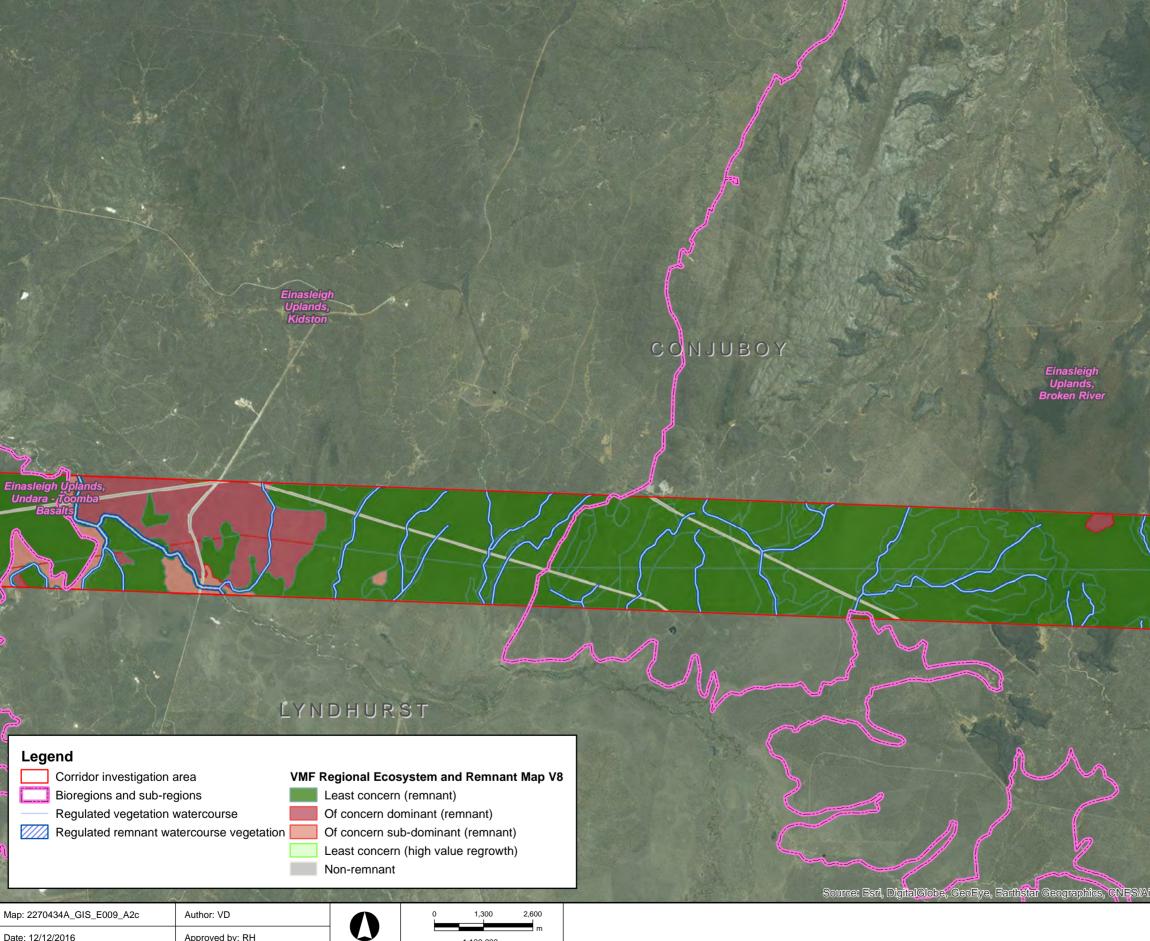
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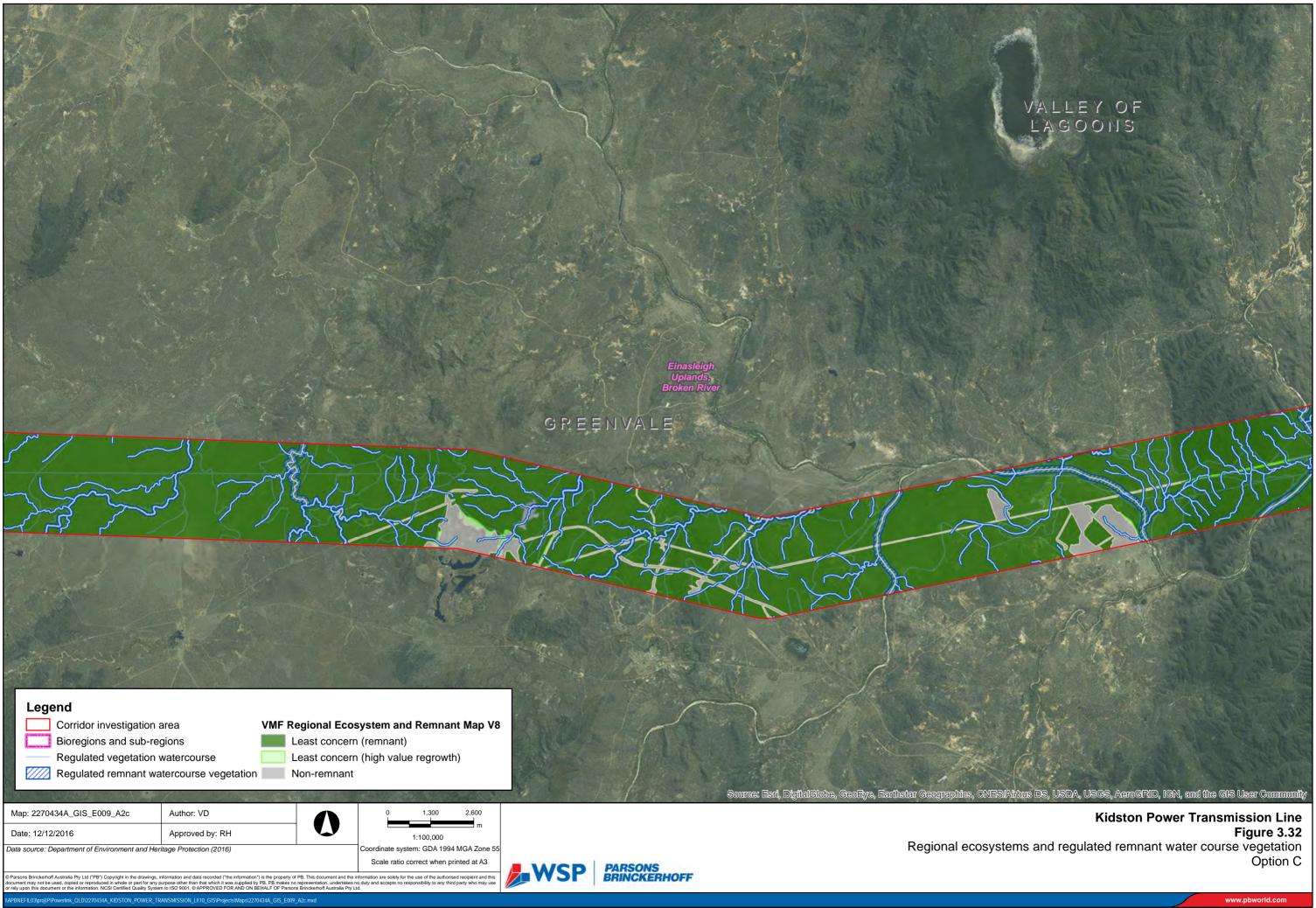
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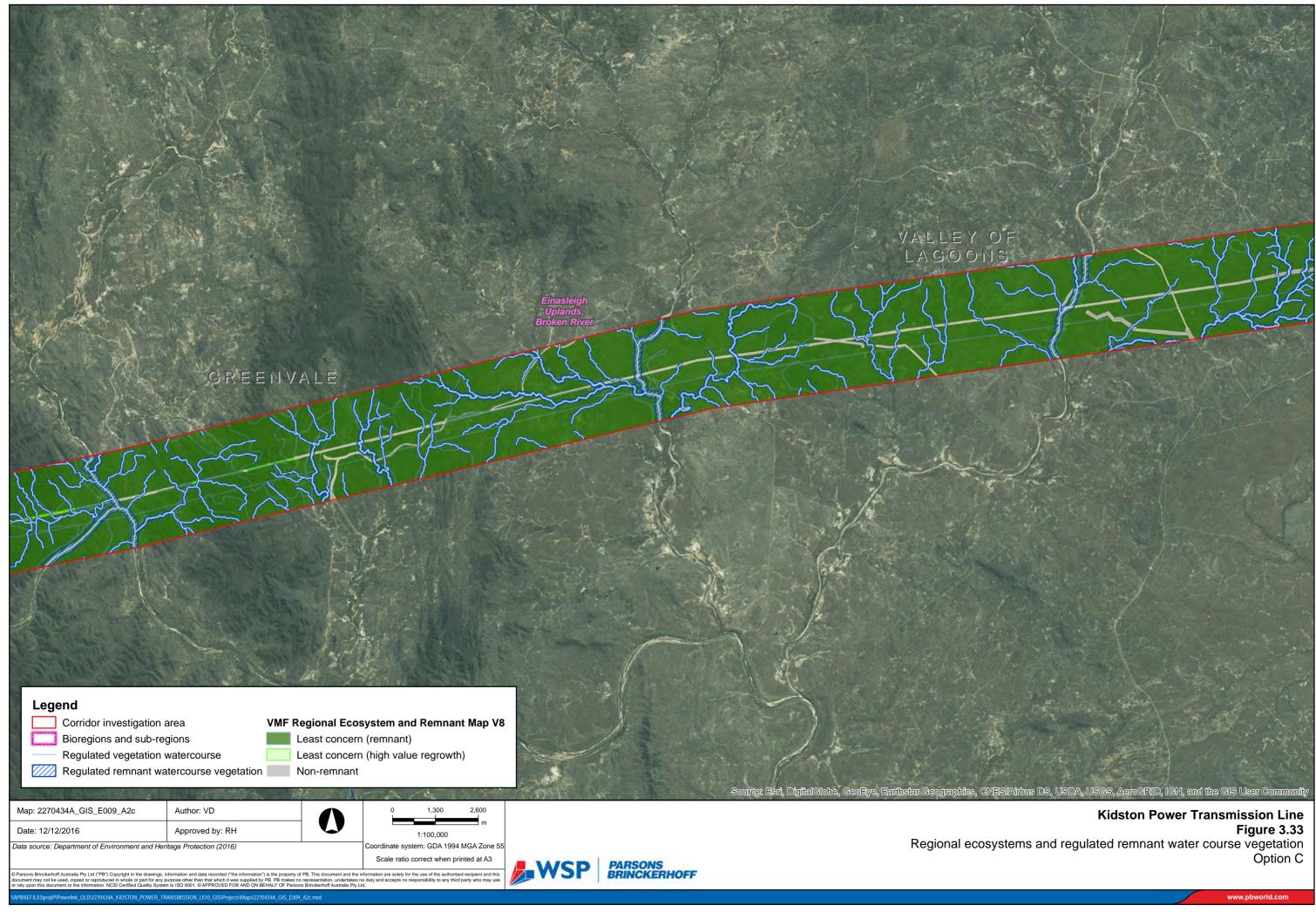
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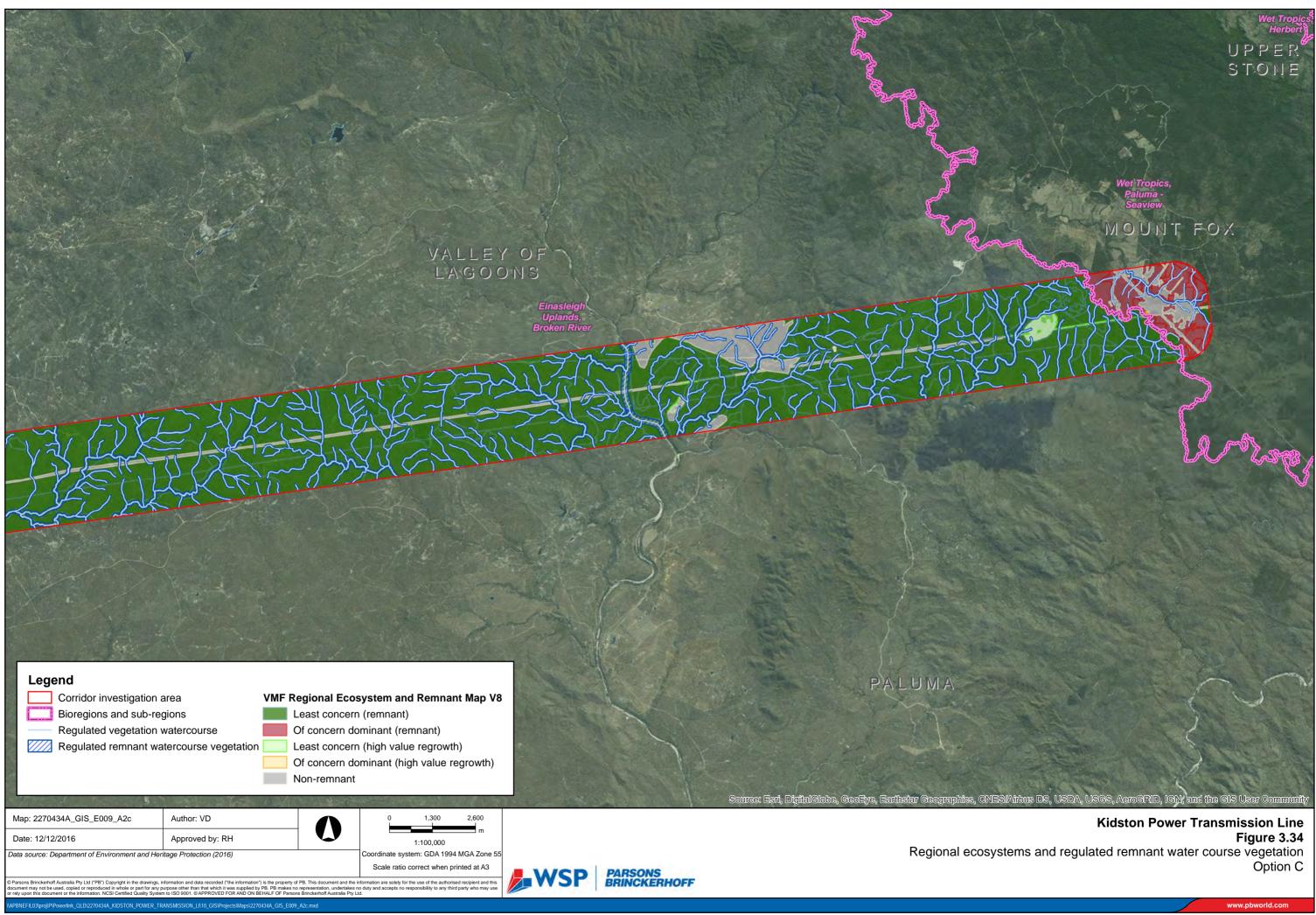
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## PROPOSED GENEX KIDSTON CONNECTION PROJECT

## **Corridor Selection Report**

PREPARED BY

QUEENSLAND ELECTRICITY TRANSMISSION CORPORATION LIMITED

(ACN 078 849 233) trading as "POWERLINK"

The woody vegetation assessment performed by Virtual GIS, used GIS remote sensing to predict the extent of woody remnant vegetation, and the potential extent of woody remnant vegetation and regrowth vegetation in each corridor investigation area. The results of the woody vegetation mapping is useful for identifying potential areas of inconsistency between the woody vegetation assessment and the DNRM mapping, for which such areas would be targeted as part of the ground-truthing surveys during the next phase of the Project.

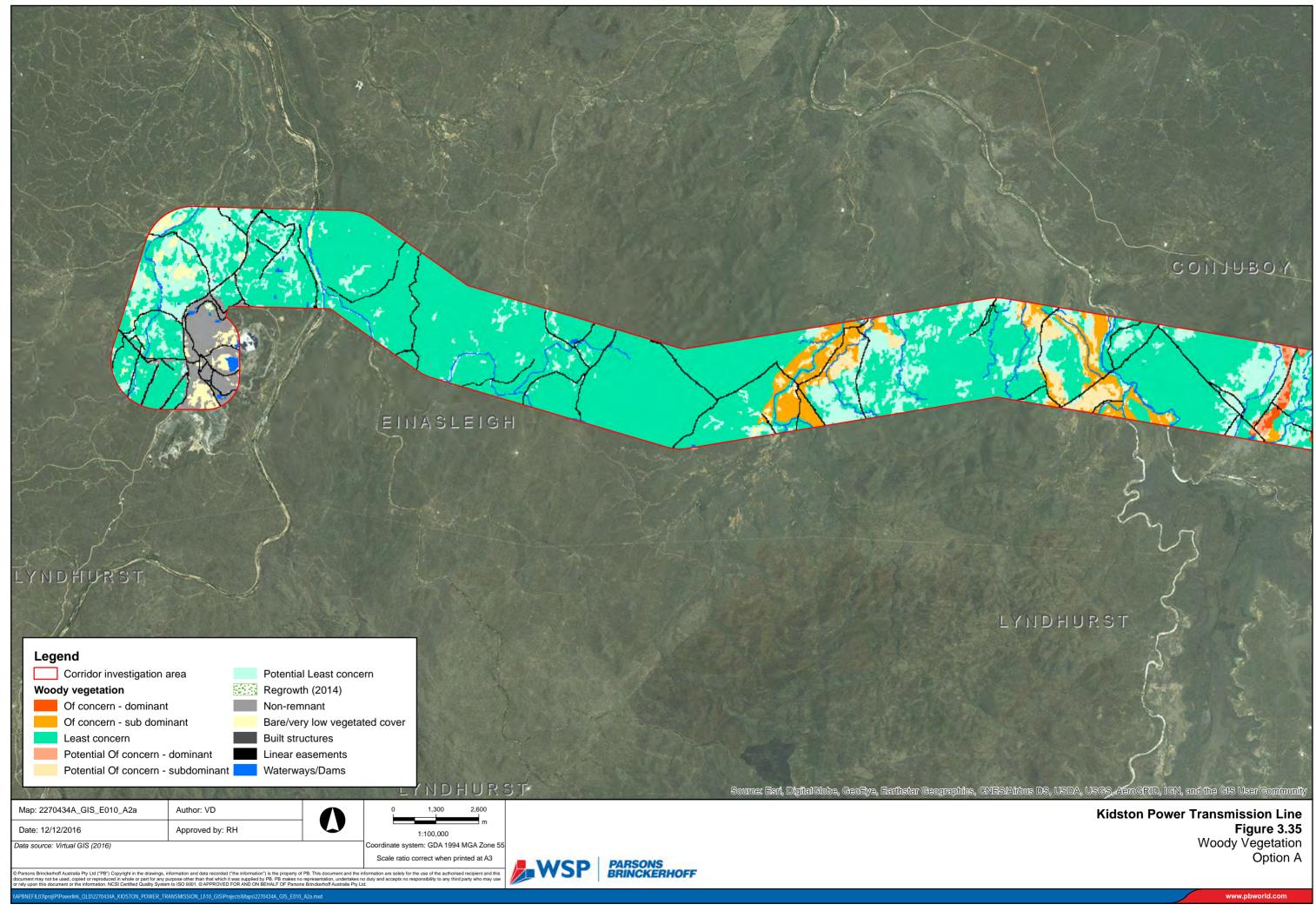
The woody vegetation that is of potential relevance to each corridor option, as an output of remote sensing of woody vegetation, is presented on Figure 3.35–Figure 3.49.

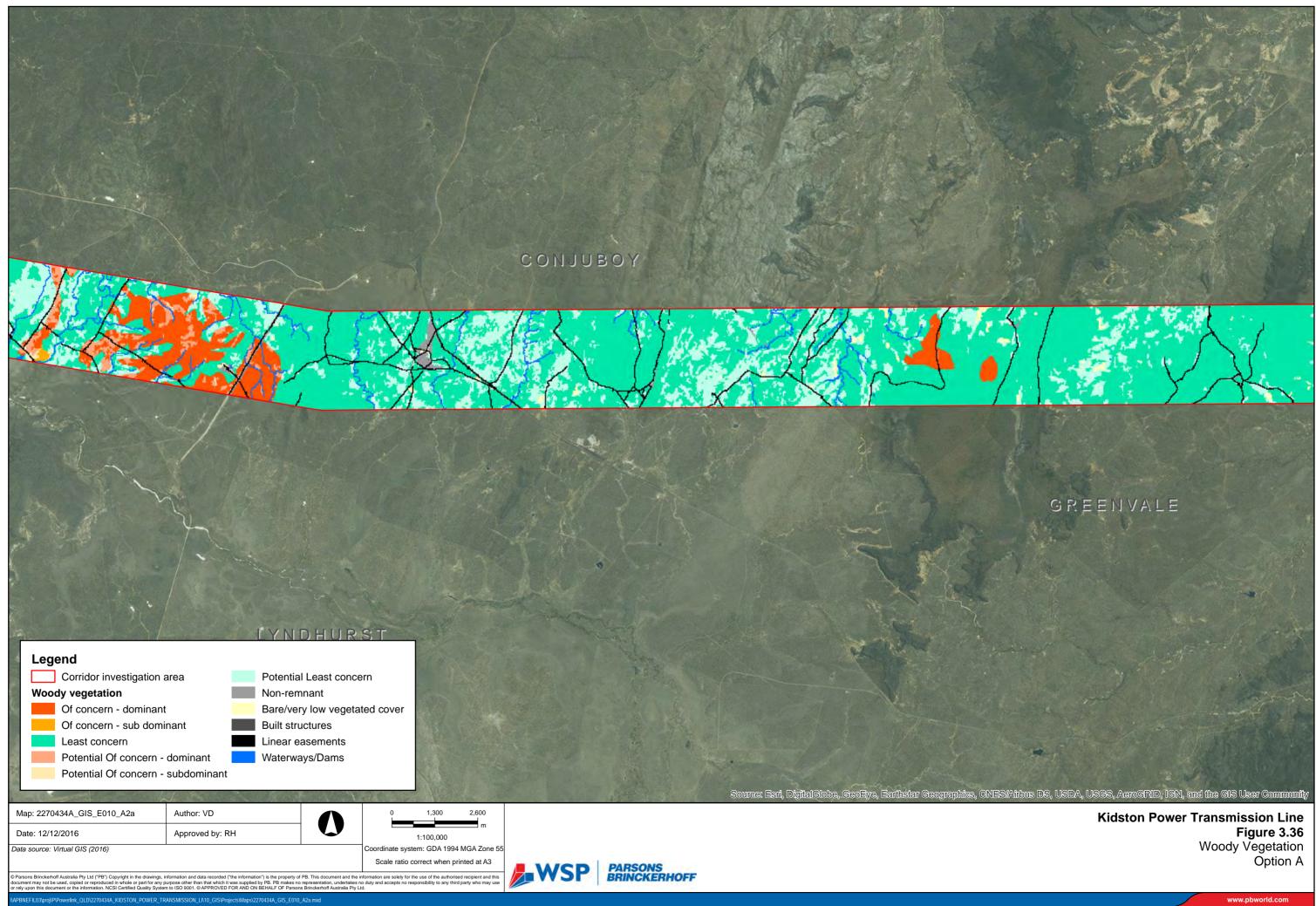
The area (ha) of dominant and sub-dominant of concern regional ecosystems that is of potential relevance to each corridor investigation area is presented in Table 3.5. The results of the analysis of woody vegetation for each corridor option is presented in Appendix G.

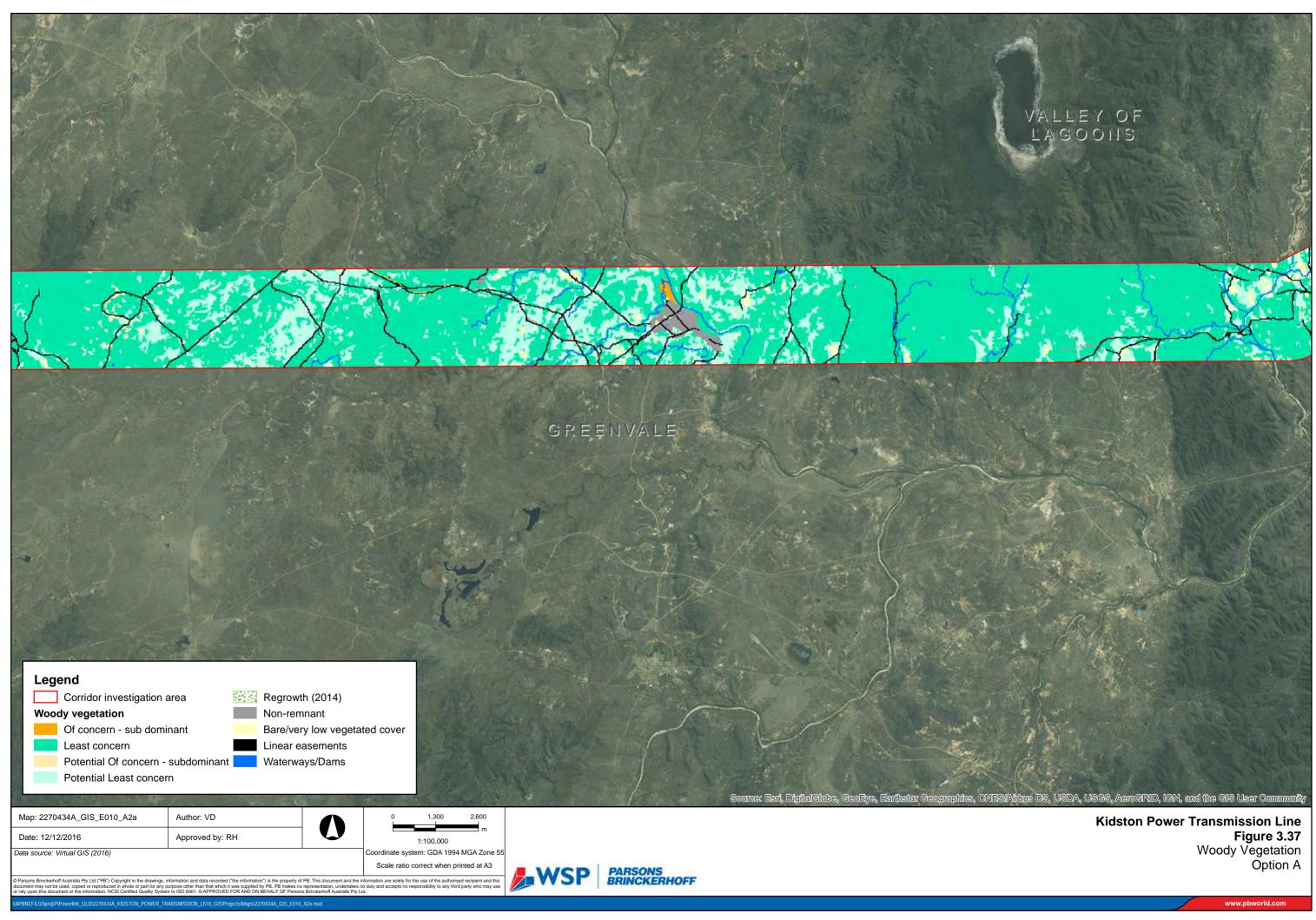
In reference to the predicted and potential woody remnant vegetation in Table 3.5, Option C is of least constraint in terms of potential impacts upon potential and predicted of concern dominant. On the other hand, Option B is of least constraint in terms of potential impacts upon potential and predicted of concern subdominant regional ecosystems. However, a greater emphasis should be applied to the predicted of concern dominant remnant vegetation being of greater significance, which therefore suggests that Option C is of least ecological constraint.

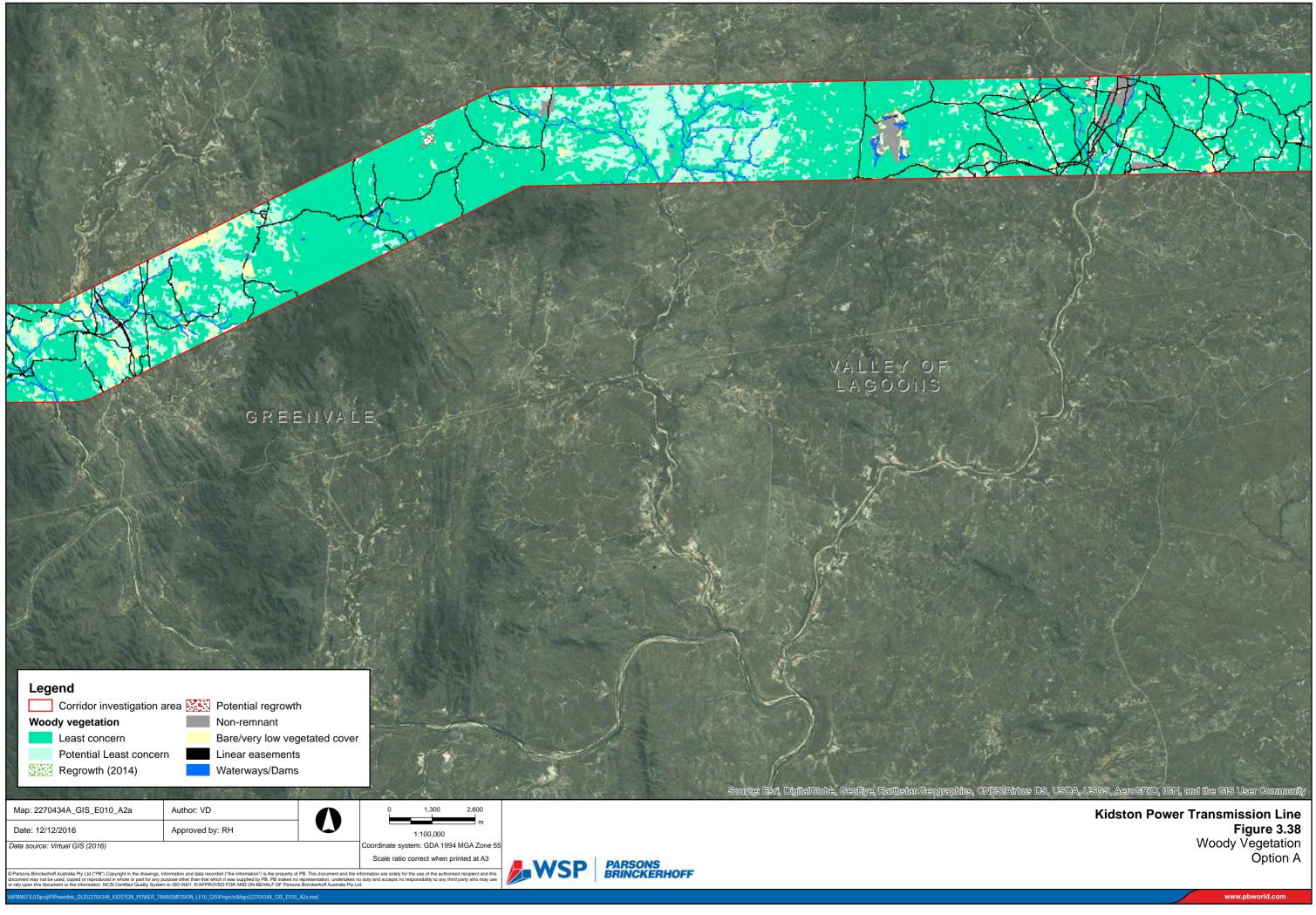
CORRIDOR	PREDICTED WOODY REMNANT VEGETATION (HA)		POTENTIAL WOODY REMNANT VEGETATION (HA)			POTENTIAL REGROWTH			NON-REMNANT		
	OF CONCERN DOMINANT	OF CONCERN SUB- DOMINANT	LEAST CONCERN	OF CONCERN DOMINANT	OF CONCERN SUB- DOMINANT	LEAST CONCERN	OF CONCERN DOMINANT	LEAST CONCERN	UNCLASSIFIED REGROWTH	VEGETATION	BARE EARTH, WATER AND EASEMENT
Option A	1,535	549	37,602	240	364	9,002	6.9	48	235	2,174	4,670
Option B	1,460	295	37,969	344	87	8,439	6.6	214	219	1,812	5,428
Option C	1,223	618	38,808	431	37	8,494	28.8	172	291	1,718	5,771

#### Table 3.5 Remotely sensed predicted and potential extent of woody remnant and regrowth vegetation of relevance to each corridor option

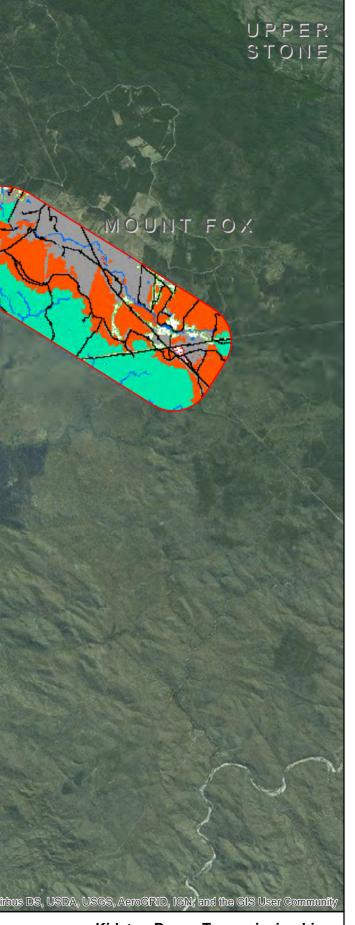




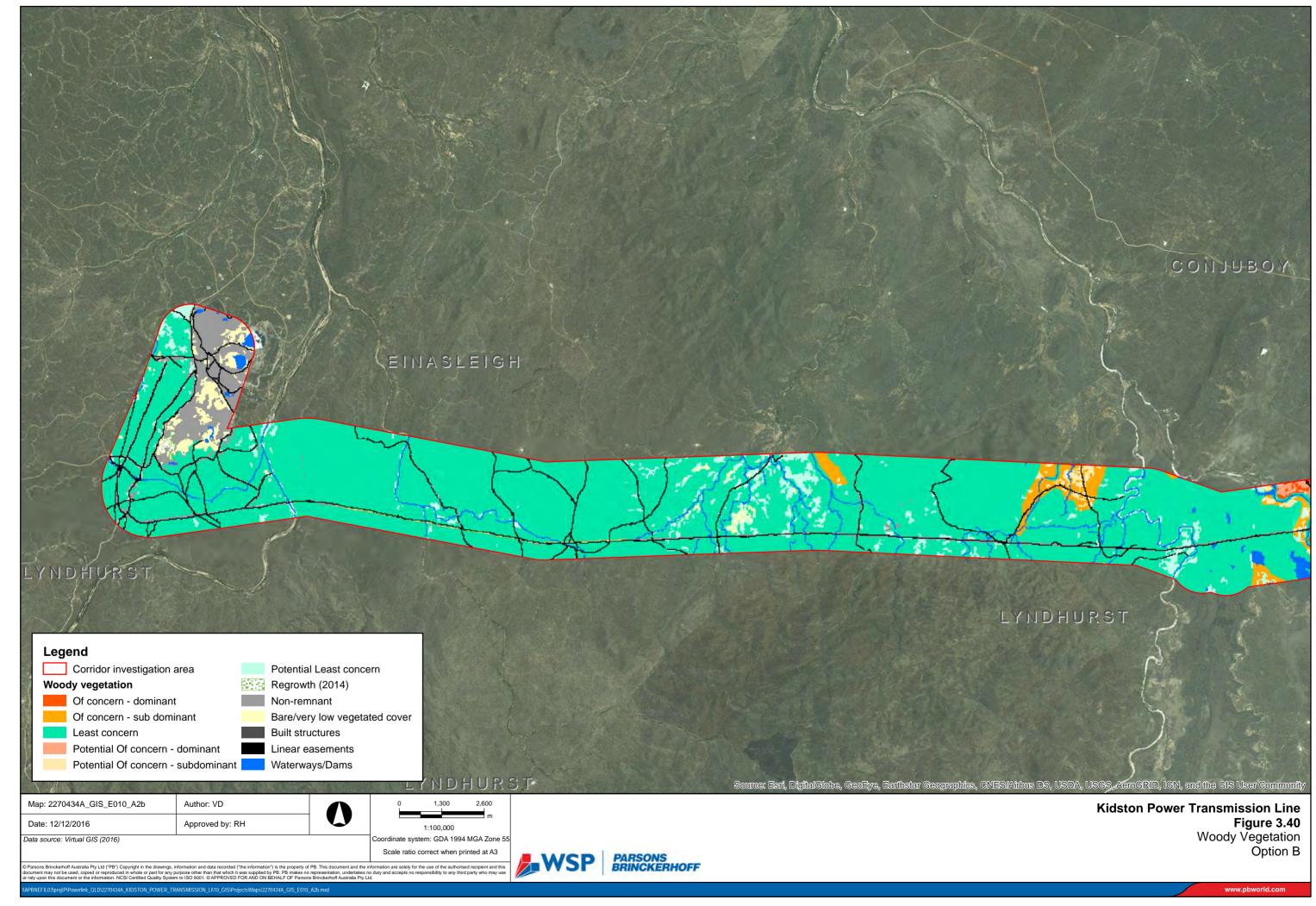


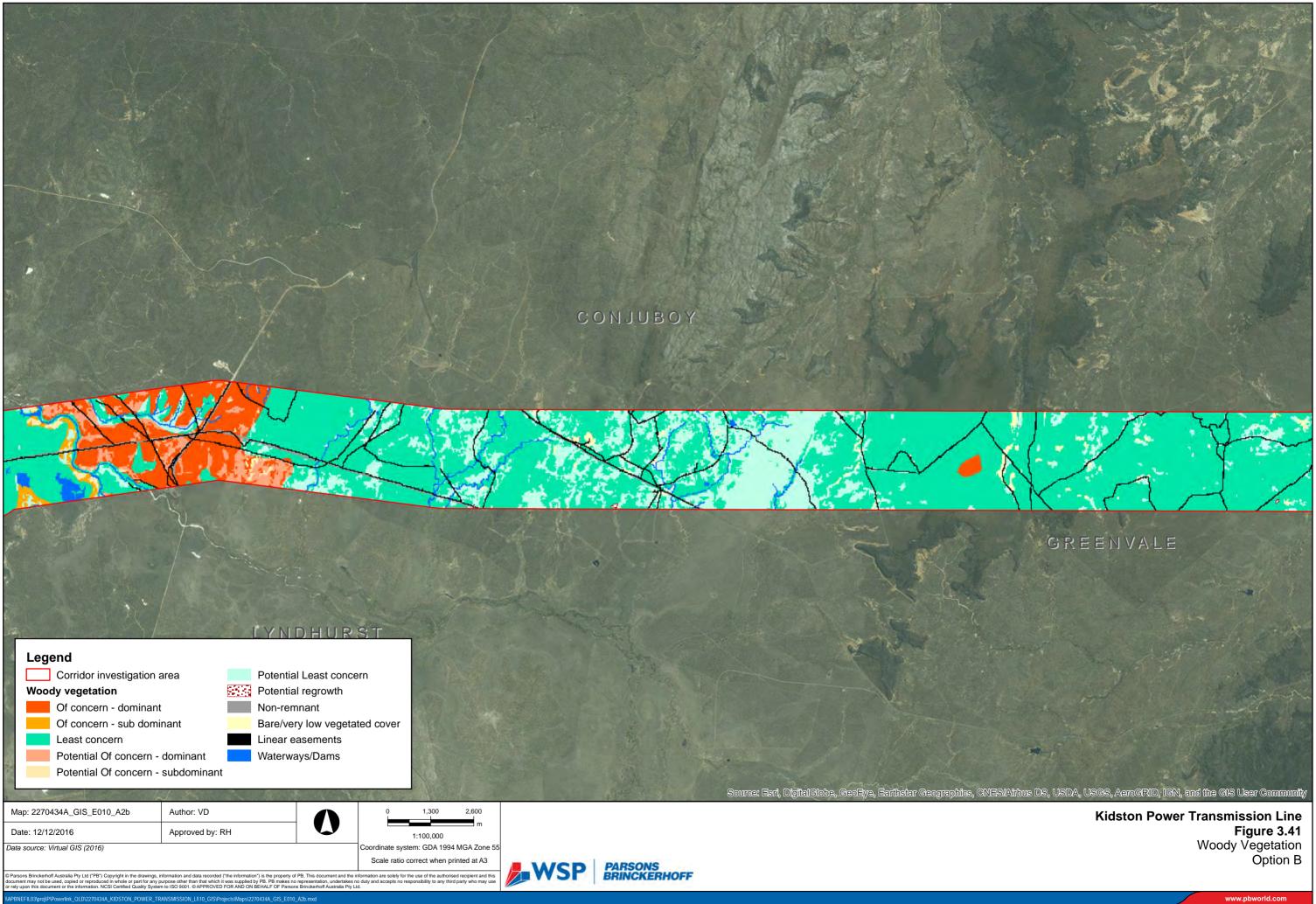


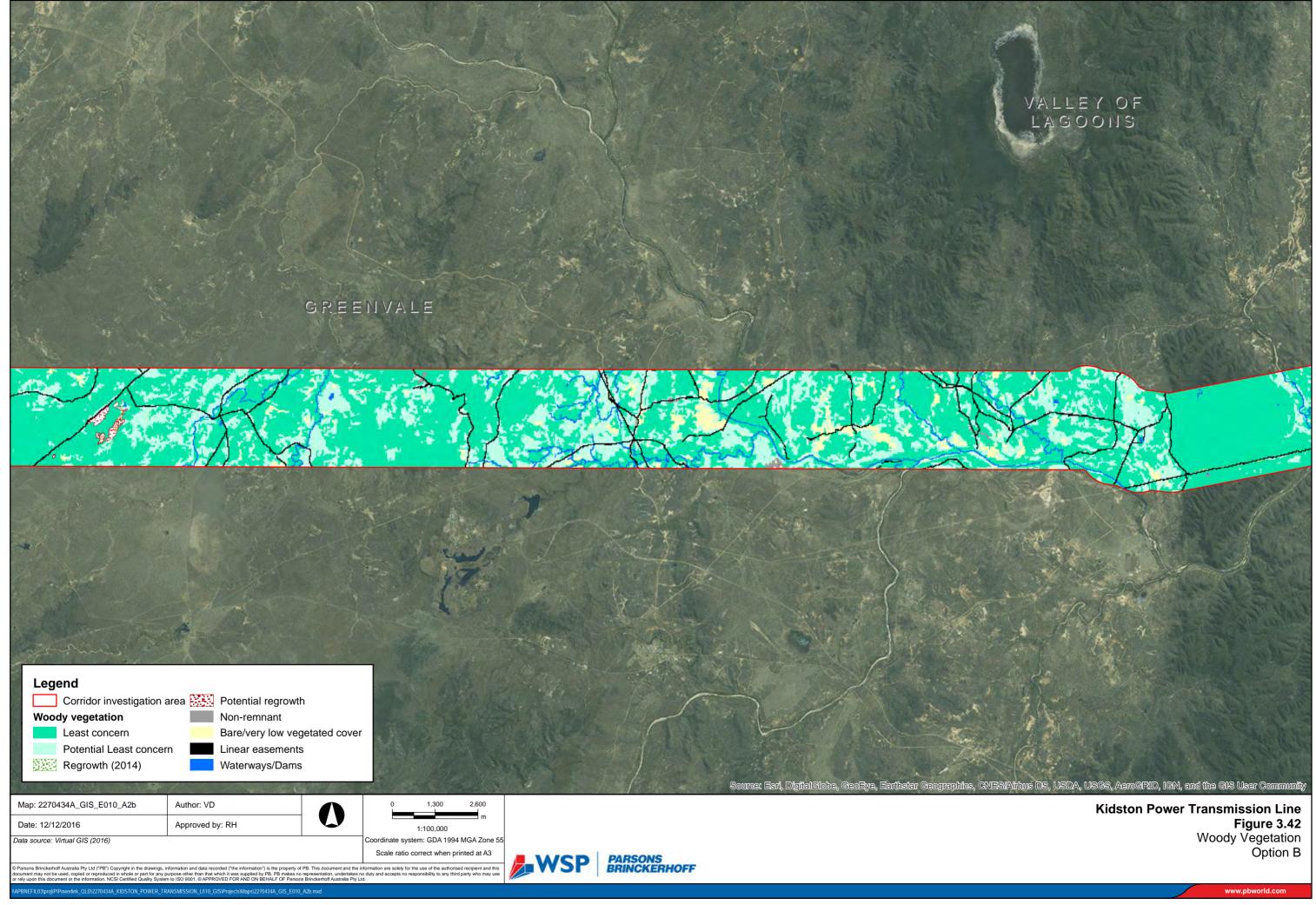
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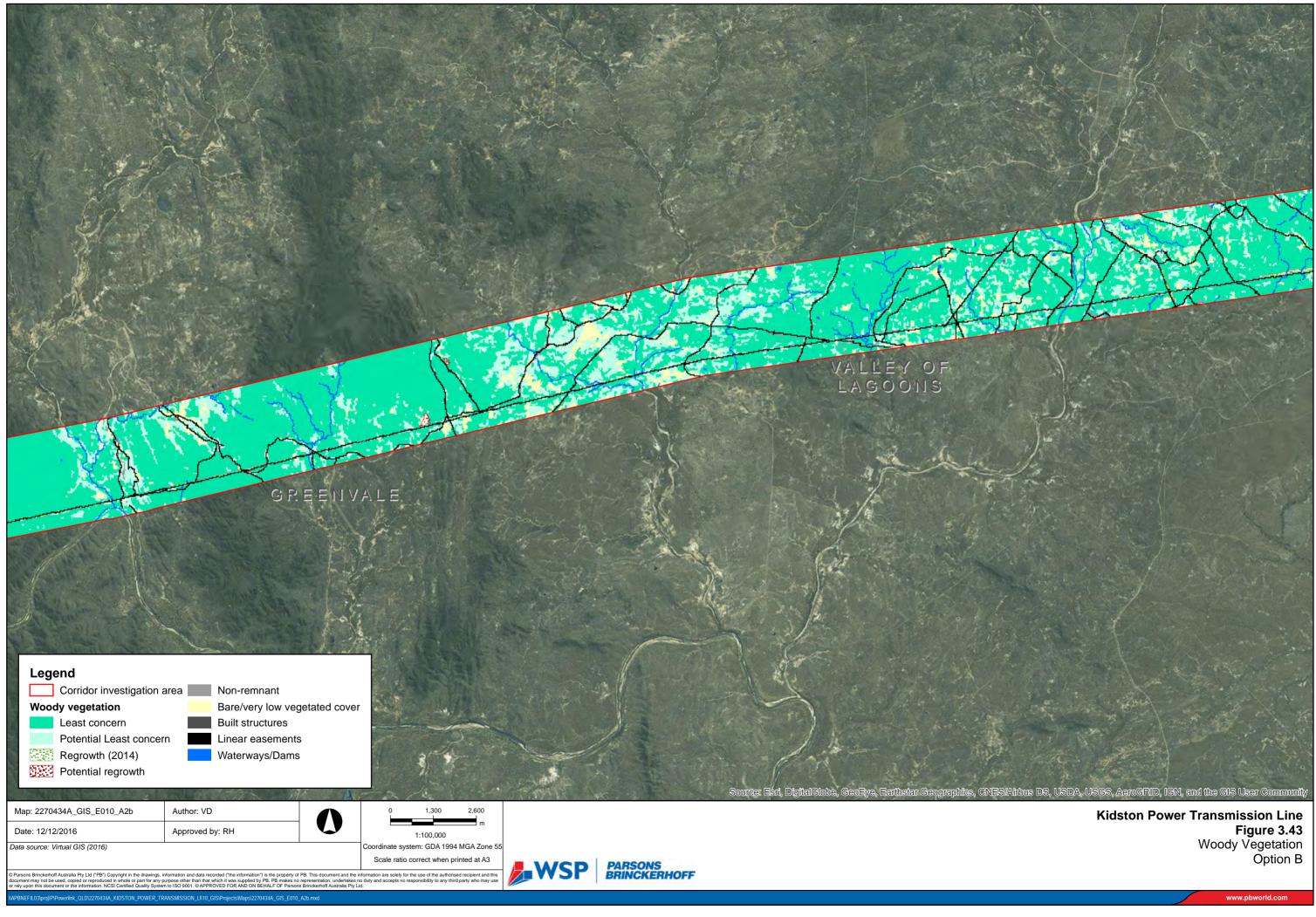


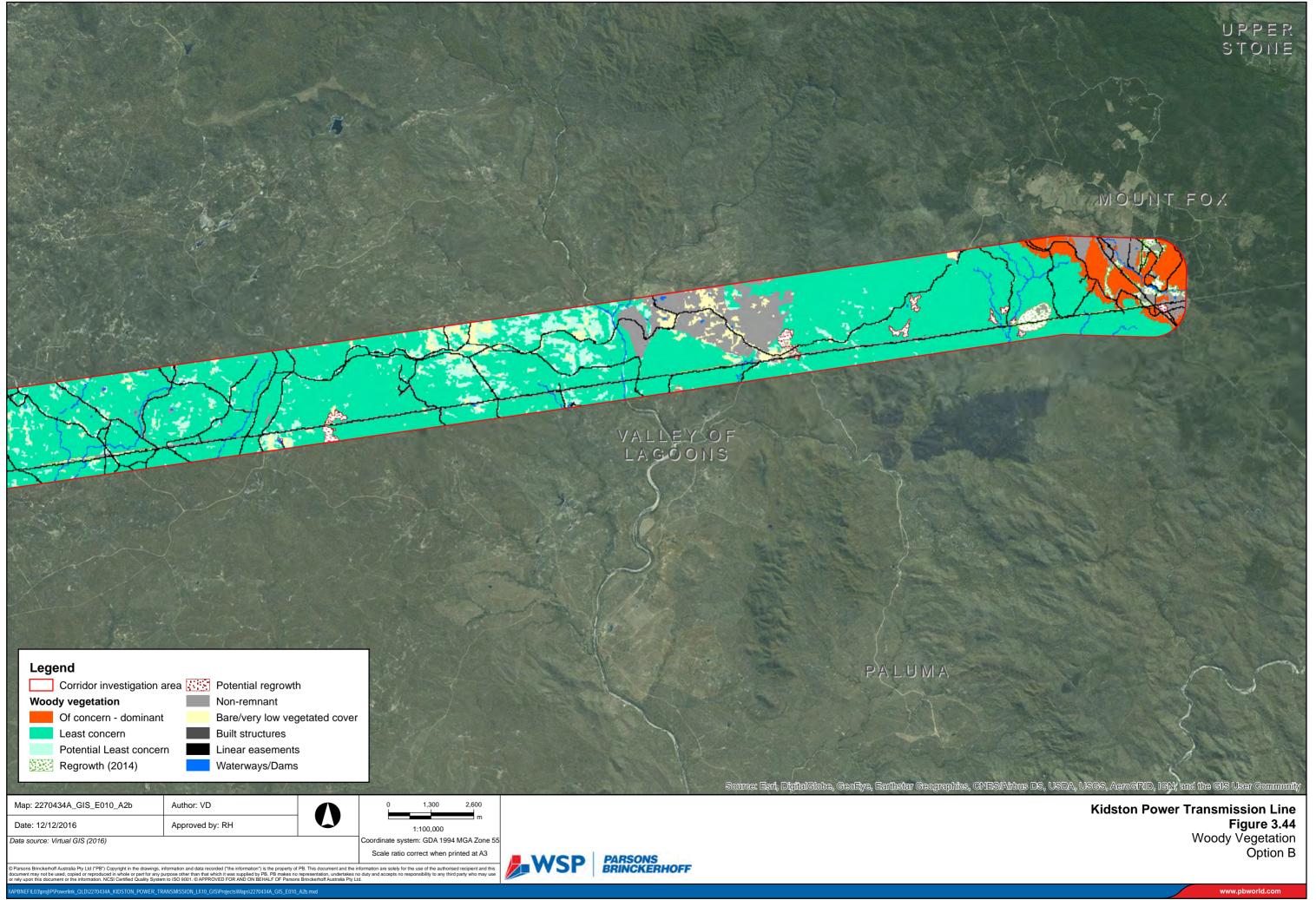
Kidston Power Transmission Line Figure 3.39 Woody Vegetation Option A

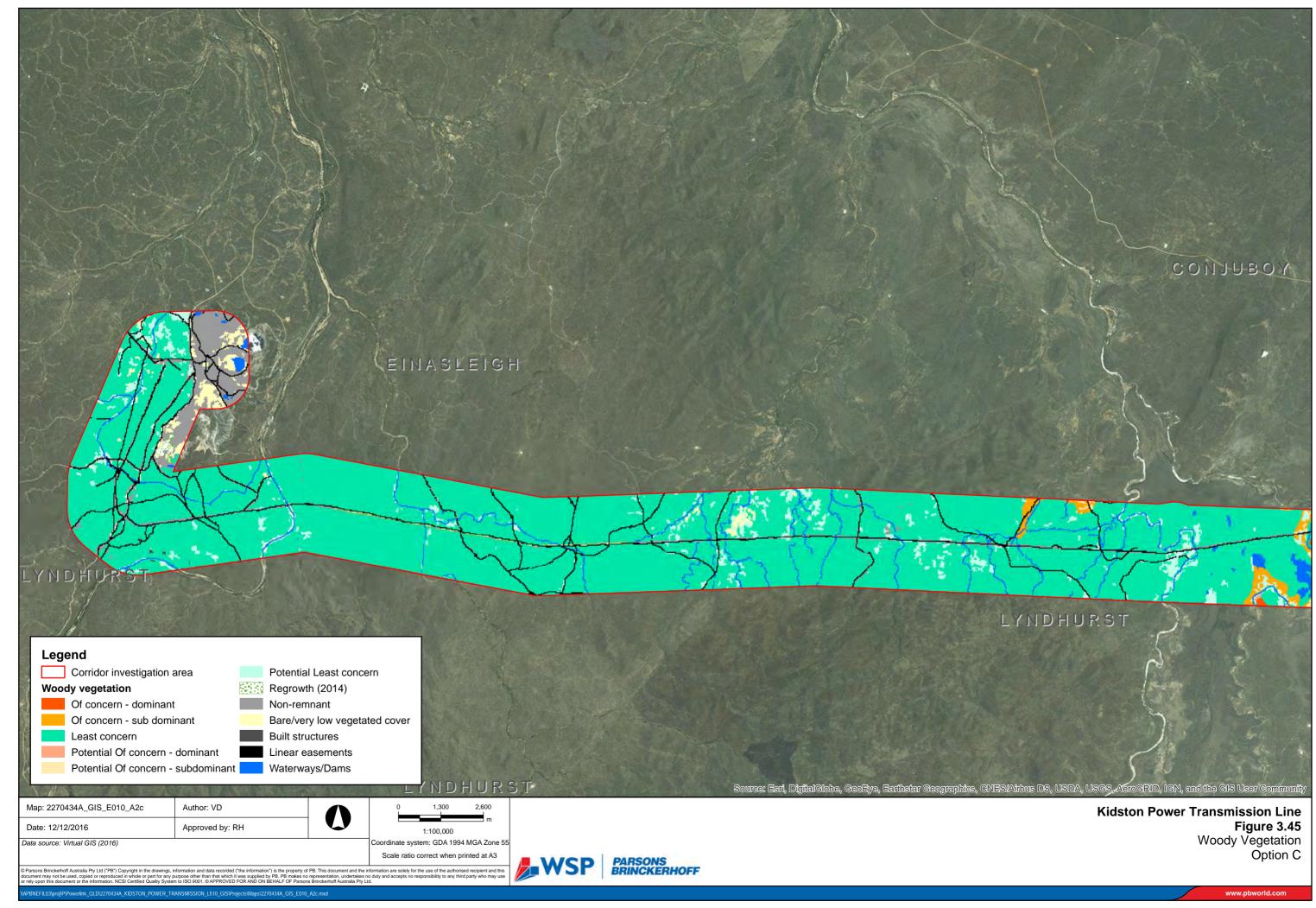


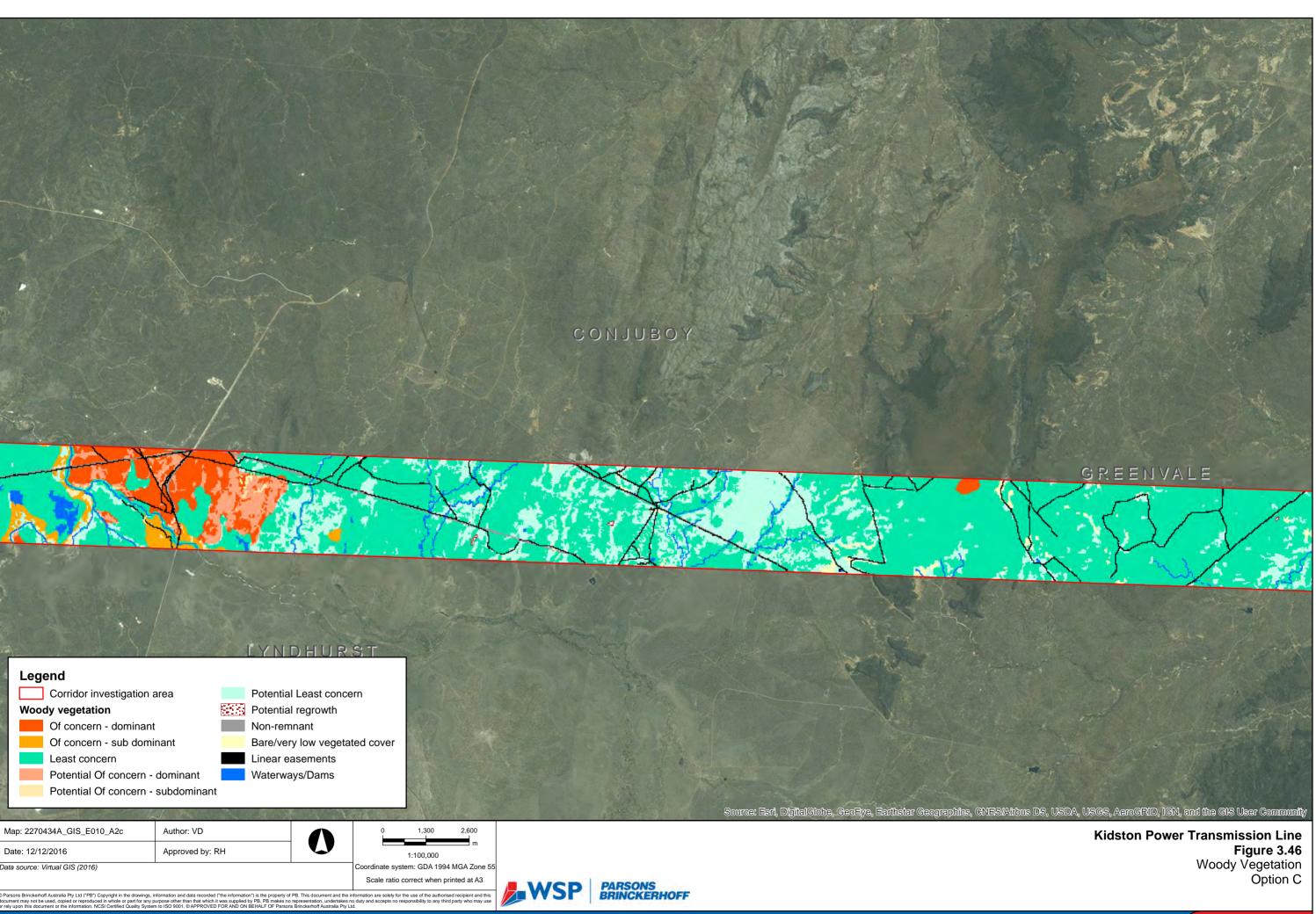






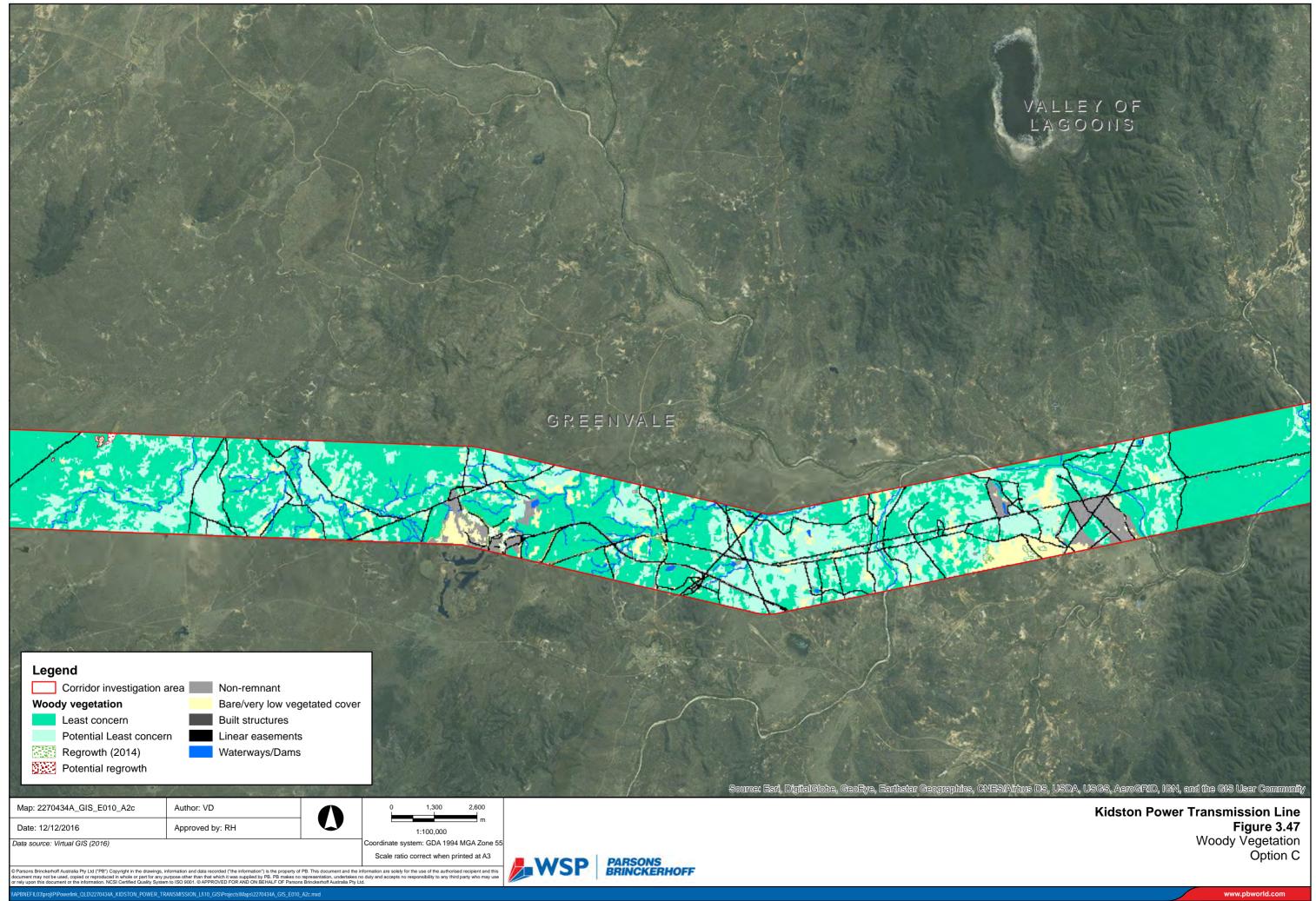


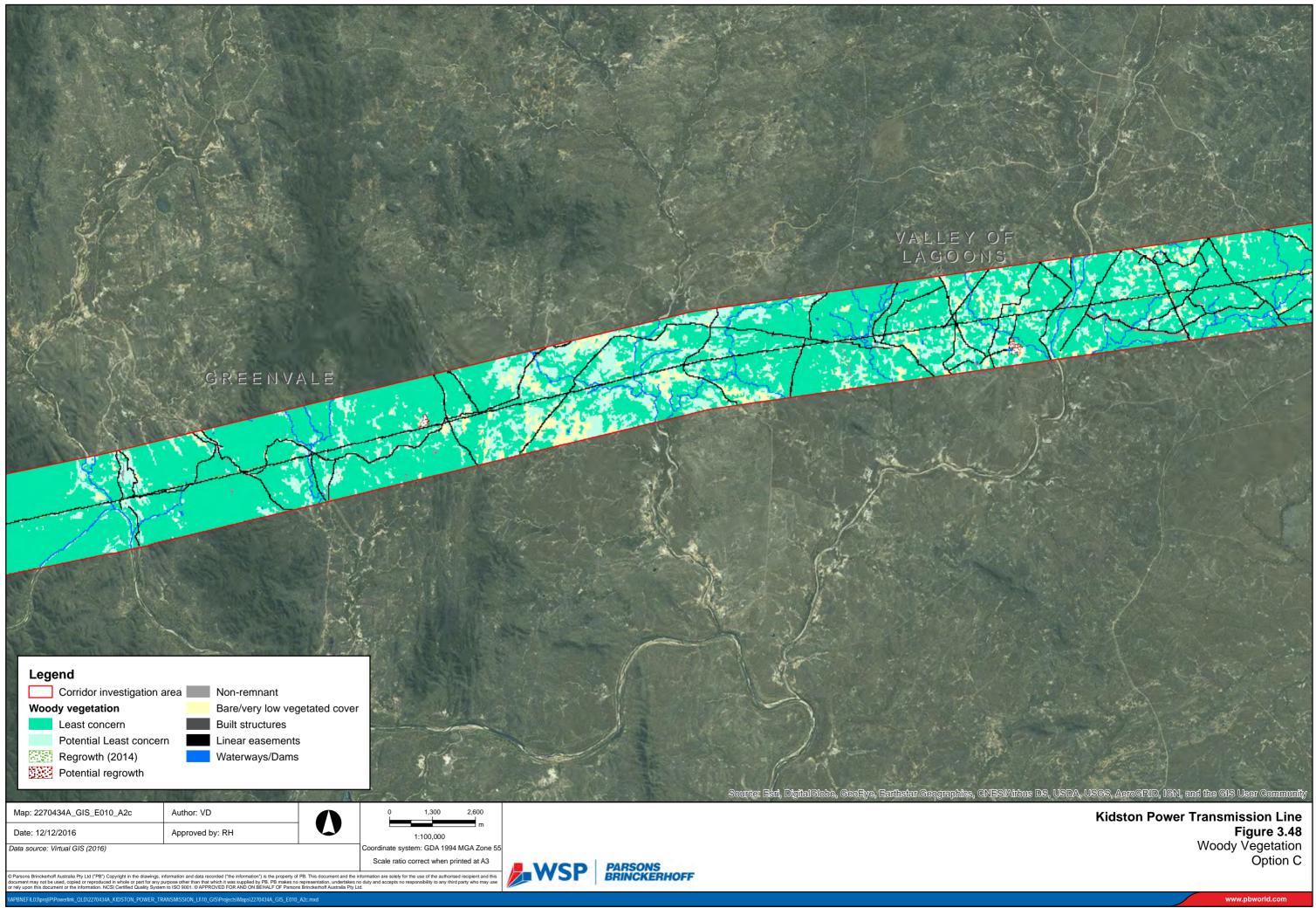


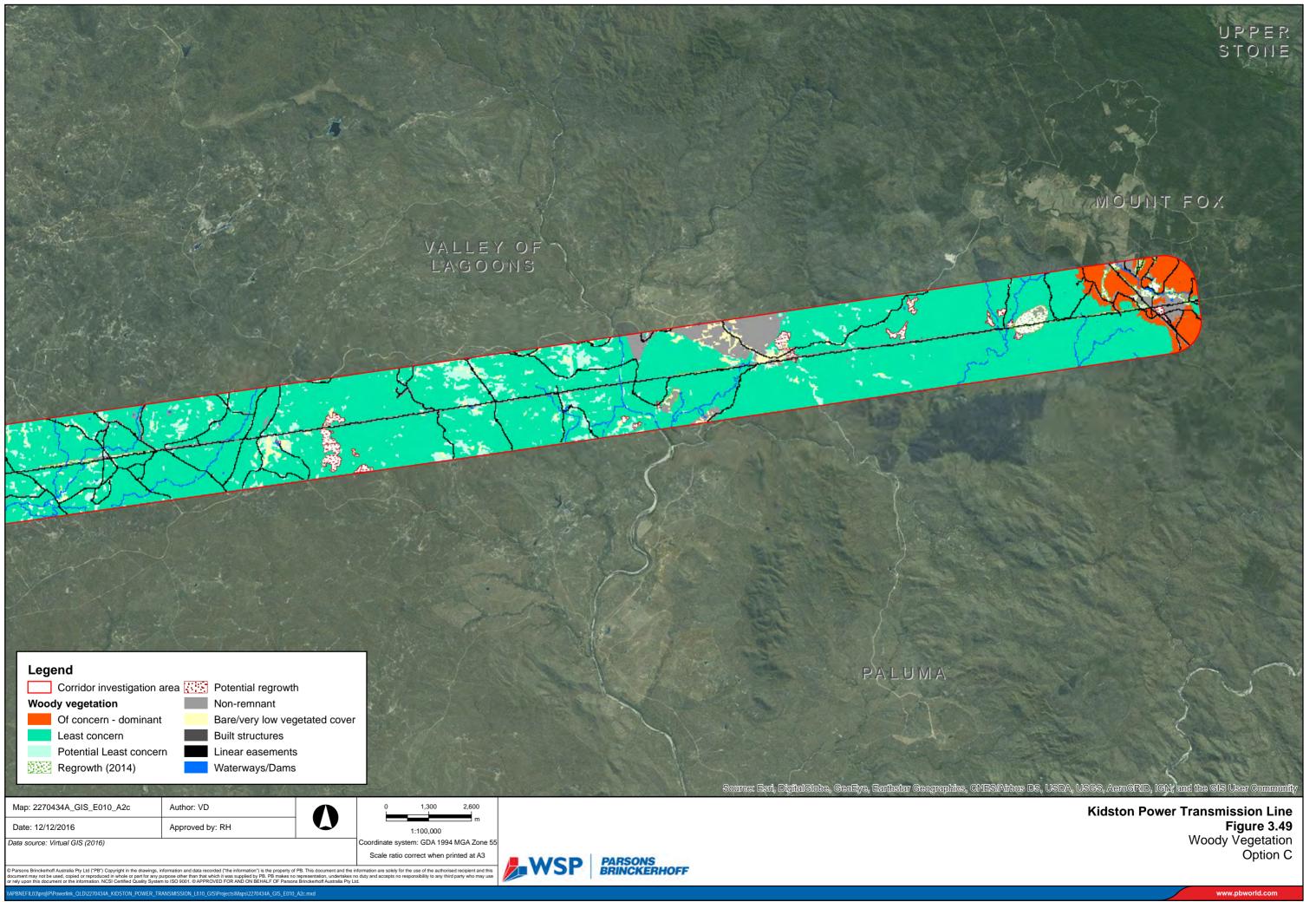


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#### 3.1.5 Regulated remnant watercourse vegetation

All three corridor investigation areas intersect remnant watercourse vegetation on the 1:100,000 and 1:250,000 scale vegetation management watercourse maps, as indicated above on Figure 3.20–Figure 3.34.

The eastern third of Options A, B and C are located in the coastal Wet Tropics bioregion and for those part of each option, the defined distances for a remnant regional ecosystem in a coastal bioregion applies (refer Table 2.1). The remaining two-thirds of Options A, B and C are located in the non-coastal Einasleigh Uplands bioregion and therefore are aligned to the defined distances for a remnant regional ecosystem in a non-coastal bioregion (refer Table 2.2).

The area (ha) of potential impact of each corridor option upon regulated remnant watercourse vegetation in the coastal Wet Tropics bioregion is presented in Table 3.6.

The results of the analysis of regulated remnant watercourse vegetation for each corridor option is presented in Appendix G.

Table 3.6	The extent (ha) of potential impact of each corridor option upon regulated remnant watercourse
	vegetation in the coastal Wet Tropics bioregion

WATERCOURSE STREAM ORDER	DISTANCE FROM THE DEFINING BANK (M)	OPTION A (HA)	OPTION B (HA)	OPTION C (HA)
1 or 2	10	65	30	23
3 or 4	25	53	20	22
5 or greater	50	0	0	0
Totals		118	50	45

In reference to Table 3.6, Option C is of least constraint in terms of potential impacts upon regulated remnant watercourse vegetation within the defined distances of watercourses in the coastal Wet Tropics bioregion.

The extent (ha) of potential impact of each corridor option upon regulated remnant watercourse vegetation in the non-coastal Einasleigh Uplands bioregion, is presented in Table 3.7.

## Table 3.7The extent (ha) of potential impact of each corridor option upon regulated remnant watercourse<br/>vegetation in the non-coastal Einasleigh Uplands bioregion

WATERCOURSE STREAM ORDER	DISTANCE FROM THE DEFINING BANK (M)	OPTION A (HA)	OPTION B (HA)	OPTION C (HA)
1 or 2	25	3,619	3,518	3,511
3 or 4	50	1,301	1,325	1,379
5 or greater	100	665	704	603
Totals		5,585	5,547	5,493

In reference to Table 3.7, Option C is of least constraint in terms of potential impacts upon regulated remnant watercourse vegetation within the defined distances of watercourses in the non-coastal Einasleigh Uplands bioregion.

#### 3.1.6 Regulated remnant wetland vegetation

All three corridor investigation areas intersect remnant wetland vegetation, as indicated above on Figure 3.20–Figure 3.34.

The area (ha) of potential impact of each corridor option upon regulated remnant wetland vegetation is presented in Table 3.8.

The results of the analysis of regulated remnant watercourse vegetation for each corridor option is presented in Appendix G.

In reference to Table 3.8, Option A is of least constraint in terms of potential impacts upon regulated remnant wetland vegetation that is within a 50 m of the defined bank of mapped regulated vegetation wetlands.

Table 3.8The extent (ha) of potential impact of each corridor option upon regulated remnant wetland<br/>vegetation

CORRIDOR	<b>REGULATED REMNANT WETLAND VEGETATION (HA)</b>
Option A	4.7
Option B	42.3
Option C	27.7

#### 3.1.7 Protected areas

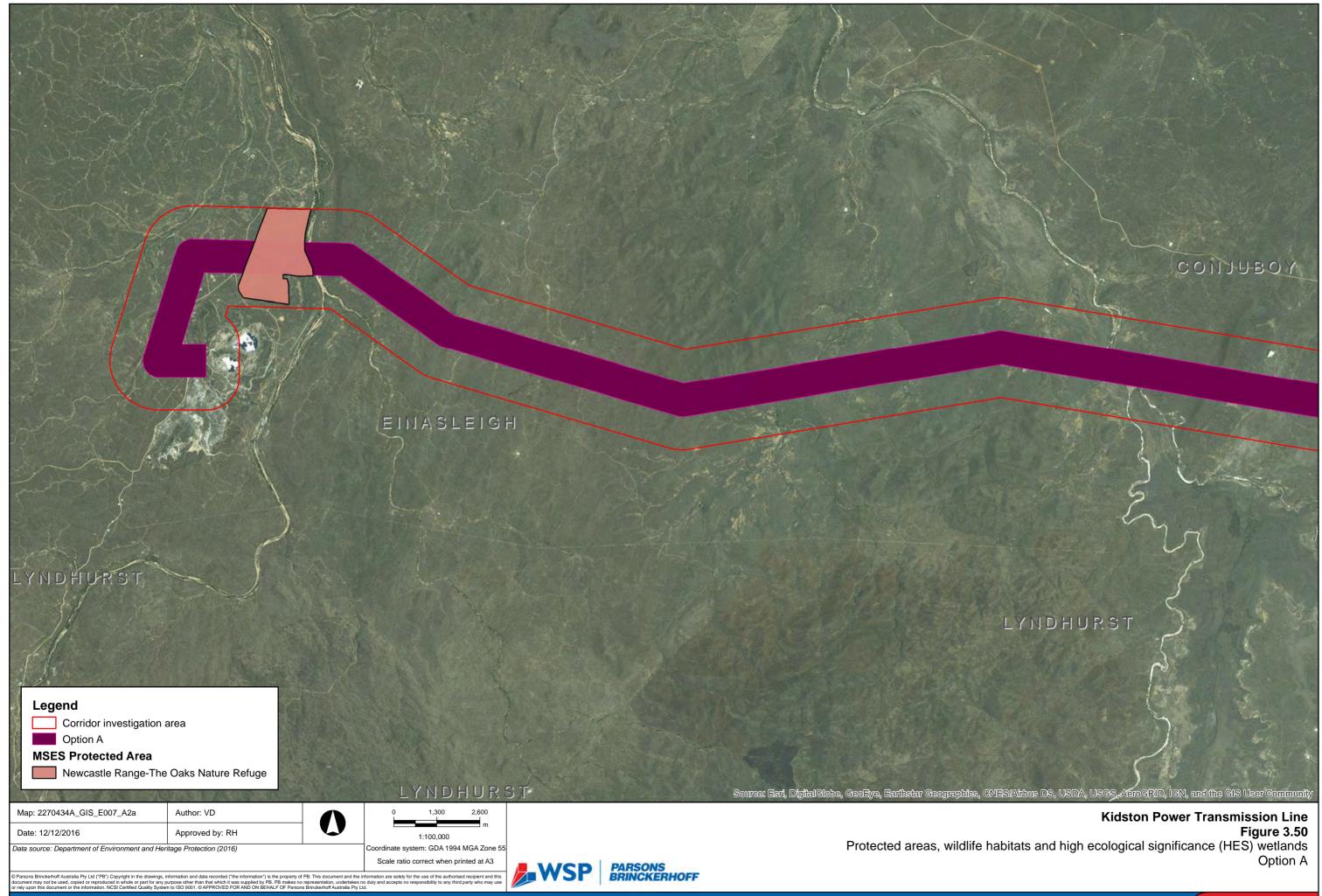
All three corridor investigation areas intersect with sections of protected areas, as indicated on Figure 3.50–Figure 3.64. The area (ha) of protected areas associated with each corridor investigation area is presented in Table 3.9.

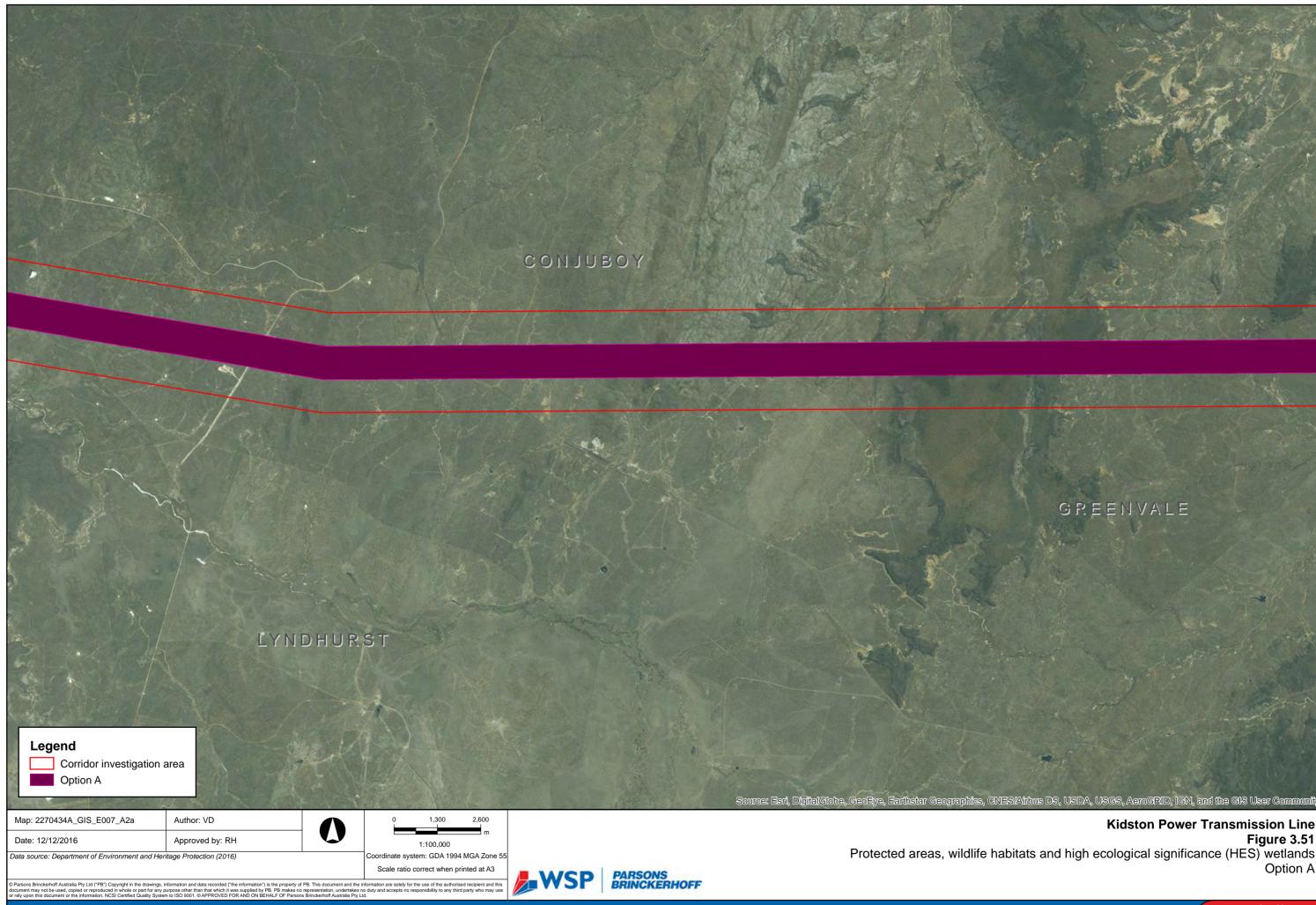
The spreadsheet that contains the analysis of protected areas is presented in Appendix G.

In reference to Table 3.9, Option C is of least constraint in terms of potential impacts upon MSES Protected Areas.

#### Table 3.9 Amount (ha) of protected areas associated with each corridor investigation area

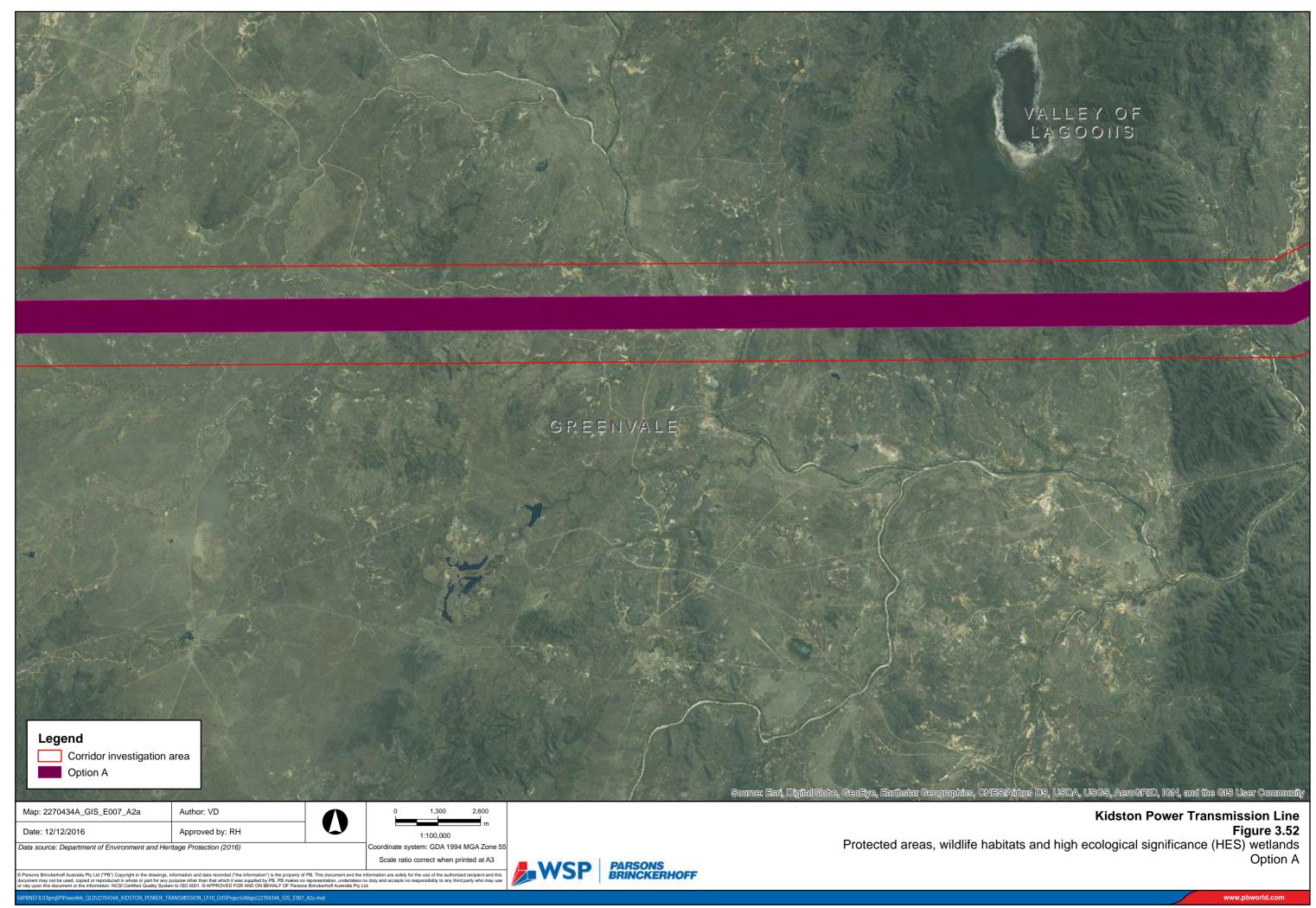
PROTECTED AREA	OPTION A (HA)	OPTION B (HA)	OPTION C (HA)
Newcastle Range – The Oaks Nature Refuge	439.8	0	0
Liefway Nature Refuge	4.7	0	3.0
Sub-totals – Nature Refuges	444.5	0	3.0
Girringun National Park	213.7	214.4	142.3
Sub-total – National Parks	213.7	214.4	142.3
Total – Protected Areas	658	214	145

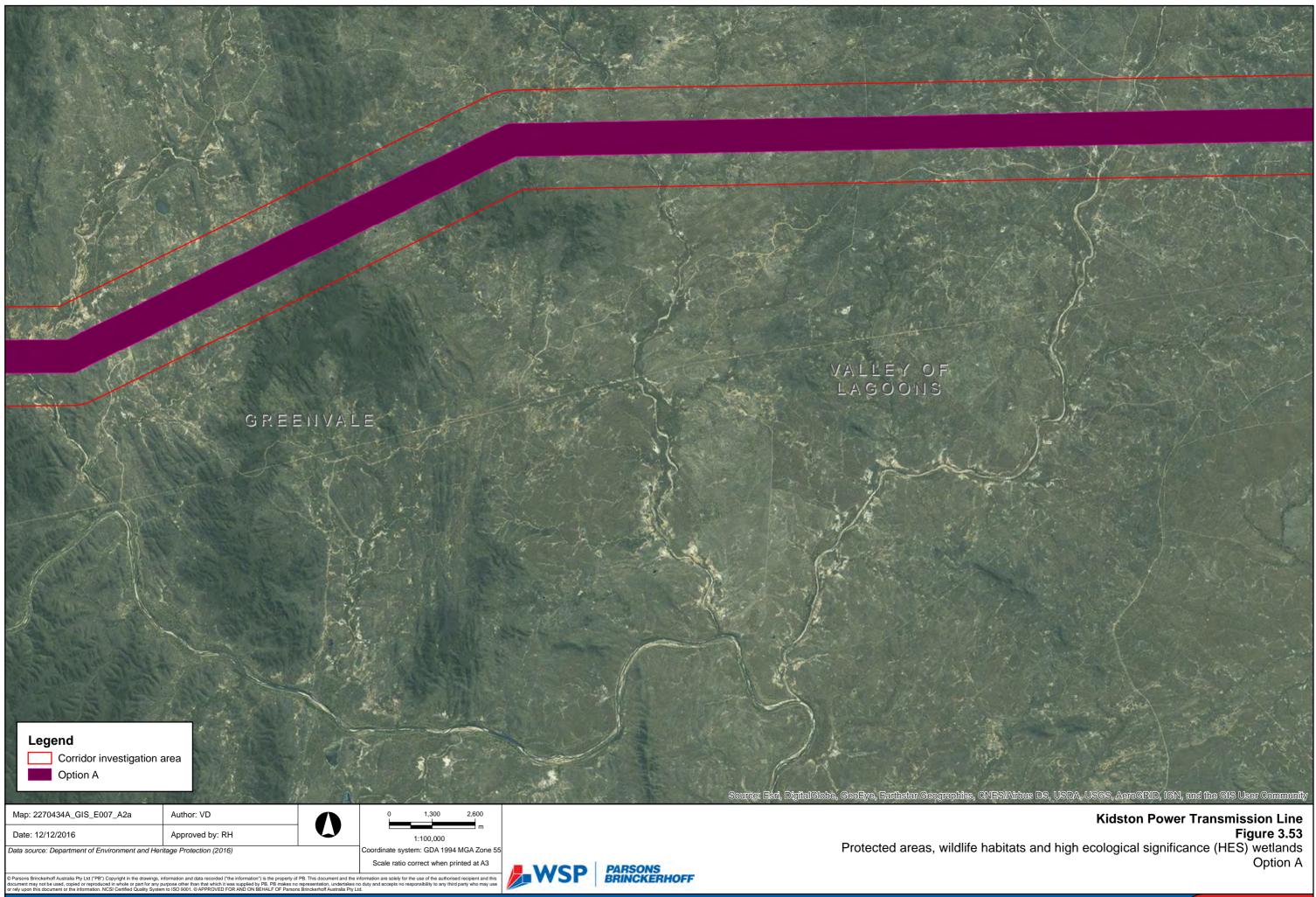


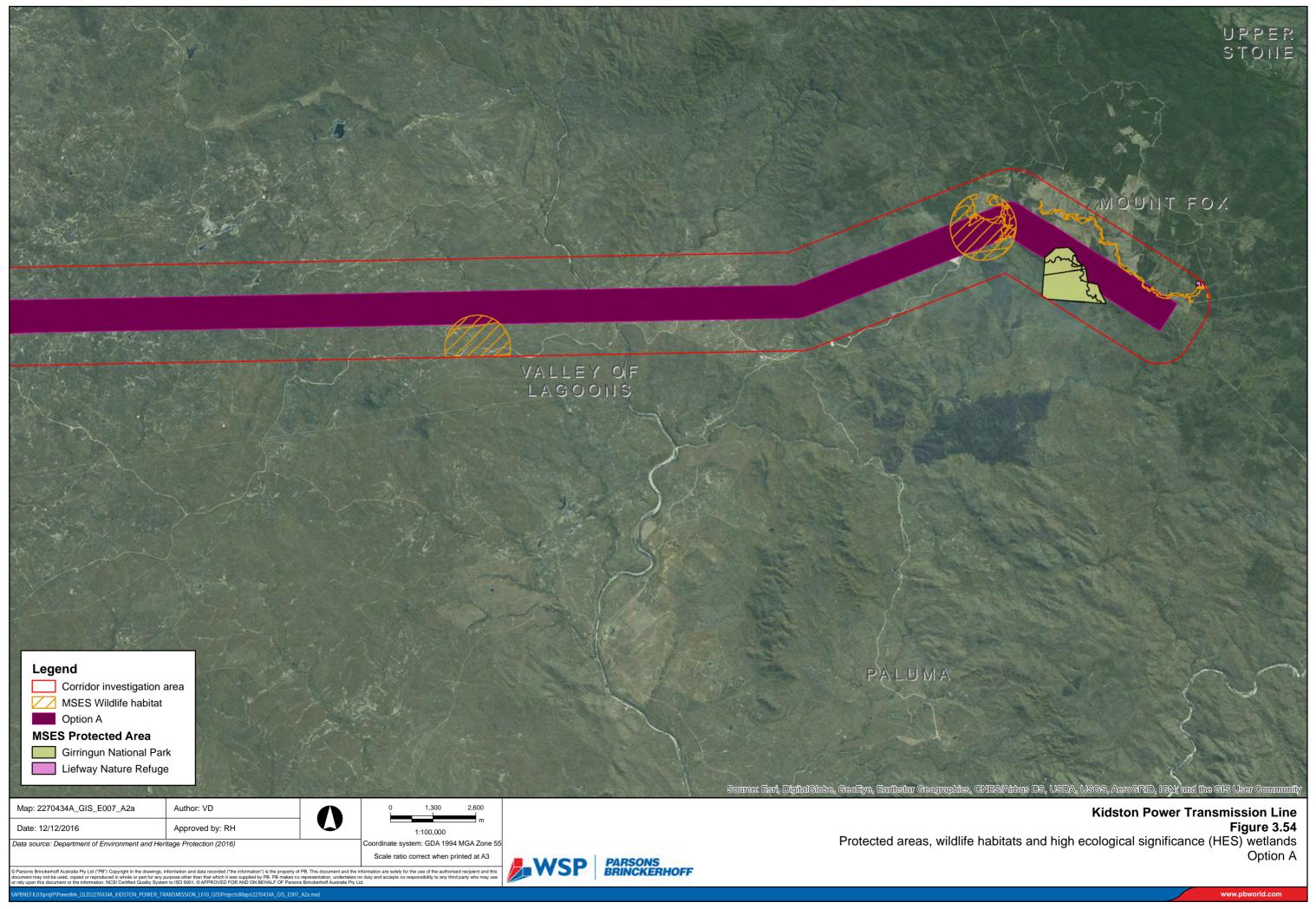


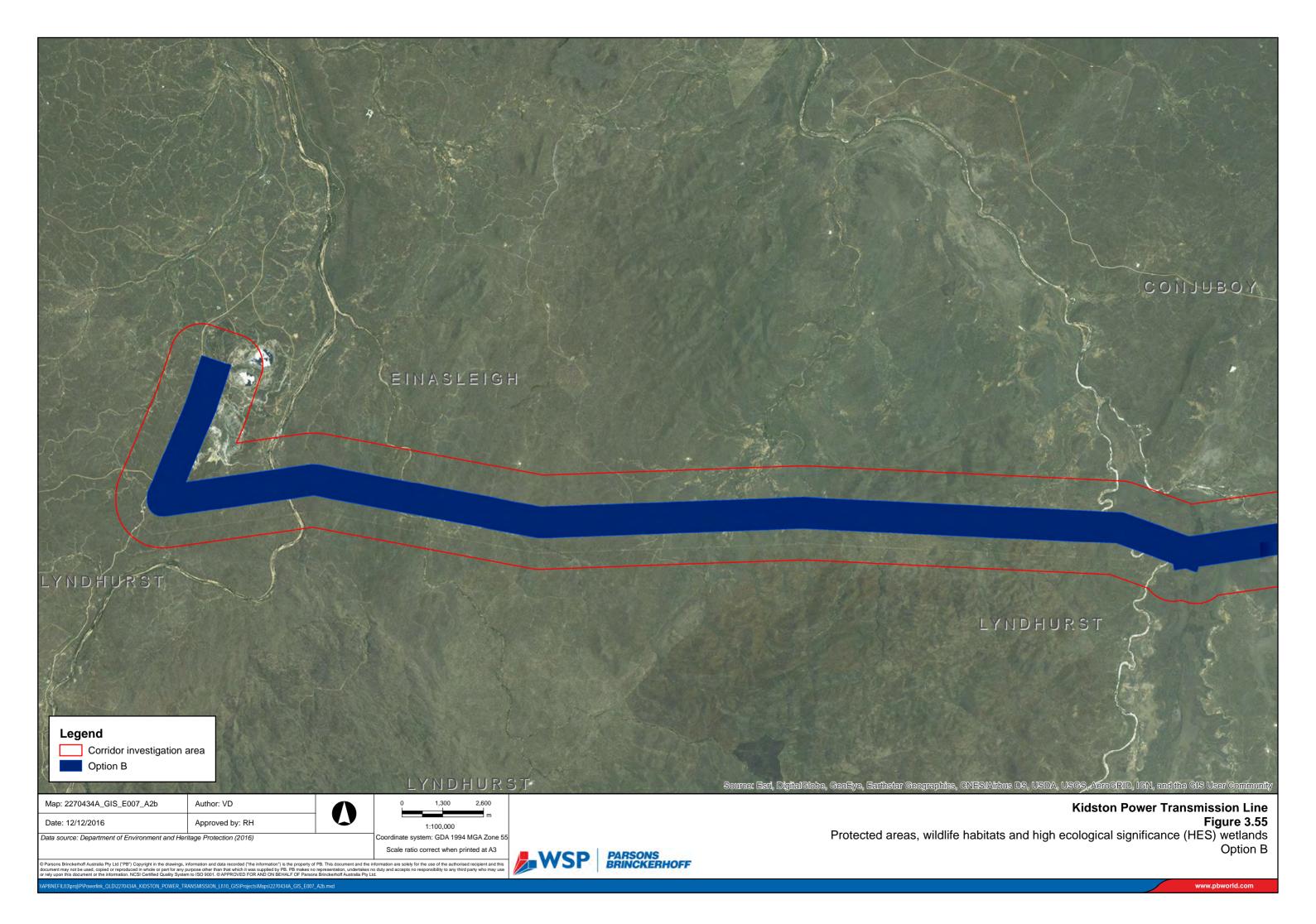


Kidston Power Transmission Line Figure 3.51 Protected areas, wildlife habitats and high ecological significance (HES) wetlands Option A





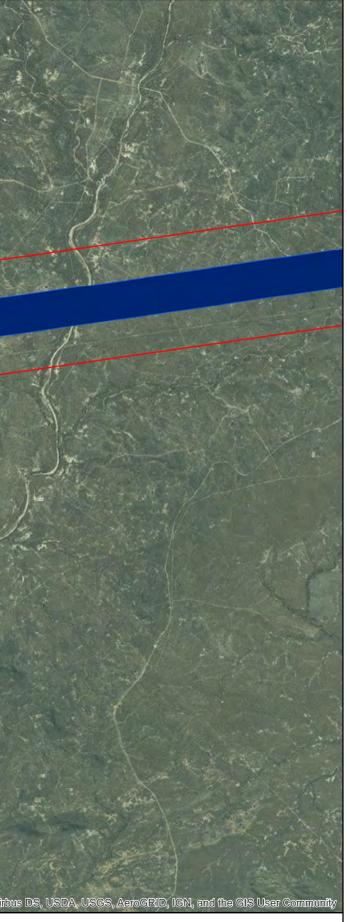




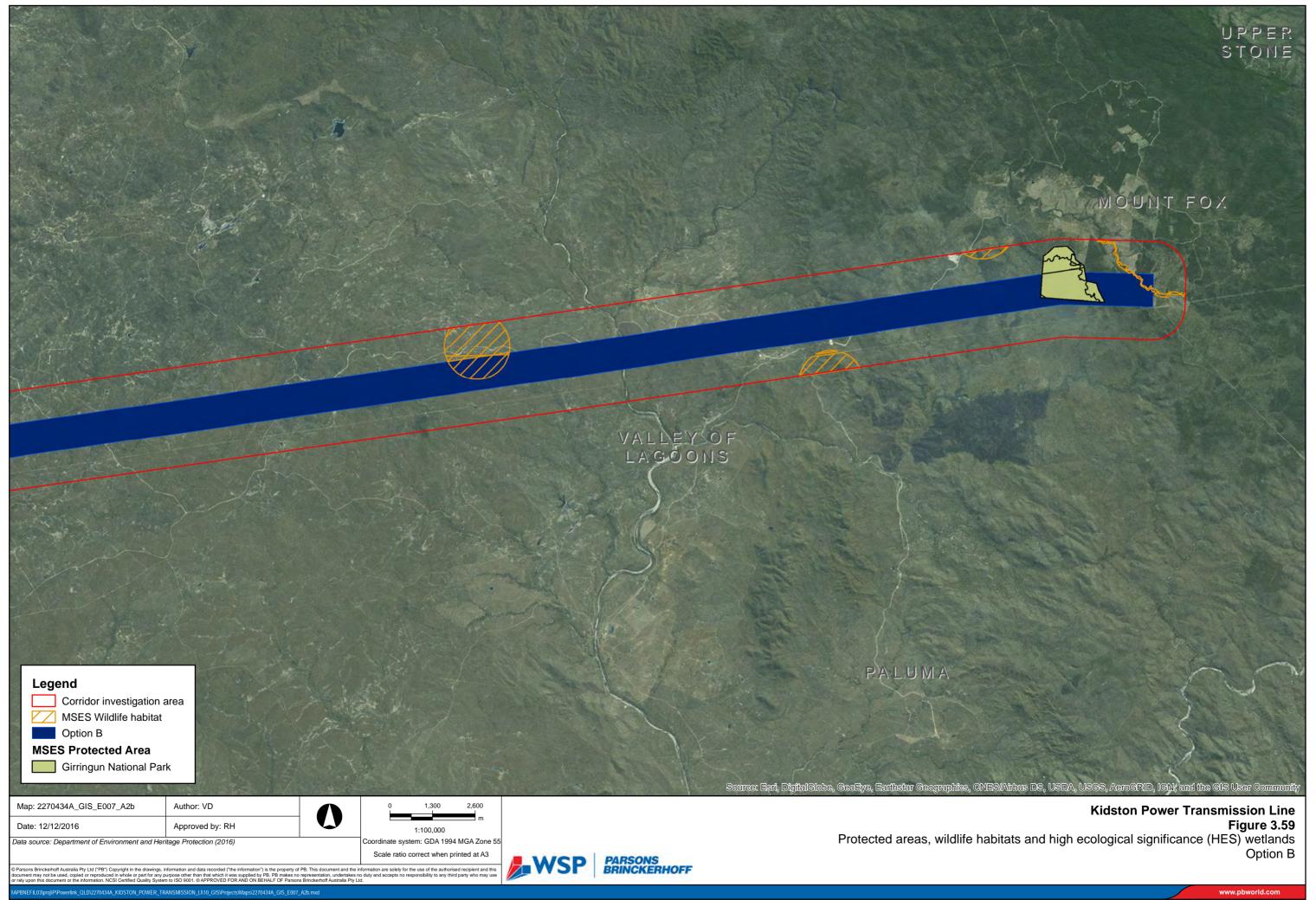


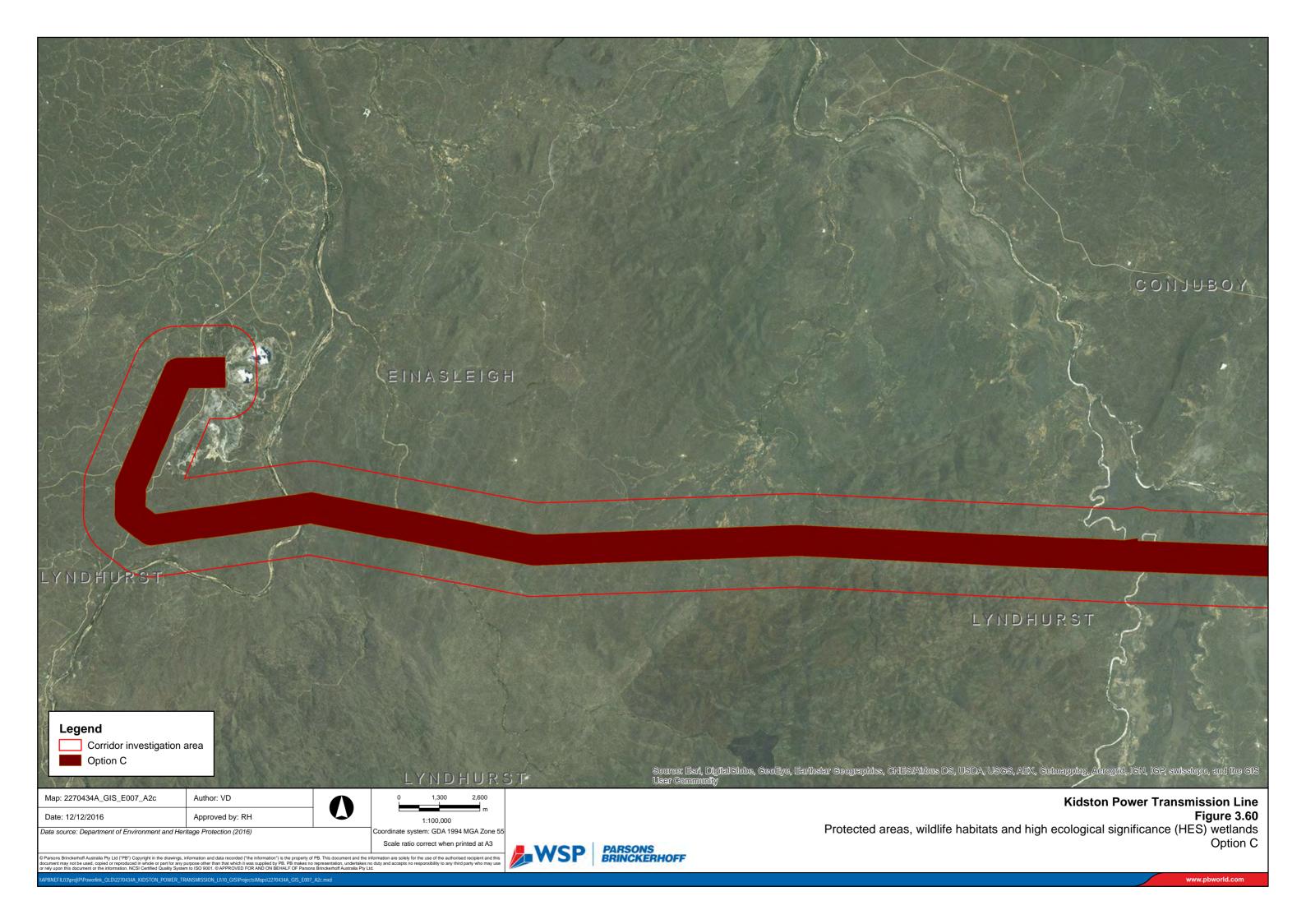


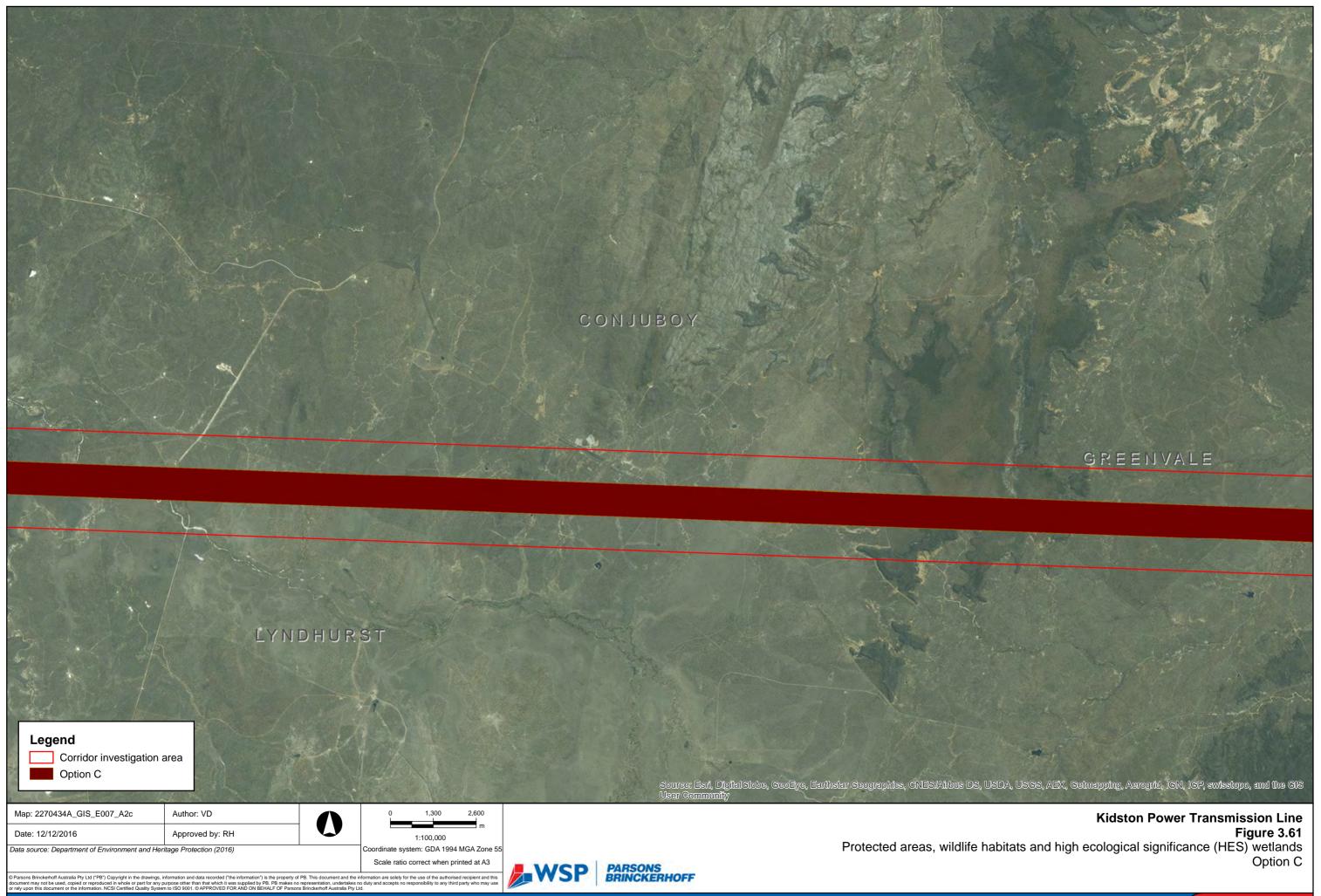
				VALLEN LAGOO	OF DNS
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Legend Corridor investigation Option B	n area				
Map: 2270434A_GIS_E007_A2b Date: 12/12/2016 Data source: Department of Environment and H © Parsons Brinckerhoff Australia Pty Ltd (*PB*) Copyright in the drawing document may not be used, copied or reproduced in whole or part for a or rely upon this document or the information. NCSI Certified Quality Sy	Author: VD Approved by: RH leritage Protection (2016) s, information and data recorded ('the information') is the property of PB. This document and ry purpose other than that which it was supplied by PB. PB makes no representation, undertail stem to ISO 9001. © APPROVED FOR AND ON BEHALF OF Parsons Brinckerholf Australia f	0 1,300 2,600 m 1:100,000 Coordinate system: GDA 1994 MGA Zone 55 Scale ratio correct when printed at A3 the information are solely for the use of the authorised recipient and this kees no duty and accepts no responsibility to any third party who may use Pty Ltd.	WSP PARSONS BRINCKERHOFF	Source: Esri, DigitalGlobe, GeoEye, Earths Protected a	tar Geographies, CNES#

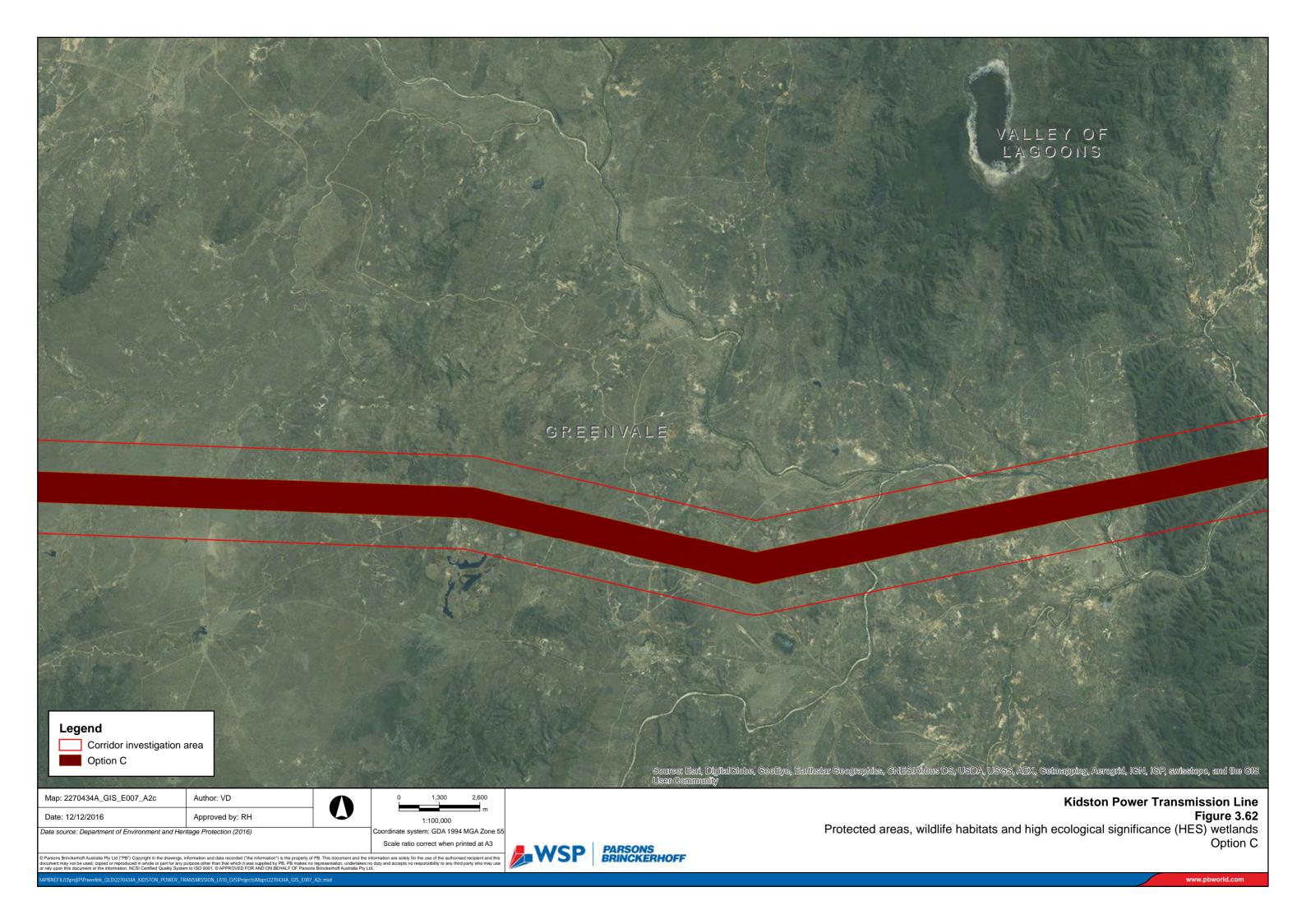


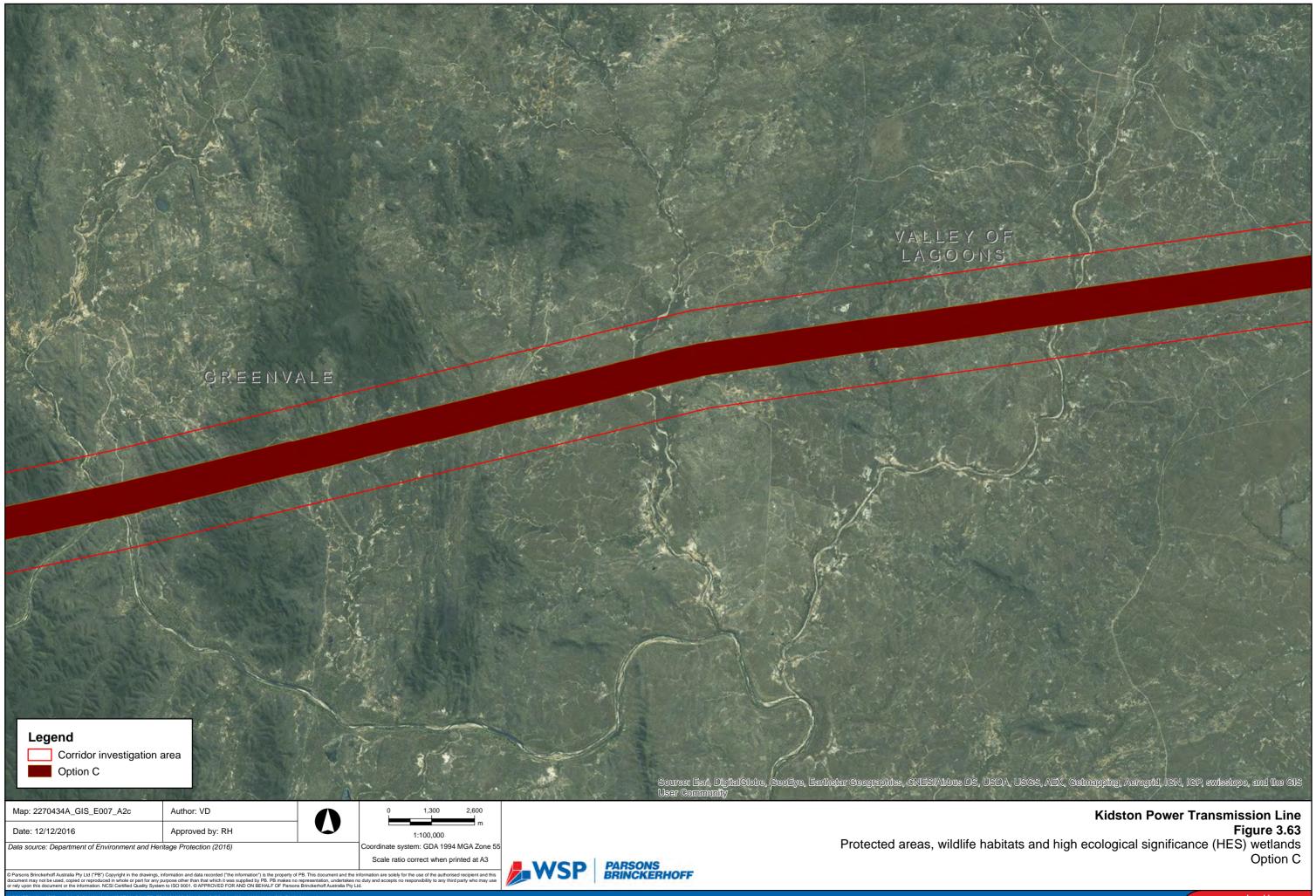
Kidston Power Transmission Line Figure 3.58 tats and high ecological significance (HES) wetlands Option B

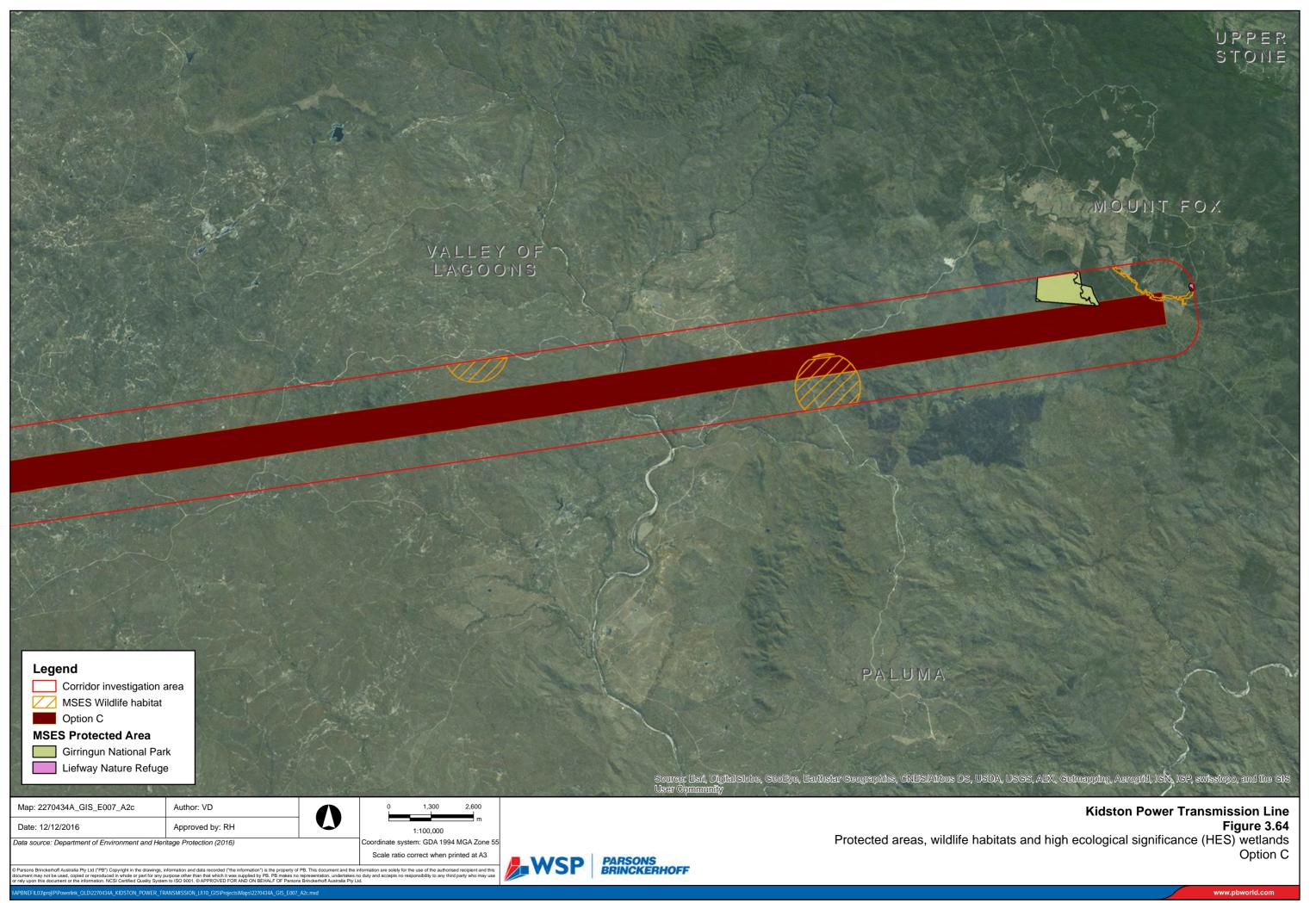












#### 3.1.8 Wildlife habitats

All three corridor investigation areas intersect mapped wildlife habitats, as indicated on Figure 3.50– Figure 3.64. The amount (ha) protected areas associated with each corridor investigation area is presented in Table 3.10. The spreadsheet that contains the analysis of wildlife habitat is presented in Appendix G.

In reference to Table 3.10, Option C is of least constraint in terms of potential impacts upon MSES Wildlife Habitat.

 Table 3.10
 Amount (ha) of wildlife habitats associated with each corridor investigation area

CORRIDOR	WILDLIFE HABITATS (HA)
Option A	522
Option B	411
Option C	385

#### 3.1.9 Strategic environmental areas

Strategic environmental areas are not relevant to any of the three corridor options.

#### 3.1.10 High ecological significance (HES) wetlands

Of the three corridor investigation areas, only Option B intersects high ecological significance wetlands, as indicated on Figure 3.50–Figure 3.64. The area (ha) of high ecological significance wetlands associated with each corridor investigation area is presented in Table 3.11. The results of the analysis of HES wetlands is presented in Appendix G.

CORRIDOR	HIGH ECOLOGICAL SIGNIFICANCE WETLANDS (HA)
Option A	0
Option B	1.1
Option C	0

#### 3.1.11 High ecological value waters (wetland)

High ecological value waters (wetland) are not relevant to any of the three corridor options.

#### 3.1.12 High ecological value waters (watercourses)

High ecological value waters (watercourses) are not relevant to any of the three corridor options.

#### 3.1.13 Environmental offset areas

Environmental offset areas are not relevant to any of the three corridor options.

#### 3.1.14 Connectivity

Connectivity is assessed significant residual impacts under the Queensland Government's *Environmental Offsets Policy version 1.1* (2014), once a project footprint has been determined. This impact is assed using GIS and the Queensland Government's *Landscape Fragmentation and Connectivity tool* for an individual impact area. Essentially, the assessment is based upon the level of fragmentation to remnant vegetation as a result of a project's impact area(s).

In reference to the amount of remnant vegetation that occurs within each corridor investigation area, all three corridor options would result in impacts to connectivity. On this basis once a preferred corridor option is selected and project impact area(s) are defined, the potential impact upon connectivity would need to be assessed using the Landscape Fragmentation and Connectivity tool.

#### 3.2 Matters of National Environmental Significance

The following MNES have been assessed in the following sections:

- → threatened species and migratory species
- → threatened ecological communities
- → heritage properties of international and national significance.

#### 3.2.1 EPBC Act Protected Matters Search Tool results

The PMST database search results revealed the MNES of potential relevance to the corridor options, including one TEC, 33 threatened species and 15 migratory species. The search also identified world, national and indigenous heritage properties.

The results of the PMST database search for each corridor investigation area is presented in Table 3.12. The complete PMST reports for each corridor option are presented in Appendix A.

MNES	SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	OPTION A	OPTION B	OPTION C
TECs	Broad leaf tea-tree ( <i>Mela</i> woodlands in high rainfall Queensland		E	-	х	х	х
Total					1	1	1
Threatened	l species						
Birds	Calidris ferruginea	Curlew Sandpiper	CE (M)	SL	х	х	х
	Casuarius casuarius johnsonii	Southern Cassowary	E	E	х	х	х
	Erythrotriorchis radiatus	Red Goshawk	V	E	х	х	х
	Erythrura gouldiae	Gouldian Finch	E	E	х	х	х
	Numenius madagascariensis	Eastern Curlew	CE	V	х	х	х
	Poephila cincta cincta	Southern Black- throated Finch	E	E	х	х	х
	Rostratula australis	Australian Painted Snipe	E	V	х	х	х
	Tyto novaehollandiae kimberli	Masked Owl (northern)	V	V	х	х	х
Frogs	Litoria dayi	Australian Lace-lid	E	E	х	х	х
	Litoria nannotis	Waterfall Frog	E	Е	х	х	х
	Litoria rheocola	Common Mistfrog	E	E	х	х	х

MNES	SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	OPTION A	OPTION B	OPTION C
Mammals	Dasyurus hallucatus	Northern Quoll	E	-	х	х	х
	Dasyurus maculatus gracilis	Spotted-tailed Quoll (north Queensland)	E	E	х	х	х
	Hipposideros semoni	Semon's Leaf-nosed Bat	E	E	х	х	х
	Macroderma gigas	Ghost Bat	V	V	х	х	x
	Mesembriomys gouldii rattoides	Black-footed Tree-rat (north Queensland)	V	-	х	х	х
	Petauroides volans	Greater Glider	V	-	х	х	х
	Petrogale sharmani	Sharman's Rock- wallaby	V	V	х	х	х
	Phascolarctos cinereus	Koala	V	V	х	х	х
	Pteropus conspicillatus	Spectacled Flying-fox	V	V	х	х	х
	Pteropus poliocephalus	Grey-headed Flying- fox	V	-	х	х	х
	Rhinolophus robertsi	Large-eared Horseshoe Bat	E	E	х	х	x
	Saccolaimus saccolaimus nudicluniatus	Bare-rumped Sheath- tailed Bat	CE	E	x	x	x
Plants	Acacia crombiei	Pink Gidgee	V	V	х	х	x
	Bulbophyllum globuliforme	Miniature Moss- orchid	V	NT	х	х	х
	Cajanus mareebensis	-	E	Е	х	х	х
	Dichanthium setosum	Bluegrass	V	-	х	х	х
	Lindsaea pulchella var. blanda	-	V	EX	х	х	x
	Marsdenia brevifolia	-	V	V	х	х	х
	Phaius australis	Lesser Swamp- orchid	E	E	х	х	х
	Tephrosia leveillei	-	V	V	х	х	х
Reptiles	Egernia rugosa	Yakka Skink	V	V	х	х	х
	Lerista vittata	Mount Cooper Striped Lerista	V	V	х	х	х
Migratory (marine)	Apus pacificus	Fork-tailed Swift	М	SL	х	х	х
Migratory	Cuculus optatus	Oriental Cuckoo	М	SL	х	х	х
(terrestrial)	Hirundapus caudacutus	White-throated Needletail	М	SL	х	х	х

MNES	SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	OPTION A	OPTION B	OPTION C
	Hirundo rustica	Barn Swallow	М	SL	х	х	х
	Monarcha melanopsis	Black-faced Monarch	М	SL	х	х	х
	Monarcha trivirgatus	Spectacled Monarch	М	SL	х	х	х
	Motacilla cinerea	Grey Wagtail	М	SL	х	х	х
	Motacilla flava	Yellow Wagtail	М	SL	х	х	х
	Myiagra cyanoleuca	Satin Flycatcher	М	SL	х	х	х
	Rhipidura rufifrons	Rufous Fantail	М	SL	х	х	х
Migratory	Calidris ferruginea	Curlew Sandpiper	M (CE)	SL	х	х	х
(wetland)	Gallinago hardwickii	Latham's Snipe	М	SL	х	х	х
	Numenius madagascariensis	Eastern Curlew	M (CE)	SL	х	х	х
	Pandion haliaetus	Osprey	М	SL	х	х	х
	Tringa nebularia	Common Greenshank	М	SL	-	х	х
Totals	Totals				47	48	48
Other MNES							
World Heritage properties - Wet Tropics of Queensland				х	-	-	
National Heritage properties (natural) - Wet Tropics of Queensland				х	-	-	
National Heritage properties (indigenous) - Wet Tropics World Heritage Area (Indigenous Values)				х	-	-	

Key: EX = extinct in the wild, CE = critically endangered, E = endangered, V = vulnerable, NT = near threatened, M = migratory, SL = special least concern

#### 3.2.2 EPBC Act threatened and migratory species records

The EPBC Act threatened and migratory species records obtained from DSITI's species profile search and Atlas of Living Australia that are of relevance to each corridor option have been quantified and mapped using GIS. The number of threatened and/or migratory species listed under the EPBC Act that have been previously recorded within each corridor investigation area is presented in Table 3.13 and mapped on Figure 3.1–Figure 3.3.

The results of the analysis of threatened species records is presented in Appendix F.

In reference to Table 3.13, Option A has the least number of EPBC Act threatened species records. However, as mentioned above, threatened species records are representative of the flora and fauna surveys that are undertaken in specific areas for developments or in association with protected areas. In respect to the remoteness of the corridor investigation areas, targeted surveys for the Project are likely to identify several threatened species that have not yet been recorded and formerly on the DSITI and ALA records databases. 
 Table 3.13
 The number of EPBC Act listed species that have been previously recorded within each corridor investigation area

EPBC ACT STATUS	OPTION A	OPTION B	<b>OPTION C</b>			
Threatened flora species						
Critically endangered	-	_	-			
Endangered	-	-	-			
Vulnerable	1	1	1			
Threatened fauna species	Threatened fauna species					
Critically endangered	-	1	1			
Endangered	1	1	1			
Vulnerable	1	2	2			
Migratory fauna species						
Migratory (marine)	-	1	-			
Migratory (terrestrial)	-	-	-			
Migratory (wetland)	-	_	_			
Total records	3	6	5			

#### 3.2.3 Threatened ecological communities

Broad Leaf Tea-tree (Melaleuca viridiflora) woodlands in high rainfall coastal north Queensland is the only TEC that was revealed by the PMST as potentially occurring within corridor option investigation areas. A map of the predicted distribution of the Broad Leaf Tea-tree TEC is presented in Appendix H. The 'likely' and 'may' occur categories of the TEC's distribution is based upon Queensland Government regional ecosystem mapping. No other TECs were revealed by the PMST.

The likely to occur category has been assigned to the following regional ecosystems of the Wet Tropics and Central Mackay Coast Bioregions, including:

- → 7.3.8a Floodplain (other than floodplain wetlands). *Melaleuca viridiflora* open forest to open woodland. Includes areas of natural invasion onto former grasslands. Alluvial plains.
- → 7.3.8b Floodplain (other than floodplain wetlands). *Melaleuca viridiflora* open forest to open woodland with eucalypt emergents (or sparse eucalypt overstorey) of species such as *Corymbia clarksoniana*, *Eucalyptus platyphylla*, *Lophostemon suaveolens* and *E. drepanophylla*. Poorly drained alluvium, mostly on the coastal plains.
- → 7.3.8c Floodplain (other than floodplain wetlands). Melaleuca viridiflora and Lophostemon suaveolens open forest to woodland. Poorly drained soils of coastal lowlands.
- → 7.3.8d Floodplain (other than floodplain wetlands). Melaleuca viridiflora, Lophostemon suaveolens and Allocasuarina littoralis open shrubland. Poorly drained soils of coastal lowlands.
- → 7.5.4g Floodplain (other than floodplain wetlands). *Melaleuca viridiflora* woodland. Laterite.
- > 8.3.2 Melaleuca viridiflora woodland on seasonally inundated alluvial plains with impeded drainage
- → 8.5.2a Melaleuca viridiflora var. viridiflora open woodland to open forest (5–20 m tall). Poorly drained, duplex soils with a sandy surface.
- → 8.5.2c Melaleuca viridiflora var. viridiflora and M. nervosa open woodland to open forest. Occurs on Tertiary sand plains on gently undulating plains of lowlands, dissected by many incised streams.
- → 8.5.6 Melaleuca viridiflora +/- Allocasuarina littoralis woodland on Tertiary sand plains.

None of the above listed regional ecosystems occur in the three corridor investigation areas.

The Broad Leaf Tea-tree TEC may also occur category within regional ecosystems on Land Zones 3 and 5 of the Wet Tropics and Central Mackay Coast Bioregions. However, based upon the regional ecosystem mapping, it has been determined there is a low possibility of the Broad Leaf Tea-tree TEC occurring in the corridor investigation areas.

# 3.3 Wet Tropics of Queensland – World, National and Indigenous Heritage properties

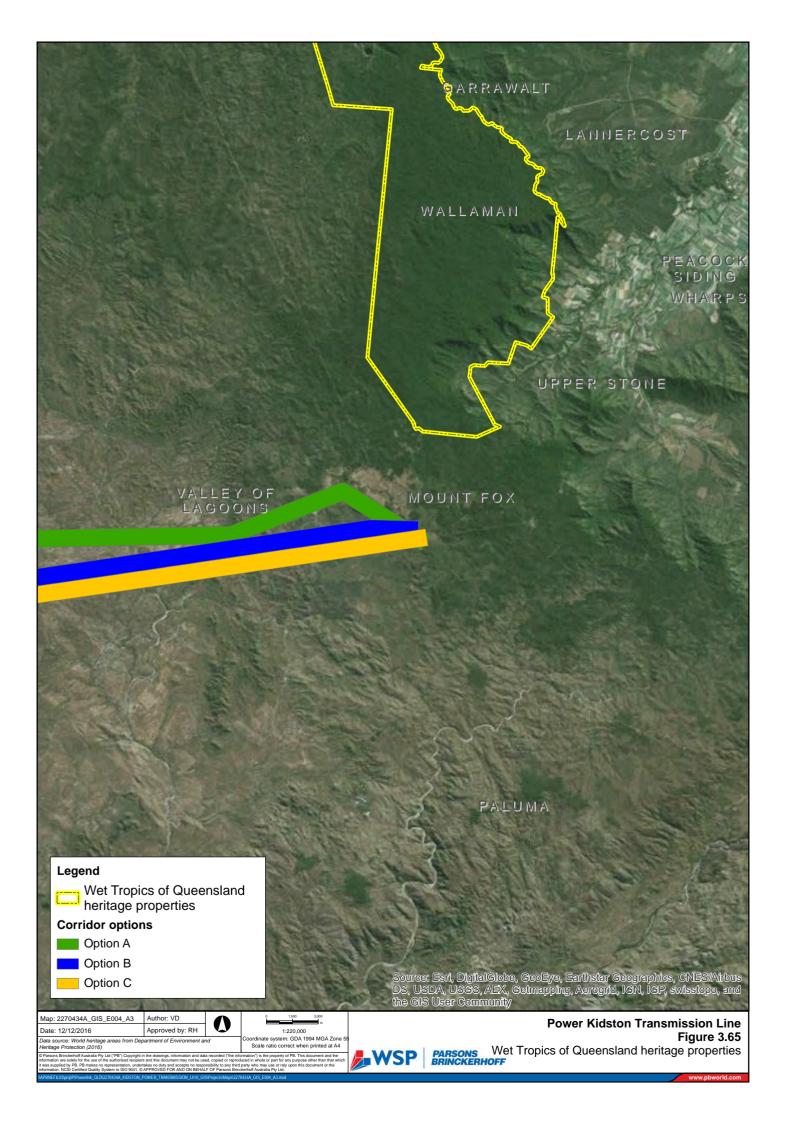
The Wet Tropics of Queensland World Heritage properties were inscribed on the World Heritage List in 1988. It was then included in the National Heritage List on 21 May 2007, as one of the one of 15 World Heritage places of Australia. On 9 November 2012 the Australian Government announced the inclusion of the national Indigenous heritage values, as part of the existing National Heritage Listing for the Wet Tropics of Queensland. A distribution map for the Wet Tropics of Queensland heritage properties, is presented in Appendix I.

None of corridor investigation areas intersect the Wet Tropics of Queensland heritage properties, as shown on Figure 3.65.

The distance of the Wet Tropics of Queensland heritage properties from each corridor investigation area is presented in Table 3.14. Based upon the distances from each corridor investigation area, the Project would be unlikely to indirectly impact the Wet Tropics of Queensland heritage properties. However this would need to be assessed in greater detail during the next phase of the Project.

OPTION	DISTANCE FROM WET TROPICS OF QUEENSLAND HERITAGE PROPERTIES (KM)
Option A	4.8
Option B	5.2
Option C	5.6

Table 3.14 Distance of each corridor option from the Wet Tropics of Queensland heritage properties



# 4 PRELIMINARY ADVICE

#### 4.1 Ecological constraints

The least amount of ecological constraints (MNES and MSES) are primarily associated with Option C, with Option B being the next preferred option with a moderate amount of ecological constraints, and Option A with the greatest amount of ecological constraints, as listed in Table 4.1.

 Table 4.1
 Summary of ecological constraint rankings for each corridor option

ECOLOGICAL CONSTRAINT	CON	CONSTRAINT RANKINGS		
	Option A	Option B	Option C	
Matters of State Environmental Significance				
NC Act threatened species records (numbers)	2	4	5	
Regulated of concern dominant vegetation (ha)	2,059	1,819	1,519	
Regulated of concern sub-dominant vegetation (ha)	1,008	442	334	
Regulated of high value regrowth vegetation (ha)	3.6	2.4	2.4	
Regulated Category R regrowth vegetation (ha)	422	296	257	
Mapped of concern dominant regional ecosystems (ha)	1,008	442	334	
Mapped of concern dominant regional ecosystems (ha)	2,059	1,891	1,519	
Predicted of concern dominant woody vegetation (ha)	1,535	1,460	1,223	
Predicted of concern sub-dominant woody vegetation (ha)	549	295	618	
Potential of concern dominant woody vegetation (ha)	240	344	431	
Potential of concern sub-dominant woody vegetation (ha)	364	87	37	
Regulated remnant watercourse Wet Tropics bioregion (ha)	118	50	45	
Regulated remnant watercourse vegetation Einasleigh Uplands bioregion (ha)	5,585	5,547	5,493	
Regulated remnant wetland vegetation (ha)	4.7	42.3	27.7	
Protected areas (ha)	658	214	145	
Wildlife habitats (ha)	522	411	385	
High ecological significance (HES) wetlands (ha)	0	1.1	0	
Matters of National Environmental Significance				
Threatened and migratory species records (numbers)	3	6	5	
Proximity to Wet Tropics of Queensland heritage properties (km)	4.8	5.2	5.6	

Table 4.1 clearly indicates that Option C would result in the least amount of impact upon MNES and MSES and would be the corridor of least ecological constraint for the Project. It also clearly indicates that Option A has the greatest amount of ecological constraint, while Option B has a moderate amount of ecological constraint, while Option C.

#### 4.2 Flora and fauna surveys

As part of the next phase of the Project, it would be necessary to undertake field surveys to verify the presence and extent of the ecological constraints outlined in Table 4.1.

Targeted flora and fauna surveys may identify ecological constraints that have not been identified by this desktop assessment. For example, the threatened species records are representative of the limited amount of ecological investigations that have been conducted across the relevant bioregions, in particular the Einasleigh Uplands bioregion. This reflects the remoteness of the corridor investigation areas where there is a low amount of large developments that require ecological investigations and environmental approvals. It is possible that targeted surveys for the Project may identify threatened species and supporting habitats that have not yet been recorded and formerly placed on the DSITI and ALA records databases.

#### 4.3 Regional ecosystem verification

Vegetation surveys would be required for the Project to verify DNRM mapped regional ecosystems. It is often the case that DNRM regional ecosystem mapping can be inaccurate at a project scale. This is a function of both scale and availability of field verified vegetation survey data.

The remote sensing analysis of woody vegetation, completed by Virtual GIS, has used a similar methodology as that used by the Queensland Herbarium. It is useful in terms of identifying areas potentially containing of concern dominant and sub-dominant regional ecosystems. However, it too can be expected to have a certain amount of inaccuracy compared to what may be revealed by field verification surveys.

#### 4.4 Significant impact assessments for MNES

Of the MNES assessed, threatened and/or migratory species are the only MNES of relevance to the three corridor options. On the other hand, MNES such as threatened ecological communities, world and national heritage properties, nationally important wetlands and the Great Barrier Reef, have been assessed as not being relevant to the three corridor options.

Under the requirements of the EPBC Act, significant impact assessments in accordance with the *Matters of National Environmental Significance: Significant impact guidelines 1.1* (2013), would need to be applied to MNES that are recorded or considered likely occurrences in areas that may be subject to project related impacts. These significant impact assessments may be required for the threatened species listed under the EPBC Act, for which records have been revealed by this assessment, including:

- → Curlew Sandpiper (*Calidris ferruginea*) Critically endangered
- → White-throated Needletail (*Hirundapus caudacutus*) Migratory (terrestrial)
- → Greater Glider (*Petauroides volans*) Vulnerable
- → Sharman's Rock-wallaby (Petrogale sharmani) Vulnerable
- → Koala (*Phascolarctos cinereus*) Vulnerable
- → Southern Black-throated Finch (Poephila cincta cincta) Endangered
- → Bluegrass (*Dichanthium setosum*) Vulnerable.

It is possible that field surveys may identify other MNES that may also require significant impacts assessments. It is recommended that significant impacts assessments be prepared in respect to predicted project impacts for the yet to be selected corridor option. The design process for the preferred option transmission line will consider opportunities to avoid or minimise impacts upon MNES. This would also be coupled with mitigation strategies to mitigate any potential impacts upon MNES, thus reducing the overall risk and extent (ha) of residual impacts upon the relevant MNES.

The outcomes of the significant impact assessments should identify whether the Project should referral is required for or not. The significant impact assessments would also identify any potential environmental offsets for the Project under the Department of the Environment and Energy's *EPBC Act Environmental Offsets Policy* (2012).

#### 4.5 Queensland – Significant residual impact assessments for MSES

In accordance with Queensland Government's *Environmental Offsets Policy version 1.1* (2014), potential impacts upon MSES need to be applied in accordance with either the Department of Environment and Heritage Protection's *Significant Residual Impact Guideline* (2014), or the Department of Development, Infrastructure and Planning's *Significant Residual Impact Guideline* (2014), which would be dependent on the approval pathway for the Project. It is important to note that significant impact assessments for an MNES that is also an MSES are not required under the Queensland Government framework.

In reference to this assessment the MSES that are likely to require significant residual impact assessments, include:

- → threatened species, including:
  - Acacia tingoorensis Vulnerable
- $\rightarrow$  regulated vegetation, including:
  - of concern regional ecosystems
  - of concern high value regrowth (to be confirmed as to why it is in the regulated vegetation layer)
  - category R regrowth vegetation (approval pathway dependant)
  - remnant watercourse vegetation
- protected areas
- → wildlife habitats
- → high ecological significance
- → connectivity.

Depending on the outcomes of the significant residual impact assessments, some if not all of the above MSES, may require environmental offsets to compensate for project related residual impacts.

Once a preferred corridor option is selected and project impact area(s) are defined, the significant residual impact upon connectivity may also need to be assessed using the Landscape Fragmentation and Connectivity tool.

#### 4.6 Summary

This initial desktop assessment and preliminary ecological constraints advice has identified Corridor Option C as containing the least amount of ecological constraints for the proposed power transmission line project.

This initial desktop assessment has identified the MNES and MSES that are of potential relevance to the Project and has provided an indication of the next steps that may be required for the Project if it is to proceed to the EIS or other approval pathway stage. Potential environmental offset commitments under Commonwealth and State environmental offset policies have also been identified.

Once a preferred corridor option is selected by the Corridor Options Report and predicted project related impacts are understood and tangible, it is recommended that Powerlink and/or Genex consider preparing significant impact assessments for MNES to determine whether an EPBC Act referral is required.

In addition, significant impact assessments (MNES) and significant residual impact assessments (MSES) should be instrumental in identifying the potential Commonwealth and State environmental offset requirements, and an approach to Commonwealth and State environmental offsets delivery.

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# Appendix A

**PROTECTED MATTERS SEARCH TOOL** 

Australian Government



Department of the Environment and Energy

# **EPBC Act Protected Matters Report**

This report provides general guidance on matters of national environmental significance and other matters protected by the EPBC Act in the area you have selected.

Information on the coverage of this report and qualifications on data supporting this report are contained in the caveat at the end of the report.

Information is available about <u>Environment Assessments</u> and the EPBC Act including significance guidelines, forms and application process details.

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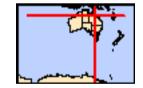
Summary Details Matters of NES Other Matters Protected by the EPBC Act Extra Information Caveat

<u>Acknowledgements</u>



This map may contain data which are ©Commonwealth of Australia (Geoscience Australia), ©PSMA 2010

Coordinates Buffer: 5.0Km



# Summary

## Matters of National Environmental Significance

This part of the report summarises the matters of national environmental significance that may occur in, or may relate to, the area you nominated. Further information is available in the detail part of the report, which can be accessed by scrolling or following the links below. If you are proposing to undertake an activity that may have a significant impact on one or more matters of national environmental significance then you should consider the <u>Administrative Guidelines on Significance</u>.

World Heritage Properties:	1
National Heritage Places:	2
Wetlands of International Importance:	None
Great Barrier Reef Marine Park:	None
Commonwealth Marine Area:	None
Listed Threatened Ecological Communities:	1
Listed Threatened Species:	33
Listed Migratory Species:	15

## Other Matters Protected by the EPBC Act

This part of the report summarises other matters protected under the Act that may relate to the area you nominated. Approval may be required for a proposed activity that significantly affects the environment on Commonwealth land, when the action is outside the Commonwealth land, or the environment anywhere when the action is taken on Commonwealth land. Approval may also be required for the Commonwealth or Commonwealth agencies proposing to take an action that is likely to have a significant impact on the environment anywhere.

The EPBC Act protects the environment on Commonwealth land, the environment from the actions taken on Commonwealth land, and the environment from actions taken by Commonwealth agencies. As heritage values of a place are part of the 'environment', these aspects of the EPBC Act protect the Commonwealth Heritage values of a Commonwealth Heritage place. Information on the new heritage laws can be found at http://www.environment.gov.au/heritage

A <u>permit</u> may be required for activities in or on a Commonwealth area that may affect a member of a listed threatened species or ecological community, a member of a listed migratory species, whales and other cetaceans, or a member of a listed marine species.

Commonwealth Land:	None
Commonwealth Heritage Places:	None
Listed Marine Species:	22
Whales and Other Cetaceans:	None
Critical Habitats:	None
Commonwealth Reserves Terrestrial:	None
Commonwealth Reserves Marine:	None

## **Extra Information**

This part of the report provides information that may also be relevant to the area you have nominated.

State and Territory Reserves:	4
Regional Forest Agreements:	None
Invasive Species:	22
Nationally Important Wetlands:	1
Key Ecological Features (Marine)	None

# Details

## Matters of National Environmental Significance

World Heritage Properties		[Resource Information]
Name	State	Status
Wet Tropics of Queensland	QLD	Declared property
National Heritage Properties		[Resource Information]
Name	State	Status
Natural		
Wet Tropics of Queensland	QLD	Listed place
Indigenous		
Wet Tropics World Heritage Area (Indigenous Values)	QLD	Within listed place

### Listed Threatened Ecological Communities

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

[Resource Information]

Name	Status	Type of Presence
Broad leaf tea-tree (Melaleuca viridiflora) woodlands in	Endangered	Community may occur
high rainfall coastal north Queensland		within area

Listed Threatened Species		[Resource Information]
Name	Status	Type of Presence
Birds		
Calidris ferruginea		
Curlew Sandpiper [856]	Critically Endangered	Species or species habitat may occur within area
Casuarius casuarius johnsonii Southern Cassowary, Australian Cassowary, Double- wattled Cassowary [25986]	Endangered	Species or species habitat known to occur within area
Erythrotriorchis radiatus Red Goshawk [942]	Vulnerable	Species or species habitat likely to occur within area



#### Frogs

Name	Status	Type of Presence
Litoria dayi Australian Lace-lid, Lace-eyed Tree Frog [86707]	Endangered	Species or species habitat likely to occur within area
<u>Litoria nannotis</u> Waterfall Frog, Torrent Tree Frog [1817]	Endangered	Species or species habitat may occur within area
<u>Litoria rheocola</u> Common Mistfrog [1802]	Endangered	Species or species habitat likely to occur within area
Mammals		
Dasyurus hallucatus Northern Quoll, Digul [331]	Endangered	Species or species habitat likely to occur within area
Dasyurus maculatus gracilis Spotted-tailed Quoll (North Queensland), Yarri [64475]	Endangered	Species or species habitat likely to occur within area
<u>Hipposideros semoni</u> Semon's Leaf-nosed Bat, Greater Wart-nosed Horseshoe-bat [180]	Endangered	Species or species habitat may occur within area
Macroderma gigas Ghost Bat [174]	Vulnerable	Species or species habitat likely to occur within area
Mesembriomys gouldii rattoides Black-footed Tree-rat (north Queensland), Shaggy Rabbit-rat [87620]	Vulnerable	Species or species habitat likely to occur within area
Petauroides volans Greater Glider [254]	Vulnerable	Species or species habitat known to occur within area
Petrogale sharmani Mount Claro Rock Wallaby, Sharman's Rock Wallaby [59281]	Vulnerable	Species or species habitat known to occur within area
Phascolarctos cinereus (combined populations of Qld, Koala (combined populations of Queensland, New South Wales and the Australian Capital Territory) [85104] Pteropus conspicillatus	NSW and the ACT) Vulnerable	Species or species habitat known to occur within area

Spectacled Flying-fox [185]	Vulnerable	Species or species habitat known to occur within area
Pteropus poliocephalus Grey-headed Flying-fox [186]	Vulnerable	Foraging, feeding or related behaviour likely to occur
Rhinolophus robertsi Large-eared Horseshoe Bat, Greater Large-eared	Endangered	within area Species or species habitat
Horseshoe Bat [87639] Saccolaimus saccolaimus nudicluniatus		likely to occur within area
Bare-rumped Sheath-tailed Bat, Bare-rumped Sheathtail Bat [66889]	Critically Endangered	Species or species habitat likely to occur within area
Plants		
Acacia crombiei Pink Gidgee [10927]	Vulnerable	Species or species habitat may occur within area
Bulbophyllum globuliforme Miniature Moss-orchid, Hoop Pine Orchid [6649]	Vulnerable	Species or species habitat likely to occur within area
<u>Cajanus mareebensis</u> [8635]	Endangered	Species or species habitat likely to occur

Name	Status	Type of Presence
		within area
Dichanthium setosum		
bluegrass [14159]	Vulnerable	Species or species habitat likely to occur within area
Lindsaea pulchella var. blanda		
[20842]	Vulnerable	Extinct within area
Marsdenia brevifolia		
[64585]	Vulnerable	Species or species habitat likely to occur within area
Phaius australis		
Lesser Swamp-orchid [5872]	Endangered	Species or species habitat likely to occur within area
<u>Tephrosia leveillei</u>		
[16946]	Vulnerable	Species or species habitat likely to occur within area
Reptiles		
Egernia rugosa		
Yakka Skink [1420]	Vulnerable	Species or species habitat may occur within area
Lerista vittata		
Mount Cooper Striped Lerista [1308]	Vulnerable	Species or species habitat likely to occur within area
Listed Migratory Species		[Resource Information]
* Species is listed under a different scientific name on	the EPBC Act - Threatenec	I Species list.
Name	Threatened	Type of Presence
Migratory Marine Birds		
Apus pacificus		
Fork-tailed Swift [678]		Species or species habitat likely to occur within area
Migratory Marine Species		
Crocodylus porosus		
Salt-water Crocodile, Estuarine Crocodile [1774]		Species or species habitat likely to occur within area
Migratory Terrestrial Species		
Cuculus optatus		
Oriental Cuckoo, Horsfield's Cuckoo [86651]		Species or species habitat

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Hirundapus caudacutus White-throated Needletail [682]

Hirundo rustica Barn Swallow [662]

Monarcha melanopsis Black-faced Monarch [609]

Monarcha trivirgatus Spectacled Monarch [610]

Motacilla cinerea Grey Wagtail [642]

Motacilla flava Yellow Wagtail [644] known to occur within area

Species or species habitat may occur within area

Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Name	Threatened	Type of Presence
Myiagra cyanoleuca		
Satin Flycatcher [612]		Species or species habitat likely to occur within area
Rhipidura rufifrons		
Rufous Fantail [592]		Species or species habitat known to occur within area
Migratory Wetlands Species		
Calidris ferruginea		
Curlew Sandpiper [856]	Critically Endangered	Species or species habitat may occur within area
<u>Gallinago hardwickii</u>		
Latham's Snipe, Japanese Snipe [863]		Species or species habitat may occur within area
Numenius madagascariensis		
Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Pandion haliaetus		
Osprey [952]		Species or species habitat likely to occur within area
Other Matters Protected by the EPBC Act		
Listed Marine Species		[Resource Information]
* Species is listed under a different scientific name on	the EPBC Act - Threatened	d Species list.
Name	Threatened	Type of Presence
Birds		
Anseranas semipalmata		
Magpie Goose [978]		Species or species habitat may occur within area
Apus pacificus		
Fork-tailed Swift [678]		Species or species habitat likely to occur within area
<u>Ardea alba</u>		
Great Egret, White Egret [59541]		Species or species habitat likely to occur within area

#### Ardea ibis



Calidris ferruginea Curlew Sandpiper [856]

<u>Cuculus saturatus</u> Oriental Cuckoo, Himalayan Cuckoo [710]

Gallinago hardwickii Latham's Snipe, Japanese Snipe [863]

Haliaeetus leucogaster White-bellied Sea-Eagle [943]

Hirundapus caudacutus White-throated Needletail [682] Species or species habitat may occur within area

Critically Endangered

Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Species or species habitat may occur within area

Name	Threatened	Type of Presence
<u>Hirundo rustica</u> Barn Swallow [662]		Species or species habitat may occur within area
<u>Merops ornatus</u> Rainbow Bee-eater [670]		Species or species habitat may occur within area
Monarcha melanopsis Black-faced Monarch [609]		Species or species habitat known to occur within area
Monarcha trivirgatus Spectacled Monarch [610]		Species or species habitat known to occur within area
Motacilla cinerea Grey Wagtail [642]		Species or species habitat may occur within area
<u>Motacilla flava</u> Yellow Wagtail [644]		Species or species habitat likely to occur within area
Myiagra cyanoleuca Satin Flycatcher [612]		Species or species habitat likely to occur within area
Numenius madagascariensis Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Pandion haliaetus Osprey [952]		Species or species habitat likely to occur within area
<u>Rhipidura rufifrons</u> Rufous Fantail [592]		Species or species habitat known to occur within area
Rostratula benghalensis (sensu lato) Painted Snipe [889]	Endangered*	Species or species habitat likely to occur within area
Reptiles		

Reptiles Crocodylus johnstoni

Freshwater Crocodile, Johnston's Crocodile, Johnston's River Crocodile [1773]

Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Species or species habitat may occur within area

Species or species habitat likely to occur within area

#### **Extra Information**

State and Territory Reserves	[Resource Information]
Name	State
Girringun	QLD
Girringun	QLD
Liefway	QLD
Newcastle Range-The Oaks	QLD

Inva	asive	Species						1	Re	sour	<u>ce Ir</u>	nforma	ation ]	
			 ~ ~		 			1.1						

Weeds reported here are the 20 species of national significance (WoNS), along with other introduced plants that are considered by the States and Territories to pose a particularly significant threat to biodiversity. The following feral animals are reported: Goat, Red Fox, Cat, Rabbit, Pig, Water Buffalo and Cane Toad. Maps from Landscape Health Project, National Land and Water Resouces Audit, 2001.

Name	Status	Type of Presence
Birds		
Columba livia		
Rock Pigeon, Rock Dove, Domestic Pigeon [803]		Species or species habitat likely to occur within area
Lonchura punctulata		
Nutmeg Mannikin [399]		Species or species habitat likely to occur within area
Passer domesticus		
House Sparrow [405]		Species or species habitat likely to occur within area
Streptopelia chinensis		
Spotted Turtle-Dove [780]		Species or species habitat likely to occur within area
Sturnus vulgaris		
Common Starling [389]		Species or species habitat likely to occur within area
Frogs		
Rhinella marina		
Cane Toad [83218]		Species or species habitat likely to occur within area
Mammals		

Bos taurus Domestic Cattle [16]

Species or species habitat likely to occur within area

Canis lupus familiaris Domestic Dog [82654]

Equus caballus Horse [5]

Felis catus Cat, House Cat, Domestic Cat [19]

Oryctolagus cuniculus Rabbit, European Rabbit [128]

Sus scrofa Pig [6]

Vulpes vulpes Red Fox, Fox [18] Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur

Name	Status	Type of Presence
Dianta		within area
Plants Acacia nilotica subsp. indica		
Prickly Acacia [6196]		Species or species habitat may occur within area
Cenchrus ciliaris		
Buffel-grass, Black Buffel-grass [20213]		Species or species habitat may occur within area
Cryptostegia grandiflora		
Rubber Vine, Rubbervine, India Rubber Vine, India Rubbervine, Palay Rubbervine, Purple Allamanda [18913]		Species or species habitat likely to occur within area
Hymenachne amplexicaulis		
Hymenachne, Olive Hymenachne, Water Stargrass West Indian Grass, West Indian Marsh Grass [3175		Species or species habitat likely to occur within area
Jatropha gossypifolia		
Cotton-leaved Physic-Nut, Bellyache Bush, Cotton-l Physic Nut, Cotton-leaf Jatropha, Black Physic Nut [7507]	leaf	Species or species habitat likely to occur within area
Lantana camara Lantana, Common Lantana, Kamara Lantana, Large leaf Lantana, Pink Flowered Lantana, Red Flowered Lantana, Red-Flowered Sage, White Sage, Wild Sa [10892] Parkinsonia aculeata	b	Species or species habitat likely to occur within area
Parkinsonia, Jerusalem Thorn, Jelly Bean Tree, Ho	rse	Species or species habitat
Bean [12301]		likely to occur within area
Parthenium hysterophorus		
Parthenium Weed, Bitter Weed, Carrot Grass, False Ragweed [19566]	9	Species or species habitat likely to occur within area
Salvinia molesta		
Salvinia, Giant Salvinia, Aquarium Watermoss, Kari Weed [13665]	ba	Species or species habitat likely to occur within area
Nationally Important Wetlands		[Resource Information]
Name		State
Poison Lake		QLD

# Caveat

The information presented in this report has been provided by a range of data sources as acknowledged at the end of the report.

This report is designed to assist in identifying the locations of places which may be relevant in determining obligations under the Environment Protection and Biodiversity Conservation Act 1999. It holds mapped locations of World and National Heritage properties, Wetlands of International and National Importance, Commonwealth and State/Territory reserves, listed threatened, migratory and marine species and listed threatened ecological communities. Mapping of Commonwealth land is not complete at this stage. Maps have been collated from a range of sources at various resolutions.

Not all species listed under the EPBC Act have been mapped (see below) and therefore a report is a general guide only. Where available data supports mapping, the type of presence that can be determined from the data is indicated in general terms. People using this information in making a referral may need to consider the qualifications below and may need to seek and consider other information sources.

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

For species where the distributions are well known, maps are digitised from sources such as recovery plans and detailed habitat studies. Where appropriate, core breeding, foraging and roosting areas are indicated under 'type of presence'. For species whose distributions are less well known, point locations are collated from government wildlife authorities, museums, and non-government organisations; bioclimatic distribution models are generated and these validated by experts. In some cases, the distribution maps are based solely on expert knowledge.

Only selected species covered by the following provisions of the EPBC Act have been mapped:

- migratory and
- marine

The following species and ecological communities have not been mapped and do not appear in reports produced from this database:

- threatened species listed as extinct or considered as vagrants
- some species and ecological communities that have only recently been listed
- some terrestrial species that overfly the Commonwealth marine area
- migratory species that are very widespread, vagrant, or only occur in small numbers

The following groups have been mapped, but may not cover the complete distribution of the species:

- non-threatened seabirds which have only been mapped for recorded breeding sites
- seals which have only been mapped for breeding sites near the Australian continent

Such breeding sites may be important for the protection of the Commonwealth Marine environment.

# Coordinates

-18.87024 144.1472,-18.87927 144.1471,-18.87901 144.1329,-18.87735 144.13,-18.87434 144.129,-18.8443 144.1393,-18.84225 144.1411,-18.84156 144.1438,-18.84368 144.1898,-18.8636 144.2184,-18.88303 144.2839,-18.87004 144.3748,-18.8979 144.5356,-18.90103 145.1652,-18.84323 145.2928,-18.83884 145.7251,-18.81633 145.7881,-18.84454 145.8358,-18.85216 145.8307,-18.82587 145.7873,-18.84778 145.7269,-18.85223 145.2943,-18.90995 145.1673,-18.90686 144.5343,-18.87912 144.375,-18.89211 144.2828,-18.87187 144.2145,-18.85232 144.1863,-18.85101 144.1469,-18.87014 144.1403,-18.87024 144.1472

# Acknowledgements

This database has been compiled from a range of data sources. The department acknowledges the following custodians who have contributed valuable data and advice:

-Office of Environment and Heritage, New South Wales

-Department of Environment and Primary Industries, Victoria

-Department of Primary Industries, Parks, Water and Environment, Tasmania

-Department of Environment, Water and Natural Resources, South Australia

-Parks and Wildlife Commission NT, Northern Territory Government

-Department of Environmental and Heritage Protection, Queensland

-Department of Parks and Wildlife, Western Australia

-Environment and Planning Directorate, ACT

-Birdlife Australia

-Australian Bird and Bat Banding Scheme

-Australian National Wildlife Collection

-Natural history museums of Australia

-Museum Victoria

-Australian Museum

-South Australian Museum

-Queensland Museum

-Online Zoological Collections of Australian Museums

-Queensland Herbarium

-National Herbarium of NSW

-Royal Botanic Gardens and National Herbarium of Victoria

-Tasmanian Herbarium

-State Herbarium of South Australia

-Northern Territory Herbarium

-Western Australian Herbarium

-Australian National Herbarium, Atherton and Canberra

-University of New England

-Ocean Biogeographic Information System

-Australian Government, Department of Defence

Forestry Corporation, NSW

-Geoscience Australia

-CSIRO

-Other groups and individuals

The Department is extremely grateful to the many organisations and individuals who provided expert advice and information on numerous draft distributions.

Please feel free to provide feedback via the <u>Contact Us</u> page.

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Department of the Environment and Energy

# **EPBC Act Protected Matters Report**

This report provides general guidance on matters of national environmental significance and other matters protected by the EPBC Act in the area you have selected.

Information on the coverage of this report and qualifications on data supporting this report are contained in the caveat at the end of the report.

Information is available about <u>Environment Assessments</u> and the EPBC Act including significance guidelines, forms and application process details.

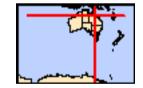
Report created: 03/11/16 14:49:40

Summary Details Matters of NES Other Matters Protected by the EPBC Act Extra Information Caveat Acknowledgements



This map may contain data which are ©Commonwealth of Australia (Geoscience Australia), ©PSMA 2010

Coordinates Buffer: 5.0Km



# Summary

#### Matters of National Environmental Significance

This part of the report summarises the matters of national environmental significance that may occur in, or may relate to, the area you nominated. Further information is available in the detail part of the report, which can be accessed by scrolling or following the links below. If you are proposing to undertake an activity that may have a significant impact on one or more matters of national environmental significance then you should consider the <u>Administrative Guidelines on Significance</u>.

World Heritage Properties:	None
National Heritage Places:	None
Wetlands of International Importance:	None
Great Barrier Reef Marine Park:	None
Commonwealth Marine Area:	None
Listed Threatened Ecological Communities:	1
Listed Threatened Species:	33
Listed Migratory Species:	16

#### Other Matters Protected by the EPBC Act

This part of the report summarises other matters protected under the Act that may relate to the area you nominated. Approval may be required for a proposed activity that significantly affects the environment on Commonwealth land, when the action is outside the Commonwealth land, or the environment anywhere when the action is taken on Commonwealth land. Approval may also be required for the Commonwealth or Commonwealth agencies proposing to take an action that is likely to have a significant impact on the environment anywhere.

The EPBC Act protects the environment on Commonwealth land, the environment from the actions taken on Commonwealth land, and the environment from actions taken by Commonwealth agencies. As heritage values of a place are part of the 'environment', these aspects of the EPBC Act protect the Commonwealth Heritage values of a Commonwealth Heritage place. Information on the new heritage laws can be found at http://www.environment.gov.au/heritage

A <u>permit</u> may be required for activities in or on a Commonwealth area that may affect a member of a listed threatened species or ecological community, a member of a listed migratory species, whales and other cetaceans, or a member of a listed marine species.

Commonwealth Land:	None
Commonwealth Heritage Places:	None
Listed Marine Species:	23
Whales and Other Cetaceans:	None
Critical Habitats:	None
Commonwealth Reserves Terrestrial:	None
Commonwealth Reserves Marine:	None

#### **Extra Information**

This part of the report provides information that may also be relevant to the area you have nominated.

State and Territory Reserves:	4
Regional Forest Agreements:	None
Invasive Species:	22
Nationally Important Wetlands:	None
Key Ecological Features (Marine)	None

# Details

## Matters of National Environmental Significance

#### Listed Threatened Ecological Communities

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

[Resource Information]

known to occur within area

Name	Status	Type of Presence
Broad leaf tea-tree (Melaleuca viridiflora) woodlands in high rainfall coastal north Queensland	Endangered	Community may occur within area
Listed Threatened Species		[Resource Information]
Name	Status	Type of Presence
Birds		
<u>Calidris ferruginea</u> Curlew Sandpiper [856]	Critically Endangered	Species or species habitat may occur within area
Casuarius casuarius johnsonii Southern Cassowary, Australian Cassowary, Double- wattled Cassowary [25986]	Endangered	Species or species habitat known to occur within area
<u>Erythrotriorchis radiatus</u> Red Goshawk [942]	Vulnerable	Species or species habitat likely to occur within area
<u>Erythrura gouldiae</u> Gouldian Finch [413]	Endangered	Species or species habitat may occur within area
Numenius madagascariensis Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Poephila cincta cincta Southern Black-throated Finch [64447]	Endangered	Species or species habitat

Rostratula australis

Australian Painted Snipe [77037]	Endangered	Species or species habitat likely to occur within area
<u>Tyto novaehollandiae kimberli</u> Masked Owl (northern) [26048]	Vulnerable	Species or species habitat likely to occur within area
Frogs		
Litoria dayi Australian Lace-lid, Lace-eyed Tree Frog [86707]	Endangered	Species or species habitat likely to occur within area
<u>Litoria nannotis</u> Waterfall Frog, Torrent Tree Frog [1817]	Endangered	Species or species habitat may occur within area

Name	Status	Type of Presence
<u>Litoria rheocola</u> Common Mistfrog [1802]	Endangered	Species or species habitat likely to occur within area
Mammals		
<u>Dasyurus hallucatus</u> Northern Quoll, Digul [331]	Endangered	Species or species habitat likely to occur within area
Dasyurus maculatus gracilis Spotted-tailed Quoll (North Queensland), Yarri [64475]	Endangered	Species or species habitat likely to occur within area
Hipposideros semoni Semon's Leaf-nosed Bat, Greater Wart-nosed Horseshoe-bat [180]	Endangered	Species or species habitat may occur within area
<u>Macroderma gigas</u> Ghost Bat [174]	Vulnerable	Species or species habitat likely to occur within area
Mesembriomys gouldii rattoides Black-footed Tree-rat (north Queensland), Shaggy Rabbit-rat [87620]	Vulnerable	Species or species habitat likely to occur within area
Petauroides volans Greater Glider [254]	Vulnerable	Species or species habitat known to occur within area
<u>Petrogale sharmani</u> Mount Claro Rock Wallaby, Sharman's Rock Wallaby [59281]	Vulnerable	Species or species habitat known to occur within area
Phascolarctos cinereus (combined populations of Qld, Koala (combined populations of Queensland, New South Wales and the Australian Capital Territory) [85104]	NSW and the ACT) Vulnerable	Species or species habitat known to occur within area
Pteropus conspicillatus Spectacled Flying-fox [185]	Vulnerable	Species or species habitat known to occur within area
Pteropus poliocephalus Grey-headed Flying-fox [186]	Vulnerable	Foraging, feeding or related behaviour likely to occur within area
<u>Rhinolophus robertsi</u> Large-eared Horseshoe Bat, Greater Large-eared Horseshoe Bat [87639]	Endangered	Species or species habitat known to occur within area
Saccolaimus saccolaimus nudicluniatus Bare-rumped Sheath-tailed Bat, Bare-rumped Sheathtail Bat [66889]	Critically Endangered	Species or species habitat likely to occur within area
Plants		
<u>Acacia crombiei</u> Pink Gidgee [10927]	Vulnerable	Species or species habitat may occur within area
Bulbophyllum globuliforme Miniature Moss-orchid, Hoop Pine Orchid [6649]	Vulnerable	Species or species habitat may occur within area
<u>Cajanus mareebensis</u> [8635]	Endangered	Species or species habitat likely to occur within area
Dichanthium setosum bluegrass [14159]	Vulnerable	Species or species habitat likely to occur within area
<u>Lindsaea pulchella var. blanda</u> [20842]	Vulnerable	Extinct within area

Name	Status	Type of Presence
<u>Marsdenia brevifolia</u> [64585]	Vulnerable	Species or species habitat likely to occur within area
Phaius australis Lesser Swamp-orchid [5872]	Endangered	Species or species habitat likely to occur within area
Tephrosia leveillei [16946]	Vulnerable	Species or species habitat likely to occur within area
Reptiles		
Egernia rugosa		
Yakka Skink [1420]	Vulnerable	Species or species habitat may occur within area
<u>Lerista vittata</u> Mount Cooper Striped Lerista [1308]	Vulnerable	Species or species habitat
		likely to occur within area
Listed Migratory Species		[Resource Information]
* Species is listed under a different scientific name or	the EPBC Act - Threatened	
Name	the EPBC Act - Threatened Threatened	d Species list. Type of Presence
Name Migratory Marine Birds		
Name		
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678]		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678] Migratory Marine Species		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678]		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678] Migratory Marine Species <u>Crocodylus porosus</u>		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774]		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within area
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus Oriental Cuckoo, Horsfield's Cuckoo [86651]		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within area
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus Oriental Cuckoo, Horsfield's Cuckoo [86651]		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat known to occur within areaSpecies or species habitat known to occur within areaSpecies or species habitat

Barn Swallow [662]

Monarcha melanopsis Black-faced Monarch [609]

Monarcha trivirgatus Spectacled Monarch [610]

Motacilla cinerea Grey Wagtail [642]

Motacilla flava Yellow Wagtail [644]

Myiagra cyanoleuca Satin Flycatcher [612]

<u>Rhipidura rufifrons</u> Rufous Fantail [592] Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat known to occur within area

Threatened	Type of Presence
Critically Endangered	Species or species habitat may occur within area
	Species or species habitat may occur within area
Critically Endangered	Species or species habitat may occur within area
	Species or species habitat likely to occur within area
	Species or species habitat may occur within area

### Other Matters Protected by the EPBC Act

Listed Marine Species		[Resource Information]
* Species is listed under a different scientific na	me on the EPBC Act - Threat	tened Species list.
Name	Threatened	Type of Presence
Birds		
Anseranas semipalmata		
Magpie Goose [978]		Species or species habitat may occur within area
Apus pacificus		
Fork-tailed Swift [678]		Species or species habitat likely to occur within area
Ardea alba		
Great Egret, White Egret [59541]		Species or species habitat likely to occur within area
Ardea ibis		
Cattle Egret [59542]		Species or species habitat may occur within area



<u>Cuculus saturatus</u> Oriental Cuckoo, Himalayan Cuckoo [710]

Gallinago hardwickii Latham's Snipe, Japanese Snipe [863]

Haliaeetus leucogaster White-bellied Sea-Eagle [943]

<u>Hirundapus caudacutus</u> White-throated Needletail [682]

Hirundo rustica Barn Swallow [662] Critically Endangered

Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Species or species habitat may occur within area

Species or species habitat may occur within area

Name	Threatened	Type of Presence
Merops ornatus		
Rainbow Bee-eater [670]		Species or species habitat may occur within area
Monarcha melanopsis		
Black-faced Monarch [609]		Species or species habitat known to occur within area
Monarcha trivirgatus		
Spectacled Monarch [610]		Species or species habitat known to occur within area
Motacilla cinerea		
Grey Wagtail [642]		Species or species habitat may occur within area
Motacilla flava		
Yellow Wagtail [644]		Species or species habitat likely to occur within area
Myiagra cyanoleuca		
Satin Flycatcher [612]		Species or species habitat likely to occur within area
Numenius madagascariensis		
Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Pandion haliaetus		
Osprey [952]		Species or species habitat likely to occur within area
Rhipidura rufifrons		
Rufous Fantail [592]		Species or species habitat known to occur within area
Rostratula benghalensis (sensu lato)		
Painted Snipe [889]	Endangered*	Species or species habitat likely to occur within area
Tringa nebularia		
Common Greenshank, Greenshank [832]		Species or species habitat may occur within area
Reptiles		
Ore ee dulue, ie broetensi		

Crocodylus johnstoni

Freshwater Crocodile, Johnston's Crocodile, Johnston's River Crocodile [1773]

Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Species or species habitat may occur within area

Species or species habitat likely to occur within area

#### **Extra Information**

State and Territory Reserves	[Resource Information]
Name	State
Girringun	QLD
Girringun	QLD
Liefway	QLD
Newcastle Range-The Oaks	QLD

Inva	asive	Species						1	Re	sour	<u>ce Ir</u>	nforma	ation ]	
			 ~ ~		 			1.1						

Weeds reported here are the 20 species of national significance (WoNS), along with other introduced plants that are considered by the States and Territories to pose a particularly significant threat to biodiversity. The following feral animals are reported: Goat, Red Fox, Cat, Rabbit, Pig, Water Buffalo and Cane Toad. Maps from Landscape Health Project, National Land and Water Resouces Audit, 2001.

Name	Status	Type of Presence
Birds		
Columba livia		
Rock Pigeon, Rock Dove, Domestic Pigeon [803]		Species or species habitat likely to occur within area
Lonchura punctulata		
Nutmeg Mannikin [399]		Species or species habitat likely to occur within area
Passer domesticus		
House Sparrow [405]		Species or species habitat likely to occur within area
Streptopelia chinensis		
Spotted Turtle-Dove [780]		Species or species habitat likely to occur within area
Sturnus vulgaris		
Common Starling [389]		Species or species habitat likely to occur within area
Frogs		
Rhinella marina		
Cane Toad [83218]		Species or species habitat likely to occur within area
Mammals		

Bos taurus Domestic Cattle [16]

Species or species habitat likely to occur within area

Canis lupus familiaris Domestic Dog [82654]

Equus caballus Horse [5]

Felis catus Cat, House Cat, Domestic Cat [19]

Oryctolagus cuniculus Rabbit, European Rabbit [128]

Sus scrofa Pig [6]

Vulpes vulpes Red Fox, Fox [18] Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur

Name	Status	Type of Presence
		within area
Plants		
Acacia nilotica subsp. indica		
Prickly Acacia [6196]		Species or species habitat may occur within area
Cenchrus ciliaris		
Buffel-grass, Black Buffel-grass [20213]		Species or species habitat may occur within area
Cryptostegia grandiflora		
Rubber Vine, Rubbervine, India Rubber Vine, India Rubbervine, Palay Rubbervine, Purple Allamanda [18913]		Species or species habitat likely to occur within area
Hymenachne amplexicaulis		Spacios or opacios habitat
Hymenachne, Olive Hymenachne, Water Stargrass, West Indian Grass, West Indian Marsh Grass [31754	.]	Species or species habitat likely to occur within area
Jatropha gossypifolia		
Cotton-leaved Physic-Nut, Bellyache Bush, Cotton-le Physic Nut, Cotton-leaf Jatropha, Black Physic Nut [7507] Lantana camara	af	Species or species habitat likely to occur within area
Lantana, Common Lantana, Kamara Lantana, Large leaf Lantana, Pink Flowered Lantana, Red Flowered Lantana, Red-Flowered Sage, White Sage, Wild Sag [10892] Parkinsonia aculeata		Species or species habitat likely to occur within area
Parkinsonia, Jerusalem Thorn, Jelly Bean Tree, Hors Bean [12301]	Se	Species or species habitat likely to occur within area
Parthenium hysterophorus		
Parthenium Weed, Bitter Weed, Carrot Grass, False Ragweed [19566]		Species or species habitat likely to occur within area
Salvinia molesta		
Salvinia, Giant Salvinia, Aquarium Watermoss, Karib Weed [13665]	а	Species or species habitat likely to occur within area

# Caveat

The information presented in this report has been provided by a range of data sources as acknowledged at the end of the report.

This report is designed to assist in identifying the locations of places which may be relevant in determining obligations under the Environment Protection and Biodiversity Conservation Act 1999. It holds mapped locations of World and National Heritage properties, Wetlands of International and National Importance, Commonwealth and State/Territory reserves, listed threatened, migratory and marine species and listed threatened ecological communities. Mapping of Commonwealth land is not complete at this stage. Maps have been collated from a range of sources at various resolutions.

Not all species listed under the EPBC Act have been mapped (see below) and therefore a report is a general guide only. Where available data supports mapping, the type of presence that can be determined from the data is indicated in general terms. People using this information in making a referral may need to consider the qualifications below and may need to seek and consider other information sources.

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

For species where the distributions are well known, maps are digitised from sources such as recovery plans and detailed habitat studies. Where appropriate, core breeding, foraging and roosting areas are indicated under 'type of presence'. For species whose distributions are less well known, point locations are collated from government wildlife authorities, museums, and non-government organisations; bioclimatic distribution models are generated and these validated by experts. In some cases, the distribution maps are based solely on expert knowledge.

Only selected species covered by the following provisions of the EPBC Act have been mapped:

- migratory and
- marine

The following species and ecological communities have not been mapped and do not appear in reports produced from this database:

- threatened species listed as extinct or considered as vagrants
- some species and ecological communities that have only recently been listed
- some terrestrial species that overfly the Commonwealth marine area
- migratory species that are very widespread, vagrant, or only occur in small numbers

The following groups have been mapped, but may not cover the complete distribution of the species:

- non-threatened seabirds which have only been mapped for recorded breeding sites
- seals which have only been mapped for breeding sites near the Australian continent

Such breeding sites may be important for the protection of the Commonwealth Marine environment.

# Coordinates

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# Acknowledgements

This database has been compiled from a range of data sources. The department acknowledges the following custodians who have contributed valuable data and advice:

-Office of Environment and Heritage, New South Wales

-Department of Environment and Primary Industries, Victoria

-Department of Primary Industries, Parks, Water and Environment, Tasmania

-Department of Environment, Water and Natural Resources, South Australia

-Parks and Wildlife Commission NT, Northern Territory Government

-Department of Environmental and Heritage Protection, Queensland

-Department of Parks and Wildlife, Western Australia

-Environment and Planning Directorate, ACT

-Birdlife Australia

-Australian Bird and Bat Banding Scheme

-Australian National Wildlife Collection

-Natural history museums of Australia

-Museum Victoria

-Australian Museum

-South Australian Museum

-Queensland Museum

-Online Zoological Collections of Australian Museums

-Queensland Herbarium

-National Herbarium of NSW

-Royal Botanic Gardens and National Herbarium of Victoria

-Tasmanian Herbarium

-State Herbarium of South Australia

-Northern Territory Herbarium

-Western Australian Herbarium

-Australian National Herbarium, Atherton and Canberra

-University of New England

-Ocean Biogeographic Information System

-Australian Government, Department of Defence

Forestry Corporation, NSW

-Geoscience Australia

-CSIRO

-Other groups and individuals

The Department is extremely grateful to the many organisations and individuals who provided expert advice and information on numerous draft distributions.

Please feel free to provide feedback via the <u>Contact Us</u> page.

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Department of the Environment and Energy

# **EPBC Act Protected Matters Report**

This report provides general guidance on matters of national environmental significance and other matters protected by the EPBC Act in the area you have selected.

Information on the coverage of this report and qualifications on data supporting this report are contained in the caveat at the end of the report.

Information is available about <u>Environment Assessments</u> and the EPBC Act including significance guidelines, forms and application process details.

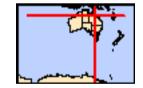
Report created: 03/11/16 14:58:24

Summary Details Matters of NES Other Matters Protected by the EPBC Act Extra Information Caveat Acknowledgements



This map may contain data which are ©Commonwealth of Australia (Geoscience Australia), ©PSMA 2010

Coordinates Buffer: 5.0Km



# Summary

#### Matters of National Environmental Significance

This part of the report summarises the matters of national environmental significance that may occur in, or may relate to, the area you nominated. Further information is available in the detail part of the report, which can be accessed by scrolling or following the links below. If you are proposing to undertake an activity that may have a significant impact on one or more matters of national environmental significance then you should consider the <u>Administrative Guidelines on Significance</u>.

World Heritage Properties:	None
National Heritage Places:	None
Wetlands of International Importance:	None
Great Barrier Reef Marine Park:	None
Commonwealth Marine Area:	None
Listed Threatened Ecological Communities:	1
Listed Threatened Species:	33
Listed Migratory Species:	16

#### Other Matters Protected by the EPBC Act

This part of the report summarises other matters protected under the Act that may relate to the area you nominated. Approval may be required for a proposed activity that significantly affects the environment on Commonwealth land, when the action is outside the Commonwealth land, or the environment anywhere when the action is taken on Commonwealth land. Approval may also be required for the Commonwealth or Commonwealth agencies proposing to take an action that is likely to have a significant impact on the environment anywhere.

The EPBC Act protects the environment on Commonwealth land, the environment from the actions taken on Commonwealth land, and the environment from actions taken by Commonwealth agencies. As heritage values of a place are part of the 'environment', these aspects of the EPBC Act protect the Commonwealth Heritage values of a Commonwealth Heritage place. Information on the new heritage laws can be found at http://www.environment.gov.au/heritage

A <u>permit</u> may be required for activities in or on a Commonwealth area that may affect a member of a listed threatened species or ecological community, a member of a listed migratory species, whales and other cetaceans, or a member of a listed marine species.

Commonwealth Land:	None
Commonwealth Heritage Places:	None
Listed Marine Species:	23
Whales and Other Cetaceans:	None
Critical Habitats:	None
Commonwealth Reserves Terrestrial:	None
Commonwealth Reserves Marine:	None

#### **Extra Information**

This part of the report provides information that may also be relevant to the area you have nominated.

State and Territory Reserves:	4
Regional Forest Agreements:	None
Invasive Species:	24
Nationally Important Wetlands:	None
Key Ecological Features (Marine)	None

# Details

## Matters of National Environmental Significance

#### Listed Threatened Ecological Communities

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

[Resource Information]

known to occur within area

Name	Status	Type of Presence
Broad leaf tea-tree (Melaleuca viridiflora) woodlands in high rainfall coastal north Queensland	Endangered	Community may occur within area
Listed Threatened Species		[Resource Information]
Name	Status	Type of Presence
Birds		
<u>Calidris ferruginea</u> Curlew Sandpiper [856]	Critically Endangered	Species or species habitat may occur within area
Casuarius casuarius johnsonii Southern Cassowary, Australian Cassowary, Double- wattled Cassowary [25986]	Endangered	Species or species habitat known to occur within area
<u>Erythrotriorchis radiatus</u> Red Goshawk [942]	Vulnerable	Species or species habitat likely to occur within area
<u>Erythrura gouldiae</u> Gouldian Finch [413]	Endangered	Species or species habitat may occur within area
Numenius madagascariensis Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Poephila cincta cincta Southern Black-throated Finch [64447]	Endangered	Species or species habitat

Rostratula australis

Australian Painted Snipe [77037]	Endangered	Species or species habitat likely to occur within area
<u>Tyto novaehollandiae kimberli</u> Masked Owl (northern) [26048]	Vulnerable	Species or species habitat likely to occur within area
Frogs		
Litoria dayi Australian Lace-lid, Lace-eyed Tree Frog [86707]	Endangered	Species or species habitat likely to occur within area
<u>Litoria nannotis</u> Waterfall Frog, Torrent Tree Frog [1817]	Endangered	Species or species habitat may occur within area

Name	Status	Type of Presence
<u>Litoria rheocola</u> Common Mistfrog [1802]	Endangered	Species or species habitat likely to occur within area
Mammals		
<u>Dasyurus hallucatus</u> Northern Quoll, Digul [331]	Endangered	Species or species habitat likely to occur within area
Dasyurus maculatus gracilis Spotted-tailed Quoll (North Queensland), Yarri [64475]	Endangered	Species or species habitat likely to occur within area
Hipposideros semoni Semon's Leaf-nosed Bat, Greater Wart-nosed Horseshoe-bat [180]	Endangered	Species or species habitat may occur within area
<u>Macroderma gigas</u> Ghost Bat [174]	Vulnerable	Species or species habitat likely to occur within area
Mesembriomys gouldii rattoides Black-footed Tree-rat (north Queensland), Shaggy Rabbit-rat [87620]	Vulnerable	Species or species habitat likely to occur within area
Petauroides volans Greater Glider [254]	Vulnerable	Species or species habitat known to occur within area
<u>Petrogale sharmani</u> Mount Claro Rock Wallaby, Sharman's Rock Wallaby [59281]	Vulnerable	Species or species habitat known to occur within area
Phascolarctos cinereus (combined populations of Qld, Koala (combined populations of Queensland, New South Wales and the Australian Capital Territory) [85104]	NSW and the ACT) Vulnerable	Species or species habitat known to occur within area
Pteropus conspicillatus Spectacled Flying-fox [185]	Vulnerable	Species or species habitat known to occur within area
Pteropus poliocephalus Grey-headed Flying-fox [186]	Vulnerable	Foraging, feeding or related behaviour likely to occur within area
<u>Rhinolophus robertsi</u> Large-eared Horseshoe Bat, Greater Large-eared Horseshoe Bat [87639]	Endangered	Species or species habitat known to occur within area
Saccolaimus saccolaimus nudicluniatus Bare-rumped Sheath-tailed Bat, Bare-rumped Sheathtail Bat [66889]	Critically Endangered	Species or species habitat likely to occur within area
Plants		
<u>Acacia crombiei</u> Pink Gidgee [10927]	Vulnerable	Species or species habitat may occur within area
Bulbophyllum globuliforme Miniature Moss-orchid, Hoop Pine Orchid [6649]	Vulnerable	Species or species habitat may occur within area
<u>Cajanus mareebensis</u> [8635]	Endangered	Species or species habitat likely to occur within area
Dichanthium setosum bluegrass [14159]	Vulnerable	Species or species habitat likely to occur within area
<u>Lindsaea pulchella var. blanda</u> [20842]	Vulnerable	Extinct within area

Name	Status	Type of Presence
<u>Marsdenia brevifolia</u> [64585]	Vulnerable	Species or species habitat likely to occur within area
Phaius australis Lesser Swamp-orchid [5872]	Endangered	Species or species habitat likely to occur within area
Tephrosia leveillei [16946]	Vulnerable	Species or species habitat likely to occur within area
Reptiles		
Egernia rugosa		
Yakka Skink [1420]	Vulnerable	Species or species habitat may occur within area
<u>Lerista vittata</u> Mount Cooper Striped Lerista [1308]	Vulnerable	Species or species habitat
		likely to occur within area
Listed Migratory Species		[Resource Information]
* Species is listed under a different scientific name or	the EPBC Act - Threatened	
Name	the EPBC Act - Threatened Threatened	d Species list. Type of Presence
Name Migratory Marine Birds		
Name		
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678]		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678] Migratory Marine Species		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678]		Type of Presence Species or species habitat
Name Migratory Marine Birds <u>Apus pacificus</u> Fork-tailed Swift [678] Migratory Marine Species <u>Crocodylus porosus</u>		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774]		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species		Type of Presence Species or species habitat likely to occur within area Species or species habitat
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within area
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus Oriental Cuckoo, Horsfield's Cuckoo [86651]		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within area
Name Migratory Marine Birds Apus pacificus Fork-tailed Swift [678] Migratory Marine Species Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Migratory Terrestrial Species Cuculus optatus Oriental Cuckoo, Horsfield's Cuckoo [86651]		Type of PresenceSpecies or species habitat likely to occur within areaSpecies or species habitat likely to occur within areaSpecies or species habitat known to occur within areaSpecies or species habitat known to occur within areaSpecies or species habitat

Barn Swallow [662]

Monarcha melanopsis Black-faced Monarch [609]

Monarcha trivirgatus Spectacled Monarch [610]

Motacilla cinerea Grey Wagtail [642]

Motacilla flava Yellow Wagtail [644]

Myiagra cyanoleuca Satin Flycatcher [612]

<u>Rhipidura rufifrons</u> Rufous Fantail [592] Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat known to occur within area

Threatened	Type of Presence
Critically Endangered	Species or species habitat may occur within area
	Species or species habitat may occur within area
Critically Endangered	Species or species habitat may occur within area
	Species or species habitat likely to occur within area
	Species or species habitat may occur within area

### Other Matters Protected by the EPBC Act

Listed Marine Species		[Resource Information]
* Species is listed under a different scientific na	me on the EPBC Act - Threat	tened Species list.
Name	Threatened	Type of Presence
Birds		
Anseranas semipalmata		
Magpie Goose [978]		Species or species habitat may occur within area
Apus pacificus		
Fork-tailed Swift [678]		Species or species habitat likely to occur within area
Ardea alba		
Great Egret, White Egret [59541]		Species or species habitat likely to occur within area
Ardea ibis		
Cattle Egret [59542]		Species or species habitat may occur within area



<u>Cuculus saturatus</u> Oriental Cuckoo, Himalayan Cuckoo [710]

Gallinago hardwickii Latham's Snipe, Japanese Snipe [863]

Haliaeetus leucogaster White-bellied Sea-Eagle [943]

<u>Hirundapus caudacutus</u> White-throated Needletail [682]

Hirundo rustica Barn Swallow [662] Critically Endangered

Species or species habitat may occur within area

Species or species habitat known to occur within area

Species or species habitat may occur within area

Species or species habitat likely to occur within area

Species or species habitat may occur within area

Species or species habitat may occur within area

Name	Threatened	Type of Presence
Merops ornatus		
Rainbow Bee-eater [670]		Species or species habitat may occur within area
Monarcha melanopsis		
Black-faced Monarch [609]		Species or species habitat known to occur within area
Monarcha trivirgatus		
Spectacled Monarch [610]		Species or species habitat known to occur within area
Motacilla cinerea		
Grey Wagtail [642]		Species or species habitat may occur within area
Motacilla flava		
Yellow Wagtail [644]		Species or species habitat likely to occur within area
Myiagra cyanoleuca		
Satin Flycatcher [612]		Species or species habitat likely to occur within area
Numenius madagascariensis		
Eastern Curlew, Far Eastern Curlew [847]	Critically Endangered	Species or species habitat may occur within area
Pandion haliaetus		
Osprey [952]		Species or species habitat likely to occur within area
Rhipidura rufifrons		
Rufous Fantail [592]		Species or species habitat known to occur within area
Rostratula benghalensis (sensu lato)		
Painted Snipe [889]	Endangered*	Species or species habitat likely to occur within area
Tringa nebularia		
Common Greenshank, Greenshank [832]		Species or species habitat may occur within area
Reptiles		
Ore ee dulue, ie broetensi		

Crocodylus johnstoni

Freshwater Crocodile, Johnston's Crocodile, Johnston's River Crocodile [1773]

Crocodylus porosus Salt-water Crocodile, Estuarine Crocodile [1774] Species or species habitat may occur within area

Species or species habitat likely to occur within area

#### **Extra Information**

State and Territory Reserves	[Resource Information]
Name	State
Girringun	QLD
Girringun	QLD
Liefway	QLD
Newcastle Range-The Oaks	QLD

Inva	asive	Species						1	Re	sour	<u>ce Ir</u>	nforma	ation ]	
			 ~ ~		 			1.1						

Weeds reported here are the 20 species of national significance (WoNS), along with other introduced plants that are considered by the States and Territories to pose a particularly significant threat to biodiversity. The following feral animals are reported: Goat, Red Fox, Cat, Rabbit, Pig, Water Buffalo and Cane Toad. Maps from Landscape Health Project, National Land and Water Resouces Audit, 2001.

Name	Status	Type of Presence
Birds		
Columba livia		
Rock Pigeon, Rock Dove, Domestic Pigeon [803]		Species or species habitat likely to occur within area
Lonchura punctulata		
Nutmeg Mannikin [399]		Species or species habitat likely to occur within area
Passer domesticus		
House Sparrow [405]		Species or species habitat likely to occur within area
Streptopelia chinensis		
Spotted Turtle-Dove [780]		Species or species habitat likely to occur within area
Sturnus vulgaris		
Common Starling [389]		Species or species habitat likely to occur within area
Frogs		
Rhinella marina		
Cane Toad [83218]		Species or species habitat likely to occur within area
Mammals		

Bos taurus Domestic Cattle [16]

Species or species habitat likely to occur within area

Canis lupus familiaris Domestic Dog [82654]

Equus caballus Horse [5]

Felis catus Cat, House Cat, Domestic Cat [19]

Feral deer Feral deer species in Australia [85733]

Oryctolagus cuniculus Rabbit, European Rabbit [128]

Rattus rattus Black Rat, Ship Rat [84] Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur within area

Species or species habitat likely to occur

Name	Status	Type of Presence
		within area
Sus scrofa		
Pig [6]		Species or species habitat likely to occur within area
Vulpes vulpes		
Red Fox, Fox [18]		Species or species habitat likely to occur within area
Plants		
Acacia nilotica subsp. indica		
Prickly Acacia [6196]		Species or species habitat may occur within area
Cenchrus ciliaris		
Buffel-grass, Black Buffel-grass [20213]		Species or species habitat may occur within area
Cryptostegia grandiflora		
Rubber Vine, Rubbervine, India Rubber Vine, India Rubbervine, Palay Rubbervine, Purple Allamanda [18913]		Species or species habitat likely to occur within area
Hymenachne amplexicaulis		
Hymenachne, Olive Hymenachne, Water Stargrass, West Indian Grass, West Indian Marsh Grass [31754		Species or species habitat likely to occur within area
Jatropha gossypifolia		
Cotton-leaved Physic-Nut, Bellyache Bush, Cotton-le Physic Nut, Cotton-leaf Jatropha, Black Physic Nut [7507] Lantana camara	eaf	Species or species habitat likely to occur within area
Lantana, Common Lantana, Kamara Lantana, Large leaf Lantana, Pink Flowered Lantana, Red Flowered Lantana, Red-Flowered Sage, White Sage, Wild Sag [10892] Parkinsonia aculeata		Species or species habitat likely to occur within area
Parkinsonia, Jerusalem Thorn, Jelly Bean Tree, Hors Bean [12301]	se	Species or species habitat likely to occur within area
Parthenium hysterophorus Parthenium Weed, Bitter Weed, Carrot Grass, False Ragweed [19566]		Species or species habitat likely to occur within area
Salvinia molesta		
Solvinia Ciant Solvinia Aquarium Watermana Karib		Species or species babitat

Salvinia, Giant Salvinia, Aquarium Watermoss, Kariba

Species or species habitat likely to occur within area

Weed [13665]

# Caveat

The information presented in this report has been provided by a range of data sources as acknowledged at the end of the report.

This report is designed to assist in identifying the locations of places which may be relevant in determining obligations under the Environment Protection and Biodiversity Conservation Act 1999. It holds mapped locations of World and National Heritage properties, Wetlands of International and National Importance, Commonwealth and State/Territory reserves, listed threatened, migratory and marine species and listed threatened ecological communities. Mapping of Commonwealth land is not complete at this stage. Maps have been collated from a range of sources at various resolutions.

Not all species listed under the EPBC Act have been mapped (see below) and therefore a report is a general guide only. Where available data supports mapping, the type of presence that can be determined from the data is indicated in general terms. People using this information in making a referral may need to consider the qualifications below and may need to seek and consider other information sources.

For threatened ecological communities where the distribution is well known, maps are derived from recovery plans, State vegetation maps, remote sensing imagery and other sources. Where threatened ecological community distributions are less well known, existing vegetation maps and point location data are used to produce indicative distribution maps.

For species where the distributions are well known, maps are digitised from sources such as recovery plans and detailed habitat studies. Where appropriate, core breeding, foraging and roosting areas are indicated under 'type of presence'. For species whose distributions are less well known, point locations are collated from government wildlife authorities, museums, and non-government organisations; bioclimatic distribution models are generated and these validated by experts. In some cases, the distribution maps are based solely on expert knowledge.

Only selected species covered by the following provisions of the EPBC Act have been mapped:

- migratory and
- marine

The following species and ecological communities have not been mapped and do not appear in reports produced from this database:

- threatened species listed as extinct or considered as vagrants
- some species and ecological communities that have only recently been listed
- some terrestrial species that overfly the Commonwealth marine area
- migratory species that are very widespread, vagrant, or only occur in small numbers

The following groups have been mapped, but may not cover the complete distribution of the species:

- non-threatened seabirds which have only been mapped for recorded breeding sites
- seals which have only been mapped for breeding sites near the Australian continent

Such breeding sites may be important for the protection of the Commonwealth Marine environment.

# Coordinates

-18.87006 144.148,-18.8791 144.148,-18.87909 144.1369,-18.90823 144.1233,-18.91321 144.1231,-18.91545 144.1259,-18.90989 144.1736,-18.92307 144.2417,-18.92151 144.3212,-18.92745 144.4211,-18.92693 144.4254,-18.92767 144.4255,-18.95179 144.9292,-18.97166 145.0154,-18.90435 145.3466,-18.84082 145.8332,-18.84976 145.8346,-18.91328 145.348,-18.95105 145.1765,-18.98079 145.0149,-18.96079 144.9283,-18.93055 144.3214,-18.93213 144.2409,-18.91898 144.1732,-18.9246 144.1236,-18.91696 144.1138,-18.90601 144.114,-18.87068 144.1314,-18.87006 144.148

# Acknowledgements

This database has been compiled from a range of data sources. The department acknowledges the following custodians who have contributed valuable data and advice:

-Office of Environment and Heritage, New South Wales

-Department of Environment and Primary Industries, Victoria

-Department of Primary Industries, Parks, Water and Environment, Tasmania

-Department of Environment, Water and Natural Resources, South Australia

-Parks and Wildlife Commission NT, Northern Territory Government

-Department of Environmental and Heritage Protection, Queensland

-Department of Parks and Wildlife, Western Australia

-Environment and Planning Directorate, ACT

-Birdlife Australia

-Australian Bird and Bat Banding Scheme

-Australian National Wildlife Collection

-Natural history museums of Australia

-Museum Victoria

-Australian Museum

-South Australian Museum

-Queensland Museum

-Online Zoological Collections of Australian Museums

-Queensland Herbarium

-National Herbarium of NSW

-Royal Botanic Gardens and National Herbarium of Victoria

-Tasmanian Herbarium

-State Herbarium of South Australia

-Northern Territory Herbarium

-Western Australian Herbarium

-Australian National Herbarium, Atherton and Canberra

-University of New England

-Ocean Biogeographic Information System

-Australian Government, Department of Defence

Forestry Corporation, NSW

-Geoscience Australia

-CSIRO

-Other groups and individuals

The Department is extremely grateful to the many organisations and individuals who provided expert advice and information on numerous draft distributions.

Please feel free to provide feedback via the <u>Contact Us</u> page.

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# Appendix B

**PROTECTED MATTERS SEARCH TOOL – OPTIONS COLLATED** 

#### PMST results for each Corridor Option

MNES Category	Scientific name	Common name	Status	Option A	Option B	Option C
TECs	Broad leaf tea-tree ( <i>Melaleuca viridiflora</i> ) Queensland	E	х	х	х	
Threatened species						
•	Calidris ferruginea	Curlew Sandpiper	CE (M)	Х	х	Х
	Casuarius casuarius johnsonii	Southern Cassowary	E	Х	х	Х
	Erythrotriorchis radiatus	Red Goshawk	V	х	х	Х
Diada	Erythrura gouldiae	Gouldian Finch	E	х	х	Х
Birds	Numenius madagascariensis	Eastern Curlew	CE	х	х	Х
	Poephila cincta cincta	Southern Black-throated Finch	E	х	х	Х
	, Rostratula australis	Australian Painted Snipe	E	х	х	Х
	Tyto novaehollandiae kimberli	Masked Owl (northern)	V	х	х	Х
	Litoria dayi	Australian Lace-lid	E	х	х	Х
Frogs	Litoria nannotis	Waterfall Frog	E	х	х	Х
5	Litoria rheocola	Common Mistfrog	E	х	х	Х
	Dasyurus hallucatus	Northern Quoll	E	x	X	X
	Dasyurus maculatus gracilis	Spotted-tailed Quoll (North Queensland)	E	x	X	X
	Hipposideros semoni	Semon's Leaf-nosed Bat	E	X	X	X
	Macroderma gigas	Ghost Bat	V	X	X	X
	Mesembriomys gouldii rattoides	Black-footed Tree-rat (north Queensland)	V	X	X	X
	Petauroides volans	Greater Glider	V	x	X	X
Mammals	Petrogale sharmani	Mount Claro Rock Wallaby	V	X	X	X
	Phascolarctos cinereus	Koala	V	X	X	X
	Pteropus conspicillatus	Spectacled Flying-fox	V	x	X	X
	Pteropus poliocephalus	Grey-headed Flying-fox	V	x	X	X
	Rhinolophus robertsi	Large-eared Horseshoe Bat	E	X	X	X
	Saccolaimus saccolaimus nudicluniatus	Bare-rumped Sheath-tailed Bat	CE	X	x	X
	Acacia crombiei	Pink Gidgee	V	x	x	X
	Bulbophyllum globuliforme	Miniature Moss-orchid	V	x	x	X
	Cajanus mareebensis		E	x	x	X
	Dichanthium setosum	Bluegrass	V	x	X	X
Plants	Lindsaea pulchella var. blanda		V	x	x	X
	Marsdenia brevifolia		V	X	x	X
	Phaius australis	Lesser Swamp-orchid	E	x	x	X
	Tephrosia leveillei		V	X	x	X
	Egernia rugosa	Yakka Skink	V	x	x	X
Reptiles	Lerista vittata	Mount Cooper Striped Lerista	V	x	X	X
Migratory (marine)	Apus pacificus	Fork-tailed Swift	M	x	X	X
ivingratory (marine)	Cuculus optatus	Oriental Cuckoo	M	x	x	X
	Hirundapus caudacutus	White-throated Needletail	M	x	x	X
	Hirundo rustica	Barn Swallow	M	X	x	X
	Monarcha melanopsis	Black-faced Monarch	M	X	X	X
Migratory (terrestrial)		Spectacled Monarch	M	x	x	X
ivingratory (terrestrial)	Motacilla cinerea	Grey Wagtail	M	X	x	X
	Motacilla flava	Yellow Wagtail	M	X	X	X
	Myiagra cyanoleuca	Satin Flycatcher	M	X	X	X
	Rhipidura rufifrons	Rufous Fantail	M	x	X	X
	Calidris ferruginea	Curlew Sandpiper	M (CE)	X	X	X
	Gallinago hardwickii	Latham's Snipe	M	X	X	х Х
Migratory (wetland)	Numenius madagascariensis	Eastern Curlew	M (CE)	X		X
(wettand)	Pandion haliaetus	Osprey	M	X	X X	х Х
	Tringa nebularia	Common Greenshank	M	^		
	n inga nebulaha		111	-	Х	Х

Other MNES	Option A	Option B	Option C
World Heritage properties - Wet Tropics of Queensland	Х	-	-
National Heritage properties (natural) - Wet Tropics of Queensland	х	-	-
National Heritage properties (indigenous) - Wet Tropics World Heritage Area (Indigenous Values)	х	-	-

# Appendix C

WILDLIFE ONLINE SEARCHES

Search Crit(Species List for a Specified Point

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.8813
Longitude:	144.1254
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:49:02
Date extracted:	Tuesday 08 Nov 2016 13:50:03
The number of records retrieved = 287	

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Feedback about Wildlife Online should be emailed to wildlife.online@science.dsitia.qld.gov.au

Description of the CODES

I - Y indicates that the taxon is introduced to Queensland and has naturalised.

Indicates the Queensland conservation status of each taxon under the Nature Conservation Act

 Q 1992. The codes are Extinct in the Wild (PE)
 Endangered (E)

 Vulnerable (V)
 Near Threatened (NT)
 Least Concern (C) or Not Protected ().

 A Indicates the Australian conservation status of each taxon under the Environment Protection and Biodiversity Conservation Act 1999. The values of EPBC are Conservation Dependent (CD)
 Critically Endangered (CE)
 Endangered (E)

Records – The first number indicates the total number of reco Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting ReSpecimen Recor	ds
plants	cycads	Cycadaceae	Cycas cairnsiana		V	V	5 5	

Search Crit/Species List for a Specified Point

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.9068
Longitude:	144.3146
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:49:26
Date extracted:	Tuesday 08 Nov 2016 13:50:11
The number of records retrieved = 167	

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Description of the CODES

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1-	Y indicates that the taxon is introduced to Indicates the Queensland conservation status of each taxon under the Nature Conservation Act 1992. The codes are	o Queensland and has naturalise	ed.	
Q -	Extinct in the Wild (PE)	Endangered (E)		
	Vulnerable (V)	Near Threatened (NT)	Least Concern (C) or Not	Protected ().
A -	Indicates the Australian conservation stat Conservation Dependent (CD)	us of each taxon under the Envi Critically Endangered (CE)	ronment Protection and Bi Endangered (E)	odiversity Conservation Act 1999. The values of EPBC are Extinct (EX)

Records – The first number indicates the total numb Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom Class	Family	Scientific Name	Common Name	NC Act	EPBC Act Si	ighting ReSpecimen Records
animals birds	Estrildidae	Poephila cincta cincta	black-throated finch (white-rumped subspecies)	E	E	3 0

Search Criteria: Species List for a Specified Point

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.9175
Longitude:	144.5056
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:49:45
Date extracted:	Tuesday 08 Nov 2016 13:50:07
The number of records retrieved = 353	-

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Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act Sig	hting Re Speci	men Records
animals	birds	Estrildidae	Poephila cincta cincta	black-throated finch (white-rumped subspecies)	E	E	5	0
animals	mammals	Muridae	Mesembriomys gouldii	black-footed tree-rat	С	V	1	0
animals	mammals	Tachyglossidae	Tachyglossus aculeatus	short-beaked echidna	SL		1	0

Search Criti Species List for a Specified Point

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.9253
Longitude:	144.697
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:50:05
Date extracted:	Tuesday 08 Nov 2016 14:00:43
The number of records retrieved = 198	

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Critically Endangered (CE) Endangered (E)

Records - The first number indicates the total number of recc Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting ReSpecimen Records
animals birds	Estrildidae	Poephila cincta cincta	black-throated finch (white-rumped subspecies)	E	E	2 0

Search Crit Species List for a Specified Point

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.9309
Longitude:	144.8886
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:50:31
Date extracted:	Tuesday 08 Nov 2016 14:00:17
The number of records retrieved = 347	

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 Vulnerable (V)
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A - Indicates the Australian conservation status of each taxon under the Environment Protection and Biodiversity Conservation Act 1999. The values of EPBC are Conservation Dependent (CD) Critically Endangered (CE) Endangered (E) Extinct (EX)

#### Records – The first number indicates the total number of recc Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act S	ighting Re Specim	en Records
animals	birds	Estrildidae	Erythrura gouldiae	Gouldian finch	E	E	1	0
animals	birds	Estrildidae	Poephila cincta cincta	black-throated finch (white-rumped subspecies)	E	E	2	0
animals	birds	Laridae	Hydroprogne caspia	Caspian tern	SL		1	0
animals	birds	Scolopacidae	Calidris ruficollis	red-necked stint	SL		1	0

Search Criteria: Species List for a Specified Point

Species:	All		
Type:	Native		
Status:	All		
Records:	All		
Date:	All		
Latitude:	-18.9345		
Longitude:	145.0795		
Distance:	18		
Email:	datov@pbworld.com		
Date submitted:	Tuesday 08 Nov 2016 13:50:54		
Date extracted:	Tuesday 08 Nov 2016 14:00:10		
The number of records retrieved = 335			

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Description of the CODES

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	Vulnerable (V)	Near Threatened (NT)	Least Concern (C) or Not Prote	ected ().	
A -	Indicates the Australian conservation status of each	n taxon under the Environment	Protection and Biodiversity Con	servation Act 1999. The values of EPBC a	re
	Conservation Dependent (CD)	Critically Endangered (CE)	Endangered (E)	Extinct (EX)	

Records – The first number indicates the total number of recoi Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting Re Specir	nen Records
animals	birds	Estrildidae	Poephila cincta cincta	black-throated finch (white-rumped subspecies)	E	E	1	0
animals	birds	Estrildidae	Erythrura gouldiae	Gouldian finch	E	E	1	0
animals	birds	Laridae	Hydroprogne caspia	Caspian tern	SL		1	0
animals	birds	Rhipiduridae	Rhipidura rufifrons	rufous fantail	SL		1	0
animals	birds	Scolopacidae	Calidris ruficollis	red-necked stint	SL		1	0
plants	higher dicots	Myrtaceae	Leptospermum pallidum		NT		5	5
plants	higher dicots	Sapindaceae	Arytera dictyoneura		NT		1	1
plants	monocots	Poaceae	Lepturus minutus		V		1	1

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.8973
Longitude:	145.2662
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:51:19
Date extracted:	Tuesday 08 Nov 2016 14:00:46
The number of records retrieved = 89	

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### Description of the CODES

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Q -

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> Indicates the Queensland conservation status of each taxon under the Nature Conservation Act

1992. The codes are Extinct in the Wild (PE)

Endangered (E) Vulnerable (V) Near Threatened (NT)

Least Concern (C) or Not Protected (). Indicates the Australian conservation status of each taxon under the Environment Protection and Biodiversity Conservation Act 1999. The values of EPBC are

Α-Conservation Dependent (CD) Critically Endangered (CE) Endangered (E) Extinct (EX)

### Records - The first number indicates the total number of recc Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting Re Spec	imen Records
animals	birds	Rhipiduridae	Rhipidura rufifrons	rufous fantail	SL		1	0
plants	higher dicots	Apiaceae	Oenanthe javanica		NT		1	1
plants	higher dicots	Sapindaceae	Arytera dictyoneura		NT		1	1

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.8737
Longitude:	145.4555
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:51:40
Date extracted:	Tuesday 08 Nov 2016 14:00:49
The number of records retrieved = 95	

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Least Concern (C) or Not Protected ().

Α-Indicates the Australian conservation status of each taxon under the Environment Protection and Biodiversity Conservation Act 1999. The values of EPBC are Conservation Dependent (CD) Critically Endangered (CE) Endangered (E) Extinct (EX)

Records - The first number indicates the total number of recor Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting Re Specime	en Records
animals	birds	Apodidae	Hirundapus caudacutus	white-throated needletail	SL		1	0
animals	mammals	Phascolarctidae	Phascolarctos cinereus	koala	V	V	1	0
animals	mammals	Tachyglossidae	Tachyglossus aculeatus	short-beaked echidna	SL		1	0
plants	higher dicots	Apiaceae	Oenanthe javanica		NT		1	1

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.8581
Longitude:	145.6465
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:52:00
Date extracted:	Tuesday 08 Nov 2016 14:00:03
The number of records retrieved = 508	

Disclaimer

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As the DSIT it is possible the information given is not complete. The information provided should only be used for the project for which it was requested and it should be appropriately acknowledged as being derived from Wildlife Online when it is used.

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Feedback about Wildlife Online should be emailed to wildlife.online@science.dsitia.qld.gov.au

Description of the CODES

### I - Y indicates that the taxon is introduced to Queensland and has naturalised.

Indicates the Queensland conservation status of each taxon under the Nature Conservation Act

- Q 1992. The codes are Extinct in the Wild (PE) Endangered (E)
- Vulnerable (V) Near Threatened (NT) Least Concern (C) or Not Protected ().
- A Indicates the Australian conservation status of each taxon under the Environment Protection and Biodiversity Conservation Act 1999. The values of EPBC are Conservation Dependent (CD) Critically Endangered (CE) Endangered (E) Extinct (EX)

Records – The first number indicates the total number of reci Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting ReSpeci	men Records
animals	birds	Apodidae	Hirundapus caudacutus	white-throated needletail	SL		1	0
animals	birds	Monarchidae	Symposiachrus trivirgatus	spectacled monarch	SL		1	0
animals	mammals	Macropodidae	Petrogale sharmani	Sharman's rock-wallaby	V	V	34	30
animals	mammals	Phascolarctidae	Phascolarctos cinereus	koala	V	V	2	0
animals	mammals	Pseudocheiridae	Petauroides volans minor	northern greater glider	С	V	1	0
plants	higher dicots	Mimosaceae	Acacia tingoorensis		V		4	4
plants	monocots	Orchidaceae	Corybas cerasinus		NT		2	2

\_\_\_\_\_

Species:	All
Туре:	Native
Status:	All
Records:	All
Date:	All
Latitude:	-18.8445
Longitude:	145.8358
Distance:	18
Email:	datov@pbworld.com
Date submitted:	Tuesday 08 Nov 2016 13:52:20
Date extracted:	Tuesday 08 Nov 2016 14:00:14
The number of records retrieved = 615	

Disclaimer

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As the DSI1 it is possible the information given is not complete. The information provided should only be used for the project for which it was requested and it should be appropriately acknowledged as being derived from Wildlife Online when it is used.

The State c nor accept responsibility for this information. Persons should satisfy themselves through independent means as to the accuracy and completeness of this information.

No statem∉ representations or warranties are made about the accuracy or completeness of this information. The State of Queensland disclaims all responsibil liability in negligence) for all expenses losses damages and costs you may incur as a result of the information being inaccurate or incomplete in any way for any reason.

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#### Description of the CODES

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I - Y indicates that the taxon is introduced to Queensland and has naturalised.

Indicates the Queensland conservation status of each taxon under the Nature Conservation Act

Q - 1992. The codes are Extinct in the Wild (PE) Endangered (E)

	Vulnerable (V)	Near Threatened (NT)	Least Concern (C) or Not Protected ().	
Α-	Indicates the Australian conservation status of each	n taxon under the Environment	Protection and Biodiversity Conservation Act 1999.	The values of EPBC are
	Conservation Dependent (CD)	Critically Endangered (CE)	Endangered (E)	Extinct (EX)

Records – The first number indicates the total number of reco Confirmed or Specimens). The second number located after the / indicates the number of specimen records for the taxon.

Kingdom	Class	Family	Scientific Name	Common Name	NC Act	EPBC Act	Sighting Re Spe	cimen Records
animals	birds	Cacatuidae	Calyptorhynchus lathami erebus	glossy black-cockatoo (northern)	V		1	0
animals	birds	Casuariidae	Casuarius casuarius johnsonii (southern population)	southern cassowary (southern population)	E	E	2	0
animals	birds	Cuculidae	Cuculus optatus	oriental cuckoo	SL		1	0
animals	birds	Monarchidae	Monarcha melanopsis	black-faced monarch	SL		20	0
animals	birds	Monarchidae	Symposiachrus trivirgatus	spectacled monarch	SL		32	0
animals	birds	Rhipiduridae	Rhipidura rufifrons	rufous fantail	SL		9	0
animals	mammals	Macropodidae	Petrogale sharmani	Sharman's rock-wallaby	V	V	34	30
animals	mammals	Petauridae	Petaurus gracilis	mahogany glider	E	E	15	0
animals	mammals	Phascolarctidae	Phascolarctos cinereus	koala	V	V	3	0
animals	mammals	Pseudocheiridae	Petauroides volans minor	northern greater glider	С	V	1	0
plants	higher dicots	Apiaceae	Oenanthe javanica		NT		1	1
plants	higher dicots	Asteraceae	Glossocardia orthochaeta		E		1	1
plants	higher dicots	Mimosaceae	Acacia tingoorensis		V		4	4
plants	higher dicots	Myrtaceae	Corymbia leptoloma		V	V	1	1
plants	monocots	Orchidaceae	Corybas cerasinus		NT		2	2

# Appendix D

WILDLIFE ONLINE DATA COLLATION

### Wildlife Online data combined for records searches

Class	Scientific Name	Common Name	NC Act	EPBC Act
Birds	Calidris ruficollis	Red-necked Stint	SL	М
Birds	Calyptorhynchus lathami erebus	Glossy Black-cockatoo (northern)	V	-
Birds	Casuarius casuarius johnsonii (southern population)	Southern Cassowary (southern population)	E	E
Birds	Cuculus optatus	Oriental cuckoo	SL	М
Birds	Erythrura gouldiae	Gouldian Finch	E	E
Birds	Hirundapus caudacutus	White-throated Needletail	SL	М
Birds	Hydroprogne caspia	Caspian Tern	SL	М
Birds	Monarcha melanopsis	Black-faced Monarch	SL	М
Birds	Poephila cincta cincta	Black-throated Finch (white-rumped subspecies)	E	E
Birds	Rhipidura rufifrons	Rufous Fantail	SL	М
Birds	Symposiachrus trivirgatus	Spectacled Monarch	SL	М
Mammals	Mesembriomys gouldii	Black-footed Tree-rat	-	V
Mammals	Petauroides volans minor	Northern Greater Glider	-	V
Mammals	Petaurus gracilis	Mahogany Glider	E	E
Mammals	Petrogale sharmani	Sharman's Rock-wallaby	V	V
Mammals	Phascolarctos cinereus	Koala	V	V
Mammals	Tachyglossus aculeatus	Short-beaked Echidna	SL	-
Plants	Acacia tingoorensis		V	-
Plants	Arytera dictyoneura		NT	-
Plants	Cycas cairnsiana		V	V
Plants	Corybas cerasinus		NT	-
Plants	Corymbia leptoloma		V	V
Plants	Glossocardia orthochaeta		E	-
Plants	Leptospermum pallidum		NT	-
Plants	Lepturus minutus		V	-
Plants	Oenanthe javanica		NT	-

## Appendix E

**DSITI AND ALA SPECIES RECORDS SEARCHES** 

### Kidston Corridor Options - Threatened species records searches

Class	Scientific Name	Common Name	EPBC Act	NC Act	Database search	Records search	DSITI Comments
Threatened flora		1					
Plant	Acacia crombiei	Pink Gidgee	V	V	PMST	DSITI and ALA	
Plant	Acacia tingoorensis	-	-	V	Wildlife Online	DSITI and ALA	
Plant	Arytera dictyoneura	-	-	NT	Wildlife Online	DSITI and ALA	
Plant	Bulbophyllum qlobuliforme	Miniature Moss-orchid	V	NT	PMST	ALA	Confidential species; unable to download records
Plant	Cajanus mareebensis	-	E	E	PMST	DSITI and ALA	
Plant	Corybas cerasinus	-	-	NT	Wildlife Online	ALA	Confidential species; unable to download records
Plant	Corymbia leptoloma	Yellowjacket	V	V	Wildlife Online	DSITI and ALA	
Other	Cycas cairnsiana	-	V	V	Wildlife Online	ALA	Confidential species; unable to download records
Plant	Dichanthium setosum	Bluegrass	V	-	PMST	DSITI and ALA	
Plant	Glossocardia orthochaeta	-		F	Wildlife Online	DSITI and ALA	
Plant	Leptospermum pallidum	-		NT	Wildlife Online	DSITI and ALA	
Plant	Lepturus minutus	-		V	Wildlife Online	DSITI and ALA	
Plant	Lindsaea pulchella var. blanda	-	V	EX	PMST	ALA	Confidential species; unable to download records
Plant	Marsdenia brevifolia	-	V	V	PMST	DSITI and ALA	
Plant	Oenanthe javanica	-	-	NT	Wildlife Online	DSITI and ALA	
Plant	Phaius australis	Lesser Swamp-orchid	E	F	PMST	ALA	Confidential species; unable to download records
Plant	Tephrosia leveillei		V	V	PMST	DSITI and ALA	connucritial species, unable to download records
Threatened faun		-	V	V	110151		
Bird	Calidris ferruginea	Curlew Sandpiper	CE (M)	SL	PMST	DSITI and ALA	
Bird	Calyptorhynchus lathami erebus	Glossy Black-cockatoo (northern)		V	Wildlife Online	DSITI and ALA	
Bird	Casuarius casuarius johnsonii	Southern Cassowary	F	V E	Wildlife Online and PMST	DSITI and ALA	
Mammal	Dasyurus hallucatus	Northern Quoli	E	L	PMST	DSITI and ALA	
Mammal	Dasyurus maculatus gracilis	Spotted-tailed Quoll (North Queensland)	E	-	PMST	DSITI and ALA	
Reptile	Egernia rugosa	Yakka Skink	V	L V	PMST	ALA	Confidential species; unable to download records
Bird	Erythrotriorchis radiatus	Red Goshawk	V	V E	PMST	DSITI and ALA	connidential species, unable to download records
Bird	Erythrura gouldiae	Gouldian Finch	E	С С	Wildlife Online and PMST	ALA	Confidential species; unable to download records
Mammal	Hipposideros semoni	Semon's Leaf-nosed Bat	E		PMST	DSITI and ALA	connuential species, unable to download records
Reptile	Lerista vittata	Mount Cooper Striped Lerista	V	E V	PMST	DSITI and ALA	
Frog	Litoria dayi	Australian Lace-lid	F	V F	PMST	DSITI and ALA	
U			E	E F	PMST	DSITI and ALA	
Frog	Litoria nannotis	Waterfall Frog	F	E	PMST	DSITI and ALA	
Frog	Litoria rheocola	Common Mistfrog Ghost Bat	E V	E	PMST		
Vammal	Macroderma gigas		V	V	Wildlife Online and PMST	DSITI and ALA DSITI and ALA	
Vammal	Mesembriomys gouldii rattoides	Black-footed Tree-rat (north Queensland)	-	- V			
Bird	Numenius madagascariensis	Eastern Curlew	CE (M)	V	PMST	DSITI and ALA	
Mammal	Petauroides volans	Greater Glider	V	-	Wildlife Online and PMST	DSITI and ALA	
Mammal	Petaurus gracilis	Mahogany Glider	E	E V	Wildlife Online	DSITI and ALA	
Mammal	Petrogale sharmani	Sharman's Rock-wallaby	V	V	Wildlife Online and PMST	DSITI and ALA	
Mammal	Phascolarctos cinereus	Koala	V	V	Wildlife Online and PMST	DSITI and ALA	
Bird	Poephila cincta cincta	Southern Black-throated Finch	E	E	Wildlife Online and PMST	ALA	Confidential species; unable to download records
Vammal	Pteropus conspicillatus	Spectacled Flying-fox	V	V	PMST	DSITI and ALA	
Vammal	Pteropus poliocephalus	Grey-headed Flying-fox	V	-	PMST	DSITI and ALA	
Bird	Rostratula australis	Australian Painted Snipe	E (M)	V	PMST	DSITI and ALA	
Mammal	Rhinolophus robertsi	Large-eared Horseshoe Bat	E	E	PMST	ALA	Confidential species; unable to download records
Mammal	Saccolaimus saccolaimus nudicluniatus	Bare-rumped Sheath-tailed Bat	CE	E	PMST		
Birds	Tyto novaehollandiae kimberli	Masked Owl (northern)	V	V	PMST		

Migratory and spe	cial least concern fauna species					
Bird (marine)	Apus pacificus	Fork-tailed Swift	М	SL	PMST	DSITI and ALA
Bird (marine)	Calidris ruficollis	Red-necked Stint	М	SL	Wildlife Online	DSITI and ALA
Bird (terrestrial)	Cuculus optatus	Oriental cuckoo	М	SL	Wildlife Online and PMST	DSITI and ALA
Bird (terrestrial)	Hirundapus caudacutus	White-throated Needletail	М	SL	Wildlife Online and PMST	DSITI and ALA
Bird (terrestrial)	Hirundo rustica	Barn Swallow	М	SL	PMST	DSITI and ALA
Bird (marine)	Hydroprogne caspia	Caspian Tern	М	SL	Wildlife Online	DSITI and ALA
Bird (terrestrial)	Monarcha melanopsis	Black-faced Monarch	М	SL	Wildlife Online and PMST	DSITI and ALA
Bird (terrestrial)	Motacilla cinerea	Grey Wagtail	М	SL	PMST	DSITI and ALA
Bird (terrestrial)	Motacilla flava	Yellow Wagtail	М	SL	PMST	DSITI and ALA
Bird (terrestrial)	Myiagra cyanoleuca	Satin Flycatcher	М	SL	PMST	DSITI and ALA
Bird (wetland)	Gallinago hardwickii	Latham's Snipe	М	SL	PMST	DSITI and ALA
Bird (wetland)	Pandion haliaetus	Osprey	М	SL	PMST	DSITI and ALA
Bird (terrestrial)	Rhipidura rufifrons	Rufous Fantail	М	SL	Wildlife Online and PMST	DSITI and ALA
Bird (terrestrial)	Symposiachrus trivirgatus	Spectacled Monarch	М	SL	Wildlife Online and PMST	DSITI and ALA
Mammal	Tachyglossus aculeatus	Short-beaked Echidna	-	SL	Wildlife Online	DSITI and ALA
Bird (wetland)	Tringa nebularia	Common Greenshank	М	SL	PMST	DSITI and ALA

# Appendix F

**THREATENED SPECIES RECORDS** 

## Kidston Corridor Options - NC Act threatened species records for each corridor option

NC Act status	Option A	Option B	Option C
Threatened flora species		-	
Endangered			
Vulnerable	1		
Near Threatened			
Threatened fauna species			
Endangered	1	1	1
Vulnerable		1	3
Near threatened			
Special least concern		2	1

## Kidston Corridor Options - EPBC Act threatened species records for each corridor option

EPBC Act status	Option A	Option B	Option C
Threatened flora species	•		
Critically endangered			
Endangered			
Vulnerable	1	1	1
Threatened fauna species			
Critically endangered		1	1
Endangered	1	1	1
Vulnerable	1	2	2
Migratory			
Migratory (marine)		1	
Migratory (terrestrial)			
Migratory (wetland)			

# Appendix G

## **MSES SPREADSHEETS**

## Kidston Corridor Option A - Regulated vegetation

Mama	M Table1 4			NAA 1	Aron ho	
	M_Table1_4	M4_RE	M4_VM_POLY	M4_1	Area_ha 0.003842	422 Total Catagory Deserve the face On the st
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R		422 Total Category R regrowth for Option A
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.469059	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.100844	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.195803	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.067611	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	12.59236	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.334779	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	7.860174	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	12.2317	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.374102	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.483168	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.700459	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.83997	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.070627	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.043435	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.830378	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.596549	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.905288	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	12.21573	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.167738	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	7.47583	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	14.34534	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.259695	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.054286	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.01047	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	5.514016	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.734517	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.020582	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.057303	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	16.09998	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.466693	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.528204	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	7.702433	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.215899	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.797685	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.034482	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.501958	
Genex: Option A						
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	5.960026	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.250622	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.163337	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.964128	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.210867	
		<null></null>	<null></null>	Cat R	0.768095	
Genex: Option A	REGULATED VEGETATION					
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.782941	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.121253	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.128736	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.148164	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.420224	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.068233	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.311802	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.472927	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.482328	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.577801	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.002918	
		<null></null>				
Genex: Option A	REGULATED VEGETATION		<null></null>	Cat R	0.380238	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.711728	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.303884	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.12129	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.335745	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.633654	
	REGULATED VEGETATION					
Genex: Option A		<null></null>	<null></null>	Cat R	2.034378	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.745112	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.606578	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.015766	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.385356	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.398018	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.210171	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.277036	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.247222	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.135324	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.614248	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.21904	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.032117	
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.078246	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.536691	
Genex: Option A				Cat R	12.71765	
enex: Option A	REGULATED VEGETATION	<null></null>	<null></null>			
enex: Option A enex: Option A	REGULATED VEGETATION				0.028328	
Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.028328	
enex: Option A enex: Option A enex: Option A enex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	4.450142	
enex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null></null></null></null>	<null> <null> <null></null></null></null>	Cat R Cat R Cat R	4.450142 0.80573	
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null></null></null></null></null>	<null> <null> <null> <null></null></null></null></null>	Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947	
Senex: Option A Senex: Option A Senex: Option A Senex: Option A Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null></null></null></null>	<null> <null> <null></null></null></null>	Cat R Cat R Cat R	4.450142 0.80573	
Senex: Option A Senex: Option A Senex: Option A Senex: Option A Senex: Option A Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null></null></null></null></null>	<null> <null> <null> <null></null></null></null></null>	Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> &lt;<null> <td><null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></td><td>Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R</td><td>4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974</td><td></td></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null></td><td>Cat R Cat R</td><td>4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942</td><td></td></n<></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null></null></null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></td><td>Cat R Cat R</td><td>4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486</td><td></td></n<></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> </null> </td></n<></null> </null> </null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> </null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.244263 0.373997	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> </null> </td></n<></null> </null> </null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> </null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.244263 0.373997 3.391567	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> </null> </td></n<></null> </null> </null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> </null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.244263 0.373997	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> </null> </td></n<></null> </null> </null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> </null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.244263 0.373997 3.391567	
Senex: Option A Senex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> &lt;<null> </null> </null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <n< td=""><td>Cat R Cat R</td><td>4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.424486 0.373997 3.391567 0.319338</td><td></td></n<></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	Cat R Cat R	4.450142 0.80573 0.271947 0.873751 0.017459 2.311132 0.953014 0.010974 7.544942 0.424486 0.424486 0.373997 3.391567 0.319338	

0		NL P		0.10	0.46.1
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.106444
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.009107
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Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.089904
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Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	7.493591
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.2608
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.959257
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.457158
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.645064
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.423002
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	7.708124
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.075444
Genex: Option A	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.20241
Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null></null>	Cat R	0.554833
Genex: Option A Genex: Option A	REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.052662
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Genex: Option A		<null> <null></null></null>	<null> <null></null></null>	Cat R	6.461723
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Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.473492
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.720557
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.084694
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.888013
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.034725
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.476705
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.890819
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	14.13818
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	14.58422
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	5.972214
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	3.9565
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.44608
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.483626
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.630734
Genex: Option A	REGULATED VEGETATION	7.8.10b	O-dom	rem_oc	0.000002
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	3.211762
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	6.756958
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.181751
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	0.097199
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.483261 9.568999
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.3.26a 7.5.4f	O-dom O-dom	rem_oc rem_oc	0.000082
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom O-dom	rem_oc	1.2572
Genex: Option A	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	0.009287
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	1.990015
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	8.687105
Genex: Option A	REGULATED VEGETATION	7.5.1c	O-dom	rem_oc	1.185375
Genex: Option A	REGULATED VEGETATION	7.5.1c	O-dom	rem_oc	0.297036
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	11.56206
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	13.34683
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	74.67075
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	4.908792
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.000035
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.589727
	DEOLU ATED MEOETATION	7.8.18a	O-dom	rem_oc	6.715692
Genex: Option A	REGULATED VEGETATION				
Genex: Option A Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	1.013457
Genex: Option A				rem_oc rem_oc rem_oc	1.013457 11.07438 694.5311

Conov. Ontion A	REGULATED VEGETATION	9.12.10	0-dom	rom 00	100.3972
Genex: Option A Genex: Option A	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	33.51078
Genex: Option A	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	118.4246
Genex: Option A	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	23.38401
Genex: Option A	REGULATED VEGETATION	9.5.2	O-dom	rem_oc	31.74374
Genex: Option A	REGULATED VEGETATION	9.5.2	O-dom	rem_oc	103.7197
Genex: Option A	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	0.016774
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	_	0.018774
	REGULATED VEGETATION	7.5.4c		rem_oc	
Genex: Option A			O-dom	rem_oc	0.115602
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.00036
Genex: Option A	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	0.72007
Genex: Option A	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	17.55466
Genex: Option A	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	0.623486
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.15235
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.000056
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.023917
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.024483
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.063902
Genex: Option A	REGULATED VEGETATION	7.5.4c	0-dom	rem_oc	0.161254
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.23368
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.402452
		7.5.4c	O-dom		7.455134
Genex: Option A	REGULATED VEGETATION			rem_oc	
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	11.3936
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	3.36858
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.147912
Genex: Option A	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	1.962138
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.280448
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	1.301988
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem oc	14.64406
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	6.153846
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	3.000527
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	_	6.518854
				rem_oc	
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	9.905173
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	2.436735
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	1.316996
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Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	19.62831
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	1.028011
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Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	2.647046
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Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom		9.52291
				rem_oc	
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	3.28628
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	32.21729
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	4.050669
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Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	1.561025
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Genex: Option A	REGULATED VEGETATION	7.5.1c	O-dom	rem_oc	5.784952
		7.5.1c		_	
Genex: Option A	REGULATED VEGETATION		O-dom	rem_oc	0.182818
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	1.321791
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	REGULATED VEGETATION	7.8.7a	O-dom		5.251581
			0-0011	rem_oc	
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c	O-dom O-dom	rem_oc	0.414249
Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a	O-dom O-dom O-dom	rem_oc rem_oc	0.414249
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c	O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c 7.5.4c 7.5.4c	O-dom O-dom O-dom	rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c	O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c 7.5.4c 7.5.4c	O-dom O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c 7.5.4c 7.5.4c 7.5.4c	O-dom O-dom O-dom O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.8.7a 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594065 0.81655 8.567495 62.13517 4.286205
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517 4.286205 6.221108
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.3.39a 7.5.2b	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517 4.286205 6.221108 34.09352
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.3.39a 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517 4.286205 6.221108 34.09352 6.059978
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.3.39a 7.5.2b 7.8.18a 7.8.18a	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517 4.286205 6.221108 34.09352 6.059978 15.98771
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.8.39a 7.5.2b 7.8.39a 7.8.18a 7.8.18a	O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594065 8.567495 62.13511 4.286205 6.221108 34.09352 6.059978 15.98771 9.11193
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.3.39a 7.5.2b 7.8.18a 7.8.18a	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 62.13517 4.286205 6.221108 34.09352 6.059978 15.98771 9.11193
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.8.39a 7.5.2b 7.8.39a 7.8.18a 7.8.18a	O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594063 0.81655 8.567499 62.13517 4.286205 6.221106 3.4.09355 6.059977 15.98771 9.11193 0.001799
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.3.39a 7.5.2b 7.8.18a 7.8.18a 7.8.18a 7.8.18a	0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom 0-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	0.414249 0.548143 4.418021 0.594067 0.81659 6.213517 4.286205 6.221108 34.09352 6.059978 15.98771 9.11193 0.001799 11.76356
Genex: Option A Genex: Option A	REGULATED VEGETATION REGULATED VEGETATION	7.5.4f 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.4c 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.5.2b 7.8.39a 7.5.2b 7.8.18a 7.8.18a 7.8.18a 7.8.18a 7.8.18a 7.8.18a	O-dom           O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	2.078499 0.414249 0.548143 4.418021 0.594067 0.81655 8.567495 6.213517 4.286205 6.221108 34.09352 6.059978 15.98771 9.11193 0.001799 11.76356 5.563263 18.22205

Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	0.616583	
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.436127	
Genex: Option A	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.278014	
Genex: Option A	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.488125	
Genex: Option A	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	11.64185	
Genex: Option A	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.444011	
Genex: Option A	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	6.182993	
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	9.199162	
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	1.360322	
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	8.433969	
Genex: Option A	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.434568	
Genex: Option A	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	0.64519	
Genex: Option A	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	58.61665	
Genex: Option A	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	15.50554	
Genex: Option A	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	0.338929	
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	18.62776	
Genex: Option A	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	5.381855	
Genex: Option A	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	348.0026	
Genex: Option A	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	345.7832	
Genex: Option A	REGULATED VEGETATION	9.3.19a/9.3.3a/9.3.23	O-subdom	rem_oc	0.053888	
Genex: Option A	REGULATED VEGETATION	9.3.19a/9.3.3a/9.3.23	O-subdom	rem_oc	0.003095	
Genex: Option A	REGULATED VEGETATION	9.3.19a/9.3.3a/9.3.23	O-subdom	rem_oc	0.418459	
Genex: Option A	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	20.80439	
Genex: Option A	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	48.92501	
Genex: Option A	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	130.6595	
Genex: Option A	REGULATED VEGETATION	9.3.19a/9.3.3a/9.3.23	O-subdom	rem_oc	18.58643	
Genex: Option A	REGULATED VEGETATION	9.3.19a/9.3.3a/9.3.23	O-subdom	rem_oc	94.72298	

1008 Total OC-subdom remnant for Option A

## Kidston Corridor Option B - Regulated vegetation

Name	M_Table1_4	M4_RE	M4 VM POLY	M4_1	Area_ha	
						20/ Tatal Catagory Disagon th for Option D
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.004087	296 Total Category R regrowth for Option B
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.146045	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.391259	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.464253	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.080121	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.305659	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.313248	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.341295	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.323414	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.326592	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.686231	
		<null></null>	<null></null>			
Genex: Option B	REGULATED VEGETATION			Cat R	0.67304	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.257651	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.612285	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.669419	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.351457	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.106444	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.009107	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.699534	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	6.691298	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.45794	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.898714	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	13.99839	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.504503	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	21.10548	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>		21.10548	
				Cat R		
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.36551	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.019771	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.089904	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.190246	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.156919	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.046139	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.2608	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.307026	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.380483	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.872902	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.434148	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.434542	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.46824	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.709577	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.014552	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.475646	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.420251	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.185744	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.450237	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.410175	
			<null></null>		0.410175	
Genex: Option B	REGULATED VEGETATION	<null></null>		Cat R		
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.327941	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.659546	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.480604	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.861445	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.632978	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.422231	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.0623	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.827108	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.380445	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.597613	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.061373	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.011887	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.108115	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.092819	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.36721	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.400628	
Genex: Option B		<null></null>	<null></null>	Cat R	0.15454	
	REGULATED VEGETATION		<null></null>	Cat R	1.604932	
Genex: Option B	REGULATED VEGETATION	<null></null>			0.798694	
Genex: Option B Genex: Option B	REGULATED VEGETATION	<null></null>				
Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION	<null></null>	<null></null>	Cat R		
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R	1.938117	
Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION	<null></null>	<null></null>			
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	1.938117	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null></null></null></null></null>	<null> <null> <null> <null></null></null></null></null>	Cat R Cat R Cat R	1.938117 1.112028 0.45894	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<nui> <nui> <nui> <nui> <nui> <nui></nui></nui></nui></nui></nui></nui>	<null> <null> <null> <null> <null></null></null></null></null></null>	Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<pre> <null>     </null></pre> <pre> </pre>	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266 0.70253	
Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null>	<pre><null> </null></pre> <null> <null> <null> <null> <null> <null> <null> </null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266 0.70253 0.773593	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<pre> <null>     </null></pre> <pre> </pre>	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266 0.70253	
Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null>	<pre><null> </null></pre> <null> <null> <null> <null> <null> <null> <null> </null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266 0.70253 0.773593	
Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <n< td=""><td><null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null></td><td>Cat R Cat R Cat R Cat R Cat R Cat R Cat R</td><td>1.938117 1.112028 0.45894 0.659266 0.70253 0.773593 0.785154</td><td></td></n<></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R Cat R Cat R	1.938117 1.112028 0.45894 0.659266 0.70253 0.773593 0.785154	

Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.468714	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.029329	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.877687	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.262705	
Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.590311	
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Genex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.388676	
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enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.234613	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.532087	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.332087	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R Cat R	0.281566	
Genex: Option B					4.103995	
Senex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.103995 0.802553	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R		
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	6.461723	
Senex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.63187	
Senex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	42.18013	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.223331	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.832999	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.321527	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.424048	
enex: Option B	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.072922	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.084694	2.4 Total OC regrowth for Option B
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.888013	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.034725	
Genex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.476705	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.890819	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.099917	1891 Total OC-dom remnant for Option B
enex: Option B	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	1.166763	
enex: Option B	REGULATED VEGETATION	7.5.1c	O-dom	rem_oc	0.037098	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	15.35428	
Genex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	5.626586	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	1.47184	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	10.58911	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	138.478	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	195.4305	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	153.7515	
enex: Option B	REGULATED VEGETATION	9.5.2	O-dom	rem_oc	5.651613	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	19.62831	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	1.028011	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	1.132719	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	2.647046	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	0.811436	
enex: Option B	REGULATED VEGETATION	7.5.2b	0-dom	rem_oc	9.52291	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	3.28628	
enex: Option B	REGULATED VEGETATION	7.5.2b	0-dom	rem_oc	32.21729	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	4.050669	
enex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	2.310894	
enex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	1.561025	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	2.128498	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.81339	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.928591	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	1.05823	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.413503	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.925361	
enex: Option B	REGULATED VEGETATION	7.5.1c	O-dom	rem_oc	5.784952	
enex: Option B	REGULATED VEGETATION	7.5.1c	0-dom	rem_oc	0.182818	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	1.321791	
enex: Option B	REGULATED VEGETATION	7.5.4c	0-dom		1.076465	
		7.5.4c		rem_oc	0.144677	
enex: Option B	REGULATED VEGETATION		O-dom	rem_oc		
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	14.87732	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	2.728083	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	27.99035	
enex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	47.06895	
enex: Option B	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	4.770745	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	1.478203	
	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	28.1685	
	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	4.236403	
enex: Option B		7.5.2b	O-dom	rem_oc	1.381377	
enex: Option B enex: Option B	REGULATED VEGETATION			rem_oc	0.053238	
enex: Option B enex: Option B		7.3.39a	O-dom		0.000200	
enex: Option B enex: Option B enex: Option B	REGULATED VEGETATION		O-dom O-dom	rem_oc	0.50231	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION	7.3.39a				
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.3.39a 7.3.39a	O-dom	rem_oc	0.50231	
enex: Option B enex: Option B enex: Option B enex: Option B enex: Option B enex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.3.39a 7.3.39a 7.5.4c 9.12.10	O-dom O-dom O-dom	rem_oc rem_oc rem_oc	0.50231 9.653945 270.0272	
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.3.39a 7.3.39a 7.5.4c	O-dom O-dom	rem_oc rem_oc	0.50231 9.653945	

Genex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem oc	108.0731	
Genex: Option B	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	1.946469	
Genex: Option B	REGULATED VEGETATION	9.5.2	O-dom	rem_oc	26.87198	
Genex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	4.418021	
Genex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.594067	
Genex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.81655	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	8.567495	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	62.13517	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	4.286205	
Genex: Option B	REGULATED VEGETATION	7.3.39a	O-dom		6.221108	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	34.09352	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	6.059978	
Genex: Option B	REGULATED VEGETATION	7.8.18a	O-dom		15.98771	
				rem_oc	9,11193	
Senex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.001799	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc		
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	11.76356	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	5.563263	
enex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	18.22205	
enex: Option B	REGULATED VEGETATION	7.8.17c	O-dom	rem_oc	3.072896	
enex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	0.616583	
nex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.436127	
nex: Option B	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.278014	
nex: Option B	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.488125	
enex: Option B	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	11.64185	
enex: Option B	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.444011	
enex: Option B	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	6.182993	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	9.199162	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	1.360322	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	8.433969	
enex: Option B	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.434568	
enex: Option B	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	0.64519	
enex: Option B	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	58.61665	
enex: Option B	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	15.50554	
enex: Option B	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	0.338929	
enex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	18.62776	
Genex: Option B	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	5.381855	
Genex: Option B	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	187.4794	442 Total OC-subdom remnant for Option
enex: Option B	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	46.67159	
enex: Option B	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	6.946683	
enex: Option B	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	4.034855	
enex: Option B	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	28.05193	
Genex: Option B	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem oc	59.27586	
Genex: Option B	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem oc	85.42147	
				_		
Genex: Option B	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	21.51575	

## Kidston Corridor Option C - Regulated vegetation

N	44 T-61-4 4			N 4 4 4	A	
Name	M_Table1_4	M4_RE	M4_VM_POLY	M4_1	Area_ha	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.328481	257 Total Category R regrowth for Option C
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.42827	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.926819	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	12.75876	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.043894	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.638457	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.716188	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.003291	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.164262	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.767526	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.17231	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.711402	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.027479	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.042959	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.229094	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.651188	
		<null></null>	<null></null>		0.627135	
GENEX Option C	REGULATED VEGETATION			Cat R		
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.280835	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	27.73202	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.293773	
		<null></null>				
GENEX Option C	REGULATED VEGETATION		<null></null>	Cat R	0.65791	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.774417	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.458449	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.535426	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	5.619303	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.315004	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.721397	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.529604	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.378213	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.419634	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.829549	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.811617	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.522101	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.007855	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.206561	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.408106	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.040172	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.508238	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.434682	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.3229	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.777548	
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GENEX Option C	REGULATED VEGETATION					
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	2.508335	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.052041	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.423372	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.423002	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.013744	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.284381	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.331178	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	4.380483	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.434148	
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.46824	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.709577	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.014552	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.475646	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.420251	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.185744	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.450237	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.410175	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.408328	
	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.327941	
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.659546	
GENEX Option C	ACCOUNTED VEOLIATION	<null></null>				
GENEX Option C GENEX Option C	DECULATED VECETATION	12 MUUN	<null></null>	Cat R	0.480604	
GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION			Cat R	0.861445	
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null></null>	<null></null>			
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION			Cat R	0.632978	
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null></null>		0.632978	
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null></null></null></null>	<null> <null></null></null>	Cat R	0.422231	
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null></null></null></null></null>	<null> <null> <null></null></null></null>	Cat R Cat R	0.422231 0.0623	
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null></null></null></null>	<null> <null></null></null>	Cat R	0.422231	
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null></null></null></null></null></null>	<null> <null> <null> <null></null></null></null></null>	Cat R Cat R Cat R	0.422231 0.0623 0.827108	
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	<null> <null> <null> <null> <null></null></null></null></null></null>	Cat R Cat R Cat R Cat R	0.422231 0.0623 0.827108 1.380445	
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null> <null></null></null></null></null></null></null></null>	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	Cat R Cat R Cat R Cat R Cat R	0.422231 0.0623 0.827108 1.380445 0.597613	
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	<null> <null> <null> <null> <null> <null></null></null></null></null></null></null>	<null> <null> <null> <null> <null></null></null></null></null></null>	Cat R Cat R Cat R Cat R	0.422231 0.0623 0.827108 1.380445	

CENEX Option C		-Noll-	«Molls	Cot D	0.100115
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.108115
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.36721
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.400628
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.15454
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	1.604932 0.798694
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R Cat R	1.938117
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.112028
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GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.659266
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.70253
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.773593 0.785154
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.242768
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.615052
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.400008
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.468714
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.029329
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.714711
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.645599
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R Cat R	0.877687 3.262705
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.590311
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.464782
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.733858
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.113461
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.388676
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.672468
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.550556
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.543306
GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null></null>	Cat R	0.106496
GENEX Option C GENEX Option C	REGULATED VEGETATION	<nuii> <nuii></nuii></nuii>	<null> <null></null></null>	Cat R Cat R	0.075444
GENEX Option C	REGULATED VEGETATION	<nuii> <nuii></nuii></nuii>	<null> <null></null></null>	Cat R Cat R	2.20241
GENEX Option C	REGULATED VEGETATION	<nuii> <nuii></nuii></nuii>	<null></null>	Cat R Cat R	0.554833
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.052662
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.595646
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.610715
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	17.18279
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.561512
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.250953
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.162867
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	9.736478
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.068693
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.139407
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.361331
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.234613
GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<null> <null></null></null>	<null> <null></null></null>	Cat R	0.532087
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	<nuii> <nuii></nuii></nuii>	<null> <null></null></null>	Cat R Cat R	0.281566
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R Cat R	4.103995
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R Cat R	4.103995
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	6.461723
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.63187
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	42.18013
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.223331
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	3.832999
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	1.321527
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.424048
GENEX Option C	REGULATED VEGETATION	<null></null>	<null></null>	Cat R	0.072922
GENEX Option C	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.084694
GENEX Option C	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.888013
GENEX Option C	REGULATED VEGETATION	7.5.2b	O-dom	hvr_oc	0.034725
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.476705
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	hvr_oc	0.890819
GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	7.8.18a 7.5.4f	O-dom	rem_oc	3.344087 5.250493
GENEX Option C GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom O-dom	rem_oc rem_oc	2.477648
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom O-dom	rem_oc	4.760297
GENEX Option C	REGULATED VEGETATION	7.5.41 7.8.18c	O-dom	rem_oc	5.401995
GENEX Option C	REGULATED VEGETATION	7.3.39a	O-dom	rem_oc	0.746091
GENEX Option C	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.817491
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	36.82176
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	204.122
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	8.058572
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	1.779746
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	11.61961
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.046029
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	5.615905
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	20.59074
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	7.634229
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.792089
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	5 719127
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	5.718137
	REGULATED VEGETATION REGULATED VEGETATION	7.8.18c 7.3.39a	O-dom O-dom	rem_oc rem_oc	21.60491 1.304391
GENEX Option C		7.3.39a 7.8.18c	O-dom O-dom	rem_oc	4.964711
GENEX Option C				rem_oc	7.624219
GENEX Option C GENEX Option C	REGULATED VEGETATION		O-dom	UL UL	
GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	7.5.4c	O-dom O-dom	rem oc	5 251591
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a	O-dom	rem_oc rem_oc	5.251581 2.078499
GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION	7.5.4c		rem_oc rem_oc rem_oc	5.251581 2.078499 0.414249
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a 7.5.4f	O-dom O-dom	rem_oc	2.078499
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a 7.5.4f 7.5.4c	O-dom O-dom O-dom	rem_oc rem_oc	2.078499 0.414249
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a 7.5.4f 7.5.4c 7.8.7a	O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc	2.078499 0.414249 0.548143 0.053238 0.50231
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a 7.5.4f 7.5.4c 7.8.7a 7.3.39a 7.3.39a 7.5.4c	O-dom O-dom O-dom O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc rem_oc	2.078499 0.414249 0.548143 0.053238 0.50231 9.653945
GENEX Option C GENEX Option C	REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION REGULATED VEGETATION	7.5.4c 7.8.7a 7.5.4f 7.5.4c 7.8.7a 7.3.39a 7.3.39a	O-dom O-dom O-dom O-dom O-dom O-dom	rem_oc rem_oc rem_oc rem_oc rem_oc	2.078499 0.414249 0.548143 0.053238 0.50231

GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem oc	0.590697
GENEX Option C	REGULATED VEGETATION	9.12.10	0-dom	rem_oc	1.771366
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	108.0731
GENEX Option C	REGULATED VEGETATION	9.12.10	O-dom	rem_oc	1.946469
GENEX Option C	REGULATED VEGETATION	9.5.2	0-dom	rem_oc	26.87198
GENEX Option C	REGULATED VEGETATION	7.5.4c	0-dom	rem_oc	4.418021
GENEX Option C	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	0.594067
GENEX Option C	REGULATED VEGETATION	7.5.4c	0-dom	rem_oc	0.81655
GENEX Option C	REGULATED VEGETATION	7.5.2b	0-dom	rem_oc	8.567495
GENEX Option C	REGULATED VEGETATION	7.5.2b	0-dom	rem_oc	62.13517
GENEX Option C	REGULATED VEGETATION	7.5.2b	0-dom	_	4.286205
GENEX Option C	REGULATED VEGETATION	7.3.39a	0-dom	rem_oc	6.221108
		7.5.2b		rem_oc	34.09352
GENEX Option C	REGULATED VEGETATION		O-dom	rem_oc	
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	6.059978
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	15.98771
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	9.11193
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	0.001799
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	11.76356
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	5.563263
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	18.22205
GENEX Option C	REGULATED VEGETATION	7.8.17c	O-dom	rem_oc	3.072896
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	0.616583
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.436127
GENEX Option C	REGULATED VEGETATION	7.5.4f	O-dom	rem_oc	3.278014
GENEX Option C	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.488125
GENEX Option C	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	11.64185
GENEX Option C	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	4.444011
GENEX Option C	REGULATED VEGETATION	7.3.39a	O-dom	rem oc	6.182993
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem oc	9.199162
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem oc	1.360322
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem_oc	8.433969
GENEX Option C	REGULATED VEGETATION	7.8.18a	O-dom	rem oc	0.434568
GENEX Option C	REGULATED VEGETATION	7.8.18c	O-dom	rem_oc	0.64519
GENEX Option C	REGULATED VEGETATION	7.5.4c	O-dom	rem_oc	58.61665
GENEX Option C	REGULATED VEGETATION	7.8.7a	O-dom	rem_oc	15.50554
GENEX Option C	REGULATED VEGETATION	7.8.7a	O-dom	rem oc	0.338929
GENEX Option C	REGULATED VEGETATION	7.5.2b	O-dom	rem_oc	18.62776
GENEX Option C	REGULATED VEGETATION	7.5.2b	0-dom	rem_oc	5.381855
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom		58.16345
				rem_oc	
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	7.270749
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	0.873508
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	24.94581
GENEX Option C	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	11.81889
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	63.09471
GENEX Option C	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	59.27586
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	85.42147
GENEX Option C	REGULATED VEGETATION	9.3.5/9.3.3a/9.3.23	O-subdom	rem_oc	21.51575
GENEX Option C	REGULATED VEGETATION	9.3.3a/9.3.20/9.3.23	O-subdom	rem_oc	2.117612

### Kidston Corridor Option A - Regional ecosystems v8

Name	VM_POLY	VM_STATUS	Area_ha	
Genex: Option A	LeastC	hvr_leastc	14.8125	62 Total LC regrowth for Option A
Genex: Option A	LeastC	hvr_leastc	0.367214	
Genex: Option A	LeastC	hvr_leastc	1.073194	
Genex: Option A	LeastC	hvr_leastc	0.02707	
Genex: Option A	LeastC	hvr_leastc	0.725399	
Genex: Option A	LeastC	hvr_leastc	23.53707	
Genex: Option A	LeastC	hvr_leastc	6.913921	
Genex: Option A	LeastC	hvr_leastc	4.153932	
Genex: Option A	LeastC	hvr_leastc	2.201232	
Genex: Option A	LeastC	hvr_leastc	2.657052	
Genex: Option A	LeastC	hvr_leastc	1.646678	
Genex: Option A	LeastC	hvr_leastc	3.808301	
Genex: Option A	O-dom	hvr_oc	0.47349	3.6 Total OC regrowth for Option A
Genex: Option A	O-dom	hvr_oc	0.72056	
Genex: Option A	O-dom	hvr_oc	0.888013	
Genex: Option A	O-dom	hvr_oc	0.034724	
Genex: Option A	O-dom	hvr_oc	0.084695	
Genex: Option A	O-dom	hvr_oc	0.476706	
Genex: Option A	O-dom	hvr_oc	0.890819	
Genex: Option A	non-rem	non_remnant	121.2521	2675 Total non-remnant for Option A
Genex: Option A	non-rem	non_remnant	25.84752	
Genex: Option A	non-rem	non_remnant	2.629019	
Genex: Option A	non-rem	non_remnant	19.38068	
Genex: Option A	non-rem	non_remnant	0.000169	
Genex: Option A	non-rem	non_remnant	2.304268	
Genex: Option A	non-rem	non_remnant	4.975241	
Genex: Option A	non-rem	non_remnant	112.914	
Genex: Option A	non-rem	non_remnant	0.444669	
Genex: Option A	non-rem	non_remnant	31.96318	
Genex: Option A	non-rem	non_remnant	63.51757	
Genex: Option A	non-rem	non_remnant	1.279065	
Genex: Option A	non-rem	non_remnant	190.1669	
Genex: Option A	non-rem	non_remnant	13.65374	
Genex: Option A	non-rem	non_remnant	6.749425	
Genex: Option A	non-rem	non_remnant	91.04484	
Genex: Option A	non-rem	non_remnant	0.000071	
Genex: Option A	non-rem	non_remnant	122.9623	
Genex: Option A	non-rem	non_remnant	0.000227	
Genex: Option A	non-rem	non_remnant	279.2766	
Genex: Option A	non-rem	non_remnant	151.6287	
Genex: Option A	non-rem	non_remnant	7.245895	
Genex: Option A	non-rem	non_remnant	8.656713	
Genex: Option A	non-rem	non_remnant	393.879	
Genex: Option A	non-rem	non_remnant	2.125011	
Genex: Option A	non-rem	non_remnant	103.4499	
Genex: Option A	non-rem	non_remnant	8.966574	
Genex: Option A	non-rem	non_remnant	6.867542	
Genex: Option A	non-rem	non_remnant	16.25738	
Genex: Option A	non-rem	non_remnant	0.234067	
Genex: Option A	non-rem	non_remnant	282.7614	
Genex: Option A	non-rem	non_remnant	0.000204	
Genex: Option A	non-rem	non_remnant	0.000074	
Genex: Option A	non-rem	non_remnant	8.914747	
Genex: Option A	non-rem	non_remnant	1.286566	
Genex: Option A	non-rem	non_remnant	1.685735	
Genex: Option A	non-rem	non_remnant	0.526879	
Genex: Option A	non-rem	non_remnant	0.121333	
Genex: Option A	non-rem	non_remnant	3.735213	
Genex: Option A	non-rem	non_remnant	117.347	
Genex: Option A	non-rem	non_remnant	468.6799	

Genex: Option A	water	non_remnant	17.22085	20 Total water non-remnant for Option A
Genex: Option A	water	non_remnant	1.779753	
Genex: Option A	water	non_remnant	1.100773	
Genex: Option A	LeastC	rem_leastc	52.57021	50675 Total LC remnant for Option A
Genex: Option A	LeastC	rem_leastc	57.17364	
Genex: Option A	LeastC	rem_leastc	146.5935	
Genex: Option A	LeastC	rem_leastc	1.712134	
Genex: Option A	LeastC	rem_leastc	32.87388	
Genex: Option A	LeastC	rem_leastc	255.2324	
Genex: Option A	LeastC	rem_leastc	1032.305	
Genex: Option A	LeastC	rem_leastc	34.28771	
Genex: Option A	LeastC	rem_leastc	276.3749	
Genex: Option A	LeastC	rem_leastc	18.62152	
Genex: Option A	LeastC	rem_leastc	42.91812	
Genex: Option A	LeastC	rem_leastc	9.551715	
Genex: Option A	LeastC	rem_leastc	54.19333	
Genex: Option A	LeastC	rem_leastc	37.36713	
Genex: Option A	LeastC	rem_leastc	14.77693	
Genex: Option A	LeastC	rem_leastc	52.74941	
Genex: Option A	LeastC	rem_leastc	36.25535	
Genex: Option A	LeastC	rem_leastc	127.1683	
Genex: Option A	LeastC	rem_leastc	27.34685	
Genex: Option A	LeastC	rem_leastc	157.0211	
Genex: Option A	LeastC	rem_leastc	9.903885	
Genex: Option A	LeastC	rem_leastc	26.95378	
Genex: Option A	LeastC	rem_leastc	29.87643	
Genex: Option A	LeastC	rem_leastc	59.37642	
Genex: Option A	LeastC	rem_leastc	347.307 125.124	
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	8.416606	
Genex: Option A	LeastC	rem leastc	136.8527	
Genex: Option A	LeastC	rem_leastc	23.68275	
Genex: Option A	LeastC	rem_leastc	1152.16	
Genex: Option A	LeastC	rem_leastc	8.13693	
Genex: Option A	LeastC	rem leastc	21.6886	
Genex: Option A	LeastC	rem leastc	73.47771	
Genex: Option A	LeastC	rem leastc	22.65108	
Genex: Option A	LeastC	rem leastc	68.04189	
Genex: Option A	LeastC	rem leastc	42.60602	
Genex: Option A	LeastC	rem_leastc	24.16052	
Genex: Option A	LeastC	rem leastc	22.17645	
Genex: Option A	LeastC	rem_leastc	62.46455	
Genex: Option A	LeastC	rem_leastc	595.0399	
Genex: Option A	LeastC	rem_leastc	1.471328	
Genex: Option A	LeastC	rem_leastc	61.99931	
Genex: Option A	LeastC	rem_leastc	239.4195	
Genex: Option A	LeastC	rem_leastc	68.66814	
Genex: Option A	LeastC	rem_leastc	147.3336	
Genex: Option A	LeastC	rem_leastc	1.401031	
Genex: Option A	LeastC	rem_leastc	251.6928	
Genex: Option A	LeastC	rem_leastc	22.36634	
Genex: Option A	LeastC	rem_leastc	172.1274	
Genex: Option A	LeastC	rem_leastc	67.82442	
Genex: Option A	LeastC	rem_leastc	158.9092	
Genex: Option A	LeastC	rem_leastc	4.672963	
Genex: Option A	LeastC	rem_leastc	105.2463	
Genex: Option A	LeastC	rem_leastc	45.21432	
Genex: Option A	LeastC	rem_leastc	4.683257	
Genex: Option A	LeastC	rem_leastc	43.53392	
Genex: Option A	LeastC	rem_leastc	1.189481	
Genex: Option A	LeastC	rem_leastc	129.4995	
Genex: Option A	LeastC	rem_leastc	185.6366	
Genex: Option A	LeastC	rem_leastc	207.8615	

Genex: Option A	LeastC	rem_leastc	4.923067
Genex: Option A	LeastC	rem_leastc	19.29285
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Genex: Option A	LeastC	rem_leastc	406.9075
Genex: Option A	LeastC	rem leastc	396.7804
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Genex: Option A	LeastC	rem_leastc	7.393406
Genex: Option A	LeastC	rem_leastc	42.71687
Genex: Option A		rem leastc	56.29743
	LeastC	-	
Genex: Option A	LeastC	rem_leastc	77.57893
Genex: Option A	LeastC	rem_leastc	294.1476
Genex: Option A	LeastC	rem_leastc	161.5184
Genex: Option A	LeastC	rem_leastc	27.12249
Genex: Option A	LeastC	rem_leastc	1485.42
Genex: Option A	LeastC	rem_leastc	5.900959
Genex: Option A	LeastC	rem leastc	1230.668
Genex: Option A	LeastC	rem_leastc	91.84999
Genex: Option A	LeastC	rem_leastc	299.5232
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Genex: Option A	LeastC	rem_leastc	325.435
Genex: Option A	LeastC	rem_leastc	5.030852
Genex: Option A	LeastC	rem leastc	36.13185
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Genex: Option A	LeastC	rem_leastc	32.00785
Genex: Option A	LeastC	rem leastc	29.35829
Genex: Option A	LeastC	rem_leastc	179.814
Genex: Option A	LeastC	rem leastc	0.068167
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Genex: Option A	LeastC	rem_leastc	36.75109
Genex: Option A	LeastC	rem_leastc	274.0591
Genex: Option A	LeastC	rem leastc	0.614416
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Genex: Option A	LeastC	rem_leastc	11.44757
Genex: Option A	LeastC	rem_leastc	337.7416
Genex: Option A		rem leastc	
	LeastC		0.215319
Genex: Option A	LeastC	rem_leastc	78.6461
Genex: Option A	LeastC	rem leastc	92.57578
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Genex: Option A	LeastC	rem_leastc	1181.485
Genex: Option A	LeastC	rem leastc	128.7503
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Genex: Option A	LeastC	rem_leastc	395.8646
Genex: Option A	LeastC	rem leastc	14.96748
Genex: Option A	LeastC	rem_leastc	263.8201
Genex: Option A	LeastC	rem_leastc	110.4523
Genex: Option A	LeastC	rem leastc	18.81884
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Genex: Option A	LeastC	rem_leastc	26.17039
Genex: Option A	LeastC	rem_leastc	11.36333
Genex: Option A	LeastC	rem_leastc	5.997836
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Genex: Option A	LeastC	rem_leastc	44.11509
Genex: Option A	LeastC	rem_leastc	48.96257
Genex: Option A	LeastC	rem leastc	8.616055
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Genex: Option A	LeastC	rem_leastc	34.58335
Genex: Option A	LeastC	rem_leastc	6.739789
Genex: Option A	LeastC	rem leastc	159.7099
Genex: Option A	LeastC	rem_leastc	0.689372
Genex: Option A	LeastC	rem_leastc	5.06318
		1	
Genex: Option A	LeastC	rem_leastc	723.1585
Genex: Option A	LeastC	rem_leastc	90.37985
Genex: Option A	LeastC	rem_leastc	525.1697
Genex: Option A	LeastC	rem_leastc	61.11363
Genex: Option A	LeastC	rem_leastc	0.189301
Genex: Option A	LeastC	rem_leastc	131.1014
Genex: Option A	LeastC	rem_leastc	1103.557
Genex: Option A	LeastC		
	LCasil	rem leaste	
Genex: Option A		rem_leastc	105.9817
	LeastC	rem_leastc rem_leastc	11.87414
	LeastC	rem_leastc	11.87414
Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115
Genex: Option A Genex: Option A	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683
Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115
Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664
Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419
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Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22087 44.22087 191.8419 261.3817 29.57973 1586.245 132.7455
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661 94.6175
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661 94.6175
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661 94.6175 132.2248
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661 94.6175 132.2248
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	11.87414 59.61115 21.74683 590.3104 168.4664 77.45731 149.9271 98.58277 668.4066 44.27461 1.12957 44.22081 191.8419 261.3817 29.57973 1586.245 132.7455 60.33661 94.6175

Genex: Option A Genex: Option A			
	LeastC	rem_leastc	20.66312
Genex: Option A		_	23.24325
	LeastC	rem_leastc	
Genex: Option A	LeastC	rem_leastc	15.31929
Genex: Option A	LeastC	rem leastc	174.3232
Genex: Option A	LeastC	rem_leastc	212.8426
Genex: Option A	LeastC	rem_leastc	313.0483
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Genex: Option A	LeastC	rem_leastc	1.196521
Genex: Option A	LeastC	rem_leastc	108.2252
Genex: Option A	LeastC	rem_leastc	36.09727
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Genex: Option A	LeastC	rem_leastc	30.38112
Genex: Option A	LeastC	rem_leastc	8.561523
Genex: Option A	LeastC	rem_leastc	88.58386
Genex: Option A	LeastC	rem_leastc	6.175024
Genex: Option A	LeastC	rem leastc	286.5008
Genex: Option A	LeastC	rem_leastc	42.79844
Genex: Option A	LeastC	rem_leastc	11.98431
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Genex: Option A	LeastC	rem_leastc	264.8763
Genex: Option A	LeastC	rem_leastc	226.6269
Genex: Option A	LeastC	rem leastc	58.80618
Genex: Option A	LeastC	rem_leastc	5.885673
Genex: Option A	LeastC	rem leastc	1064.296
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Genex: Option A	LeastC	rem_leastc	92.68297
Genex: Option A	LeastC	rem_leastc	20.83021
Genex: Option A	LeastC	rem_leastc	37.9357
Genex: Option A	LeastC	rem_leastc	5.468407
Genex: Option A	LeastC	rem leastc	186.0467
Genex: Option A	LeastC	rem leastc	49.86623
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Genex: Option A	LeastC	rem_leastc	297.9879
Genex: Option A	LeastC	rem leastc	153.8439
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Genex: Option A	LeastC	rem_leastc	180.351
Genex: Option A	LeastC	rem leastc	660.1058
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Genex: Option A	LeastC	rem_leastc	100.4104
Genex: Option A	LeastC	rem_leastc	62.69539
Genex: Option A	LeastC	rem_leastc	26.27379
Genex: Option A	LeastC	rem_leastc	161.7457
Genex: Option A	LeastC	rem_leastc	1.141023
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Genex: Option A	LeastC	rem_leastc	67.52318
Genex: Option A	LeastC	rem leastc	182.8862
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Genex: Option A	LeastC	rem_leastc	34.12553
Genex: Option A	LeastC	rem_leastc	25.81363
Genex: Option A	LeastC	rem leastc	1075.062
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Genex: Option A	LeastC	rem_leastc	155.934
Genex: Option A	LeastC	rem leastc	12.85443
Genex: Option A	LeastC	rem_leastc	3539.709
Genex: Option A	LeastC	rem leastc	695.5142
Genex: Option A	LeastC	rem leastc	28.58443
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Genex: Option A	LeastC	rem_leastc	69.74197
Genex: Option A	LeastC	rem_leastc	0.853428
		Tem_lease	
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	LeastC	rem_leastc	0.18627
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Genex: Option A	LeastC	rem_leastc	0.18627 0.001726
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Genex: Option A	LeastC	rem_leastc	0.18627 0.001726
Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924
Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885
Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424
Genex: Option A Genex: Option A	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186 167.1931
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186 167.1931 18.74343
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186 167.1931
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186 167.1931 18.74343 80.59614
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	0.18627 0.001726 309.9822 11.33468 32.23451 59.20924 25.91885 39.92079 26.49348 151.5962 154.6605 226.7467 185.4138 6.451919 3.380721 473.4424 2.155563 0.978982 154.2547 912.1949 1.431118 46.05539 13.67831 0.00085 42.24882 0.250186 167.1931 18.74343

Genex: Option A	LeastC	rem_leastc	68.2613
Genex: Option A	LeastC	rem_leastc	77.81073
Genex: Option A	LeastC	rem_leastc	0.388276
Genex: Option A	LeastC	rem_leastc	109.4388
Genex: Option A	LeastC	rem_leastc	1.866744
Genex: Option A	LeastC	rem_leastc	1411.093
Genex: Option A	LeastC	rem_leastc	12.58193
Genex: Option A	LeastC	rem_leastc	134.5518
Genex: Option A	LeastC	rem_leastc	972.373
Genex: Option A	LeastC	rem_leastc	24.79335
Genex: Option A	LeastC	rem_leastc	52.52491
Genex: Option A	LeastC	rem leastc	421.2866
Genex: Option A	LeastC	rem_leastc	97.98
Genex: Option A	LeastC	rem leastc	455.7125
Genex: Option A	LeastC	rem_leastc	1.622995
Genex: Option A	LeastC	rem_leastc	38.69352
Genex: Option A	LeastC	rem_leastc	31.32427
Genex: Option A	LeastC	rem_leastc	7.188786
Genex: Option A	LeastC	rem_leastc	235.663
Genex: Option A	LeastC	rem leastc	301.155
Genex: Option A	LeastC	rem_leastc	27.47533
Genex: Option A	LeastC	rem_leastc	539.3737
Genex: Option A	LeastC	rem_leastc	6.576623
Genex: Option A	LeastC	rem_leastc	89.62867
Genex: Option A	LeastC	rem_leastc	4.997119
Genex: Option A	LeastC	rem leastc	22.16766
Genex: Option A	LeastC	rem leastc	474.5252
Genex: Option A	LeastC	rem leastc	183.6038
Genex: Option A	LeastC	rem leastc	256.3734
Genex: Option A	LeastC	rem leastc	1596.657
Genex: Option A	LeastC		
		rem_leastc	10.11799
Genex: Option A	LeastC LeastC	rem_leastc	376.1155
Genex: Option A Genex: Option A	LeastC	rem_leastc rem_leastc	591.401 187.4392
Genex: Option A			
	LeastC LeastC	rem_leastc	64.95695 44.80492
Genex: Option A	LeastC	rem_leastc rem_leastc	240.1405
Genex: Option A Genex: Option A			0.000222
	LeastC	rem_leastc	6.768497
Genex: Option A	LeastC	rem_leastc	
Genex: Option A	LeastC	rem_leastc	2.94192
Genex: Option A	LeastC	rem_leastc	4.207578
Genex: Option A	LeastC	rem_leastc	1.623035
Genex: Option A	LeastC	rem_leastc	1.398633
Genex: Option A	LeastC	rem_leastc	3.79504
Genex: Option A	LeastC	rem_leastc	2.18134
Genex: Option A	LeastC	rem_leastc	8.046262
Genex: Option A	LeastC	rem_leastc	12.96229
Genex: Option A	LeastC	rem_leastc	1.289661
Genex: Option A	LeastC	rem_leastc	79.01004
Genex: Option A	LeastC	rem_leastc	0.004241
Genex: Option A	LeastC	rem_leastc	46.31915
Genex: Option A	LeastC	rem_leastc	0.037105
Genex: Option A	LeastC	rem_leastc	10.82383
Genex: Option A	LeastC	rem_leastc	0.217139
Genex: Option A	LeastC	rem_leastc	6.178431
Genex: Option A	LeastC	rem_leastc	4.681369
Genex: Option A	LeastC	rem_leastc	7.860195
Genex: Option A	LeastC	rem_leastc	55.38093
Genex: Option A	LeastC	rem_leastc	0.075479

Genex: Option A	LeastC	rem_leastc	95.10482
Genex: Option A	LeastC	rem_leastc	81.64126
Genex: Option A	LeastC	rem_leastc	7.557274
Genex: Option A	LeastC	rem_leastc	13.22293
Genex: Option A	LeastC	rem leastc	44.78436
Genex: Option A	LeastC	rem leastc	23.87905
Genex: Option A	LeastC	rem_leastc	73.61523
Genex: Option A	LeastC	rem leastc	149.9886
Genex: Option A	LeastC	rem_leastc	74.99154
Genex: Option A	LeastC	rem_leastc	3.286117
Genex: Option A	LeastC	rem_leastc	191.8055
Genex: Option A	LeastC	rem_leastc	13.39539
Genex: Option A	LeastC	rem_leastc	60.63732
Genex: Option A	LeastC	rem_leastc	137.9151
Genex: Option A	LeastC	rem_leastc	13.08896
Genex: Option A	LeastC	rem_leastc	612.04
Genex: Option A	LeastC	rem_leastc	61.22569
Genex: Option A	LeastC	rem_leastc	0.257113
Genex: Option A	LeastC	rem_leastc	4.220058
Genex: Option A	LeastC	rem leastc	10.04527
Genex: Option A	LeastC	rem_leastc	2.741389
Genex: Option A	LeastC	rem_leastc	0.34877
Genex: Option A	LeastC	rem_leastc	0.008478
Genex: Option A	LeastC	rem_leastc	0.289362
Genex: Option A	LeastC	rem_leastc	25.32193
Genex: Option A	LeastC	rem_leastc	200.9618
Genex: Option A	LeastC	rem_leastc	53.11645
Genex: Option A	LeastC	rem leastc	37.14543
Genex: Option A	LeastC	rem leastc	32.21486
Genex: Option A	LeastC	rem_leastc	46.3546
Genex: Option A	LeastC	rem leastc	0.008122
Genex: Option A	LeastC	rem leastc	44.16289
		-	
Genex: Option A	LeastC	rem_leastc	5.923245
Genex: Option A	LeastC	rem_leastc	3.490973
Genex: Option A	LeastC	rem_leastc	131.4287
Genex: Option A	LeastC	rem_leastc	8.152129
Genex: Option A	LeastC	rem_leastc	63.09774
Genex: Option A	LeastC	rem_leastc	1.551911
Genex: Option A	LeastC	rem_leastc	6.945101
Genex: Option A	LeastC	rem_leastc	11.23512
Genex: Option A	LeastC	rem leastc	3.025342
Genex: Option A	LeastC	rem_leastc	96.39073
Genex: Option A	LeastC	rem leastc	15.80917
Genex: Option A	LeastC	rem leastc	192.9281
			0.037976
Genex: Option A	LeastC	rem_leastc	
Genex: Option A	LeastC	rem_leastc	49.50354
Genex: Option A	LeastC	rem_leastc	125.1382
Genex: Option A	LeastC	rem_leastc	20.527
Genex: Option A	LeastC	rem_leastc	2.774712
Genex: Option A	LeastC	rem_leastc	102.0444
Genex: Option A	LeastC	rem_leastc	4.059098
Genex: Option A	LeastC	rem_leastc	0.453452
Genex: Option A	LeastC	rem_leastc	12.2734
Genex: Option A	LeastC	rem_leastc	8.720471
Genex: Option A	LeastC	rem_leastc	9.605091
Genex: Option A	LeastC		0.326735
		rem_leastc	
Genex: Option A	LeastC	rem_leastc	0.727728
Genex: Option A	LeastC	rem_leastc	0.288437
Genex: Option A	LeastC	rem_leastc	13.82986
Genex: Option A	LeastC	rem_leastc	57.22385
Genex: Option A	LeastC	rem_leastc	4.842688
Genex: Option A	LeastC	rem_leastc	32.01963
Genex: Option A	LeastC	rem_leastc	0.78514
Genex: Option A	LeastC	rem_leastc	31.26611
Genex: Option A	LeastC	rem_leastc	0.826022
		rem_leastc	0.584156
(Penex. (Intion V	east(	I CITI I Casto	0.004100
Genex: Option A	LeastC		0 400001
Genex: Option A	LeastC	rem_leastc	0.600281
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	6.922604
Genex: Option A Genex: Option A Genex: Option A	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	6.922604 51.43218
Genex: Option A Genex: Option A	LeastC LeastC	rem_leastc rem_leastc	6.922604

Genex: Option A	O-dom	rem_oc	100.3972	2059 Total OC-dom remnant for Option A
Genex: Option A	O-dom	rem_oc	694.5312	
Genex: Option A	O-dom	rem_oc	33.51078	
Genex: Option A	O-dom	rem_oc	118.4246	
Genex: Option A	O-dom	rem_oc	23.384	
Genex: Option A	O-dom	rem_oc	3.956503	
Genex: Option A	O-dom	rem_oc	0.446083	
Genex: Option A	O-dom	rem_oc	7.45513	
Genex: Option A	O-dom	rem_oc	0.589727	
Genex: Option A	O-dom	rem_oc	21.69239	
Genex: Option A	O-dom	rem_oc	16.01724	
Genex: Option A	O-dom	rem_oc	0.000035	
Genex: Option A	O-dom	rem_oc	3.368585	
Genex: Option A Genex: Option A	O-dom O-dom	rem_oc	7.426551 6.518863	
		rem_oc	3.000528	
Genex: Option A Genex: Option A	O-dom O-dom	rem_oc rem_oc	6.153842	
Genex: Option A	O-dom	rem_oc	26.12089	
Genex: Option A	O-dom	rem_oc	0.000001	
Genex: Option A	O-dom	rem_oc	103.7197	
Genex: Option A	0-dom	rem_oc	31.74373	
Genex: Option A	O-dom	rem_oc	0.297039	
Genex: Option A	O-dom	rem_oc	1.185374	
Genex: Option A	O-dom	rem_oc	0.483261	
Genex: Option A	O-dom	rem_oc	3.211757	
Genex: Option A	0-dom	rem_oc	1.2572	
Genex: Option A	O-dom	rem_oc	8.687105	
Genex: Option A	O-dom	rem_oc	1.990018	
Genex: Option A	O-dom	rem_oc	0.630736	
Genex: Option A	0-dom	rem_oc	14.58421	
Genex: Option A	O-dom	rem_oc	0.181752	
Genex: Option A	O-dom	rem_oc	0.483626	
Genex: Option A	O-dom	rem_oc	17.55465	
Genex: Option A	O-dom	rem_oc	1.414196	
Genex: Option A	O-dom	rem_oc	6.715692	
Genex: Option A	O-dom	rem_oc	0	
Genex: Option A	O-dom	rem_oc	0.000082	
Genex: Option A	O-dom	rem_oc	0.946609	
Genex: Option A Genex: Option A	O-dom O-dom	rem_oc	1.013452 0.720073	
Genex: Option A		rem_oc	1.988208	
Genex: Option A	O-dom O-dom	rem_oc rem_oc	5.081188	
Genex: Option A	O-dom	rem_oc	6.756952	
Genex: Option A	O-dom	rem_oc	86.20428	
Genex: Option A	O-dom	rem oc	14.47359	
Genex: Option A	O-dom	rem_oc	9.568996	
Genex: Option A	O-dom	rem_oc	27.99034	
Genex: Option A	O-dom	rem_oc	47.06895	
Genex: Option A	O-dom	rem_oc	0.925363	
Genex: Option A	O-dom	rem_oc	0.182819	
Genex: Option A	O-dom	rem_oc	0.144678	
Genex: Option A	O-dom	rem_oc	8.038469	
Genex: Option A	O-dom	rem_oc	0.001118	
Genex: Option A	O-dom	rem_oc	5.780518	
Genex: Option A	O-dom	rem_oc	2.728088	
Genex: Option A	O-dom	rem_oc	0.792089	
Genex: Option A	O-dom	rem_oc	2.647043	
Genex: Option A	O-dom	rem_oc	1.321795	
Genex: Option A	O-dom	rem_oc	1.075075	
Genex: Option A Genex: Option A	O-dom O-dom	rem_oc rem_oc	5.615905 11.75996	
Genex: Option A	O-dom	rem oc	1.028013	
Genex: Option A	O-dom	rem_oc	4.05067	
Genex: Option A	O-dom	rem_oc	9.522925	
Genex: Option A	O-dom	rem_oc	0.001392	
Genex: Option A	O-dom	rem_oc	0.003394	
Genex: Option A	O-dom	rem_oc	20.59073	
Genex: Option A	O-dom	rem_oc	0.001035	
Genex: Option A	O-dom	rem_oc	33.69548	
Genex: Option A	O-dom	rem_oc	1.381382	
Genex: Option A	O-dom	rem_oc	0.928588	
Genex: Option A	O-dom	rem_oc	0.4135	
Genex: Option A	O-dom	rem_oc	3.286292	
Genex: Option A	O-dom	rem_oc	0.813393	
Genex: Option A	O-dom	rem_oc	4.770736	
Genex: Option A	O-dom	rem_oc	21.60491	
Genex: Option A	O-dom	rem_oc	1.561019	
Genex: Option A Genex: Option A	O-dom O-dom	rem_oc	2.309776 9.712734	
Genex: Option A	O-dom	rem_oc rem_oc	5.799722	
Schok. Option A	o doni	iciii_00	5.177122	

Genex: Option A	O-dom	rem_oc	1.058237	
enex: Option A	O-dom	rem_oc	0.046029	
enex: Option A	O-dom	rem_oc	4.964713	
enex: Option A	O-dom	rem_oc	5.369112	
enex: Option A	O-dom	rem_oc	5.718127	
enex: Option A	O-dom	rem_oc	2.128498	
enex: Option A	O-dom	rem_oc	1.30439	
enex: Option A	O-dom	rem_oc	14.87732	
enex: Option A	O-dom	rem_oc	48.60826	
enex: Option A	O-dom	rem_oc	0.43457	
enex: Option A	O-dom	rem_oc	58.61665	
enex: Option A	O-dom	rem_oc	62.13515	
enex: Option A	O-dom	rem_oc	4.286207	
enex: Option A	O-dom	rem_oc	8.433967	
Senex: Option A	O-dom	rem_oc	4.418017	
Senex: Option A	O-dom	rem_oc	0.816551	
Senex: Option A	O-dom	rem_oc	1.360325	
Senex: Option A	O-dom	rem_oc	0.61658	
Senex: Option A	O-dom	rem_oc	0.594068	
Senex: Option A	O-dom	rem_oc	9.111942	
enex: Option A	O-dom	rem_oc	8.567498	
enex: Option A	O-dom	rem_oc	5.563262	
enex: Option A	O-dom	rem_oc	58.10314	
enex: Option A	O-dom	rem_oc	11.76356	
enex: Option A	O-dom	rem_oc	15.50552	
enex: Option A	O-dom	rem_oc	4.488126	
enex: Option A	O-dom	rem_oc	3.278022	
enex: Option A	O-dom	rem_oc	3.436123	
enex: Option A	O-dom	rem_oc	11.64185	
enex: Option A	O-dom	rem_oc	18.22205	
Genex: Option A	O-dom	rem_oc	0.0018	
Genex: Option A	O-dom	rem_oc	4.444007	
Genex: Option A	O-dom	rem_oc	6.059981	
Genex: Option A	O-dom	rem_oc	0.338932	
Genex: Option A	O-dom	rem_oc	15.98738	
Genex: Option A	O-dom	rem_oc	0.645188	
Genex: Option A	O-dom	rem_oc	3.072902	
Genex: Option A	O-dom	rem_oc	0.000344	
Genex: Option A	O-dom	rem_oc	9.199158	
Genex: Option A	O-dom	rem_oc	6.182994	
Genex: Option A	O-dom	rem_oc	6.221117	
Genex: Option A	O-subdom	rem_oc	20.80435	1008 Total OC-subdom remnant for Option A
Genex: Option A	O-subdom	rem_oc	130.6595	
Genex: Option A	O-subdom	rem_oc	345.7833	
Genex: Option A	O-subdom	rem_oc	48.92495	
Genex: Option A	O-subdom	rem oc	18.5864	
Genex: Option A	O-subdom	rem oc	348.0027	
Genex: Option A	O-subdom	rem oc	95.19846	
	2 22.540111			

## Kidston Corridor Option B - Regional ecosystems v8

Name	VM_POLY	VM_STATUS	Area_ha	
Canau Ontion D		hum la sata	11 7/1//	202 Tatal I Conservation D
Genex: Option B	LeastC	hvr_leastc	11.76166	202 Total LC regrowth for Option B
Genex: Option B	LeastC	hvr_leastc	0.071373 0.032936	
Genex: Option B	LeastC LeastC	hvr_leastc	0.032936	
Genex: Option B Genex: Option B	LeastC	hvr_leastc hvr_leastc	4.030559	
Genex: Option B	LeastC	hvr_leastc	4.030339	
Genex: Option B	LeastC	hvr_leastc	2.294056	
Genex: Option B	LeastC	hvr_leastc	1.055631	
Genex: Option B	LeastC	hvr_leastc	1.583837	
Genex: Option B	LeastC	hvr_leastc	4.886182	
Genex: Option B	LeastC	hvr_leastc	0.483842	
Genex: Option B	LeastC	hvr leastc	0.022396	
Genex: Option B	LeastC	hvr_leastc	16.55398	
Genex: Option B	LeastC	hvr_leastc	0.847925	
Genex: Option B	LeastC	hvr_leastc	0.452182	
Genex: Option B	LeastC	hvr_leastc	0.355135	
Genex: Option B	LeastC	hvr_leastc	1.329568	
Genex: Option B	LeastC	hvr_leastc	7.449196	
Genex: Option B	LeastC	hvr_leastc	10.2054	
Genex: Option B	LeastC	hvr_leastc	0.252609	
Genex: Option B	LeastC	hvr_leastc	0.004932	
Genex: Option B	LeastC	hvr_leastc	2.193916	
Genex: Option B	LeastC	hvr_leastc	0.628194	
Genex: Option B	LeastC	hvr_leastc	4.798001	
Genex: Option B	LeastC	hvr_leastc	4.892764	
Genex: Option B	LeastC	hvr_leastc	0.857374	
Genex: Option B	LeastC	hvr_leastc	0.016948	
Genex: Option B	LeastC	hvr_leastc	0.026688	
Genex: Option B	LeastC	hvr_leastc	0.489348	
Genex: Option B	LeastC	hvr_leastc	0.001268	
Genex: Option B	LeastC	hvr_leastc	0.733646	
Genex: Option B	LeastC	hvr_leastc	0.001018	
Genex: Option B	LeastC	hvr_leastc	2.958507	
Genex: Option B	LeastC	hvr_leastc	2.596304	
Genex: Option B	LeastC	hvr_leastc	2.274126	
Genex: Option B	LeastC	hvr_leastc	0.860982	
Genex: Option B	LeastC	hvr_leastc	0.17804	
Genex: Option B	LeastC	hvr_leastc	1.002297	
Genex: Option B	LeastC	hvr_leastc	0.00441	
Genex: Option B	LeastC	hvr_leastc	13.17734	
Genex: Option B	LeastC	hvr_leastc	0.224692	
Genex: Option B	LeastC	hvr_leastc	1.836962	
Genex: Option B	LeastC	hvr_leastc	0.523549	
Genex: Option B	LeastC	hvr_leastc	5.202041	
Genex: Option B	LeastC	hvr_leastc	0.41416	
Genex: Option B	LeastC	hvr_leastc	0.004426	
Genex: Option B	LeastC	hvr_leastc	0.175718	
Genex: Option B	LeastC	hvr_leastc	1.371884	
Genex: Option B	LeastC	hvr_leastc	0.720414	
Genex: Option B	LeastC	hvr_leastc	0.086502	
Genex: Option B	LeastC	hvr_leastc	0.349904	
Genex: Option B	LeastC	hvr_leastc	9.197092	
Genex: Option B	LeastC	hvr_leastc	2.849699 1.297257	
Genex: Option B Genex: Option B	LeastC	hvr_leastc		
	LeastC	hvr_leastc	63.51923	
Genex: Option B	LeastC	hvr_leastc	0.000024	
Genex: Option B Genex: Option B	LeastC	hvr_leastc	0.000132	
	LeastC	hvr_leastc	0.000386	
Genex: Option B	LeastC	hvr_leastc	2.657052 1.646678	
Genex: Option B	LeastC	hvr_leastc	3.808301	
Genex: Option B Genex: Option B	LeastC	hvr_leastc	0.001706	
Genex. Option B	LeastC	hvr_leastc	0.001706	

Genex: Option B	O-dom	hvr_oc	0.888013	2.4 Total OC-dom HVR regrowth for Option B
Genex: Option B	O-dom	hvr_oc	0.034724	
Genex: Option B	O-dom	hvr_oc	0.084695	
Genex: Option B	O-dom	hvr_oc	0.476706	
Genex: Option B	O-dom	hvr_oc	0.890819	
Genex: Option B	non-rem	non_remnant	12.82744	2379 Total non-remnant for Option B
Genex: Option B	non-rem	non_remnant	1.191201	
Genex: Option B	non-rem	non_remnant	3.098019	
Genex: Option B	non-rem	non_remnant	6.002011	
Genex: Option B	non-rem	non_remnant non_remnant	10.64987 18.11507	
Genex: Option B Genex: Option B	non-rem non-rem	non_remnant	7.444218	
Genex: Option B	non-rem	non_remnant	103.7112	
Genex: Option B	non-rem	non_remnant	1.27489	
Genex: Option B	non-rem	non_remnant	1.573922	
Genex: Option B	non-rem	non_remnant	0.416775	
Genex: Option B	non-rem	non_remnant	0.284209	
Genex: Option B	non-rem	non_remnant	0.934327	
Genex: Option B	non-rem	non_remnant	32.73321	
Genex: Option B	non-rem	non_remnant	0.21711	
Genex: Option B	non-rem	non_remnant	7.341556	
Genex: Option B	non-rem	non_remnant	1.536304	
Genex: Option B Genex: Option B	non-rem non-rem	non_remnant non_remnant	1.624022 1.525838	
Genex: Option B	non-rem	non_remnant	6.35254	
Genex: Option B	non-rem	non remnant	0.311335	
Genex: Option B	non-rem	non_remnant	7.245895	
Genex: Option B	non-rem	non_remnant	8.656713	
Genex: Option B	non-rem	non_remnant	0.284912	
Genex: Option B	non-rem	non_remnant	0.799373	
Genex: Option B	non-rem	non_remnant	1.586069	
Genex: Option B	non-rem	non_remnant	393.879	
Genex: Option B	non-rem	non_remnant	106.2974	
Genex: Option B	non-rem	non_remnant	3.895228	
Genex: Option B Genex: Option B	non-rem non-rem	non_remnant non_remnant	288.9389 4.258125	
Genex: Option B	non-rem	non_remnant	0.056622	
Genex: Option B	non-rem	non_remnant	0.167595	
Genex: Option B	non-rem	non_remnant	0.535546	
Genex: Option B	non-rem	non_remnant	0.756154	
Genex: Option B	non-rem	non_remnant	103.4499	
Genex: Option B	non-rem	non_remnant	6.867542	
Genex: Option B	non-rem	non_remnant	347.132	
Genex: Option B	non-rem	non_remnant	0.000275	
Genex: Option B	non-rem	non_remnant	0.000386	
Genex: Option B Genex: Option B	non-rem	non_remnant	0.234067 282.7614	
Genex: Option B	non-rem non-rem	non_remnant non_remnant	0.000204	
Genex: Option B	non-rem	non_remnant	0.000204	
Genex: Option B	non-rem	non_remnant	8.914747	
Genex: Option B	non-rem	non_remnant	1.286566	
Genex: Option B	non-rem	non_remnant	1.685735	
Genex: Option B	non-rem	non_remnant	0.526879	
Genex: Option B	non-rem	non_remnant	0.121333	
Genex: Option B	non-rem	non_remnant	3.735213	
Genex: Option B	non-rem	non_remnant	117.347	
Genex: Option B	non-rem	non_remnant	468.6799	
Genex: Option B	non-rem	non_remnant	0.001706 87.63232	117 Total water non-remnant for Option B
Genex: Option B Genex: Option B	water water	non_remnant non_remnant	0.341308	
Genex: Option B	water	non_remnant	11.9254	
Genex: Option B	water	non_remnant	17.22085	
Genex: Option B	LeastC	rem_leastc	26.95741	51121 Total LC remnant for Option B
Genex: Option B	LeastC	rem_leastc	7.493854	
Genex: Option B	LeastC	rem_leastc	255.6705	
Genex: Option B	LeastC	rem_leastc	99.21011	
Genex: Option B	LeastC	rem_leastc	337.5413	
Genex: Option B	LeastC	rem_leastc	19.5728	
Genex: Option B	LeastC	rem_leastc	2.196989	
Genex: Option B	LeastC	rem_leastc	156.2433	

Genex: Option B	LeastC	rem_leastc	2.363897
Genex: Option B	LeastC	rem_leastc	59.46574
Genex: Option B	LeastC	rem_leastc	326.0669
Genex: Option B	LeastC	rem_leastc	42.58946
Genex: Option B	LeastC	rem_leastc	79.24012
Genex: Option B	LeastC	rem_leastc	3.026556
Genex: Option B	LeastC	rem_leastc	14.37214
Genex: Option B	LeastC	rem_leastc	79.46571
Genex: Option B	LeastC	rem_leastc	169.7709
Genex: Option B	LeastC	rem_leastc	188.8082
Genex: Option B	LeastC	rem_leastc	1.809083
Genex: Option B	LeastC	rem_leastc	4.936312
Genex: Option B	LeastC	rem_leastc	41.46687
Genex: Option B	LeastC	rem_leastc	26.04
Genex: Option B	LeastC	rem leastc	28.90649
Genex: Option B	LeastC	rem leastc	55.9325
Genex: Option B	LeastC	rem_leastc	3.377138
Genex: Option B	LeastC	rem leastc	13.57611
Genex: Option B	LeastC	rem leastc	92.17016
Genex: Option B	LeastC	rem_leastc	27.56209
Genex: Option B	LeastC	rem_leastc	24.13472
Genex: Option B	LeastC	rem leastc	36.77539
Genex: Option B	LeastC	rem leastc	26.5067
Genex: Option B	LeastC	rem_leastc	0.114053
Genex: Option B	LeastC	rem_leastc	90.00732
Genex: Option B	LeastC	rem_leastc	8.935729
Genex: Option B	LeastC	rem_leastc	62.61951
Genex: Option B	LeastC	rem_leastc	502.94
Genex: Option B	LeastC	rem_leastc	41.45126
Genex: Option B	LeastC	rem_leastc	27.55622
Genex: Option B	LeastC	rem_leastc	457.4285
Genex: Option B	LeastC	rem leastc	604.3354
Genex: Option B	LeastC	rem_leastc	13.58892
Genex: Option B	LeastC	rem_leastc	23.97449
Genex: Option B	LeastC	rem leastc	1.36276
Genex: Option B	LeastC	rem leastc	41.44455
Genex: Option B	LeastC	rem_leastc	766.9081
Genex: Option B	LeastC		7.498845
Genex: Option B		rem_leastc	1.855243
	LeastC	rem_leastc	
Genex: Option B	LeastC	rem_leastc	98.11036
Genex: Option B	LeastC	rem_leastc	11.92043
Genex: Option B	LeastC	rem_leastc	2.674089
Genex: Option B	LeastC	rem_leastc	421.3659
Genex: Option B	LeastC	rem_leastc	15.33338
Genex: Option B	LeastC	rem_leastc	0.033028
Genex: Option B	LeastC	rem_leastc	15.38248
Genex: Option B	LeastC	rem_leastc	602.6281
Genex: Option B	LeastC	rem_leastc	539.0101
Genex: Option B	LeastC	rem_leastc	45.58743
Genex: Option B	LeastC	rem_leastc	54.55858
Genex: Option B	LeastC	rem_leastc	26.25275
Genex: Option B	LeastC	rem_leastc	24.49658
Genex: Option B	LeastC	rem_leastc	4.012632
Genex: Option B	LeastC	rem_leastc	32.58558
Genex: Option B	LeastC	rem_leastc	2.735045
Genex: Option B	LeastC	rem_leastc	329.6436
Genex: Option B	LeastC	rem_leastc	78.50426
Genex: Option B	LeastC	rem_leastc	15.35245
Genex: Option B	LeastC	rem_leastc	50.55502
Genex: Option B	LeastC	rem_leastc	54.24811
Genex: Option B	LeastC	rem_leastc	106.3064
Genex: Option B	LeastC	rem_leastc	215.6079
Genex: Option B	LeastC	rem_leastc	2.555068
Genex: Option B	LeastC	rem_leastc	13.52695
Genex: Option B	LeastC	rem_leastc	100.1217
Genex: Option B	LeastC	rem_leastc	267.2011
Genex: Option B	LeastC	rem_leastc	7.561765
Genex: Option B	LeastC	rem_leastc	563.416
Genex: Option B	LeastC	rem_leastc	13.37656
Genex: Option B	LeastC	rem_leastc	698.7491
Genex: Option B	LeastC	rem_leastc	6.465764
Genex: Option B	LeastC	rem_leastc	43.71799
Genex: Option B	LeastC	rem_leastc	426.9156
0 0 11 D	LeastC	rem_leastc	142.4977
Genex: Option B	Leasic		
Genex: Option B Genex: Option B	LeastC	rem_leastc	36.24586
	LeastC LeastC	rem_leastc rem_leastc	36.24586 0.351853
Genex: Option B	LeastC		

Genex: Option B	LeastC	rem_leastc	43.78872
Genex: Option B	LeastC	rem leastc	16.76179
Genex: Option B	LeastC	rem leastc	236.8411
Genex: Option B	LeastC	rem_leastc	45.65808
Genex: Option B	LeastC	rem leastc	33.08399
Genex: Option B	LeastC	rem leastc	18.19931
Genex: Option B	LeastC	rem leastc	74.95996
Genex: Option B	LeastC	rem_leastc	6.095082
Genex: Option B	LeastC	rem leastc	77.06789
Genex: Option B	LeastC	rem leastc	80.45791
Genex: Option B	LeastC	rem_leastc	1.01019
Genex: Option B	LeastC	rem_leastc	72.92622
Genex: Option B	LeastC	rem leastc	82.15856
Genex: Option B	LeastC	rem leastc	368.5145
Genex: Option B	LeastC	rem_leastc	53.66492
Genex: Option B	LeastC	rem_leastc	40.47175
Genex: Option B	LeastC	rem_leastc	88.70847
Genex: Option B	LeastC	rem leastc	17.2582
Genex: Option B	LeastC	rem_leastc	40.4714
Genex: Option B	LeastC	rem_leastc	30.47152
Genex: Option B	LeastC	rem_leastc	4.376344
Genex: Option B	LeastC	rem_leastc	349.1754
Genex: Option B	LeastC	rem leastc	45.56187
Genex: Option B	LeastC	rem_leastc	366.7211
Genex: Option B	LeastC	rem_leastc	27.30528
Genex: Option B	LeastC	rem leastc	237.8945
Genex: Option B	LeastC	rem_leastc	28.61304
Genex: Option B	LeastC	rem_leastc	21.1362
Genex: Option B	LeastC	rem_leastc	144.4847
Genex: Option B	LeastC	rem leastc	110.4779
Genex: Option B	LeastC	rem_leastc	30.2618
Genex: Option B	LeastC	rem_leastc	9.529117
Genex: Option B	LeastC	rem_leastc	369.3495
Genex: Option B	LeastC	rem leastc	302.0024
Genex: Option B	LeastC	rem_leastc	221.8051
Genex: Option B	LeastC	rem_leastc	394.3924
Genex: Option B	LeastC	rem_leastc	813.6089
Genex: Option B	LeastC	rem_leastc	1336.278
Genex: Option B	LeastC	rem leastc	6.820337
Genex: Option B	LeastC	rem_leastc	42.57597
Genex: Option B	LeastC	rem leastc	190.512
Genex: Option B	LeastC	rem leastc	70.1625
Genex: Option B	LeastC	rem_leastc	338.8364
Genex: Option B	LeastC	rem_leastc	4.722484
Genex: Option B	LeastC	rem leastc	12.75251
Genex: Option B	LeastC	rem leastc	11.71659
Genex: Option B	LeastC	rem leastc	11.7374
Genex: Option B	LeastC	rem leastc	5.838685
Genex: Option B	LeastC	rem_leastc	19.6216
Genex: Option B	LeastC	rem_leastc	12.0518
IGCHEA, ODHOLLD	LeastC	rem leastc	
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189
Genex: Option B	LeastC	rem_leastc	
			6.919189 54.14043
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039
Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1760 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991 216.618
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991 216.618 2.482518
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991 216.618 2.482518 192.9534
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.17609 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991 216.618 2.482518 192.9534 108.3536
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.919189 54.14043 19.80428 78.62039 15.51213 117.7604 165.1769 197.739 482.4364 66.28895 0.199872 21.28374 219.6841 76.77679 10.29573 380.5413 92.48148 6.691739 526.1154 428.0187 27.47299 34.75991 216.618 2.482518 192.9534

Genex: Option B	LeastC	rem_leastc	0.004241
Genex: Option B	LeastC	rem leastc	193.8193
Genex: Option B	LeastC	rem leastc	183.4683
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Genex: Option B	LeastC	rem_leastc	105.8241
Genex: Option B	LeastC	rem_leastc	30.87396
Genex: Option B	LeastC	rem_leastc	437.6609
Genex: Option B	LeastC	rem_leastc	26.50336
Genex: Option B	LeastC	rem_leastc	43.01024
Genex: Option B	LeastC	rem_leastc	2.674355
Genex: Option B	LeastC	rem_leastc	145.642
Genex: Option B	LeastC	rem leastc	45.83799
Genex: Option B	LeastC	rem_leastc	1.750816
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Genex: Option B	LeastC	rem_leastc	7.330887
Genex: Option B	LeastC	rem_leastc	80.91222
Genex: Option B	LeastC	rem_leastc	285.9006
Genex: Option B	LeastC	rem_leastc	1.802566
Genex: Option B	LeastC	rem_leastc	59.32959
Genex: Option B	LeastC	rem_leastc	12.23004
Genex: Option B	LeastC	rem leastc	8.971527
Genex: Option B	LeastC	rem_leastc	13.00588
Genex: Option B	LeastC	rem_leastc	3.75467
		_	
Genex: Option B	LeastC	rem_leastc	178.9832
Genex: Option B	LeastC	rem_leastc	107.5745
Genex: Option B	LeastC	rem_leastc	61.24909
Genex: Option B	LeastC	rem_leastc	88.93272
Genex: Option B	LeastC	rem_leastc	8.600226
Genex: Option B	LeastC	rem_leastc	42.1064
Genex: Option B	LeastC	rem_leastc	526.2937
Genex: Option B	LeastC	rem_leastc	4.800436
Genex: Option B	-		12.7743
	LeastC	rem_leastc	
Genex: Option B	LeastC	rem_leastc	18.0793
Genex: Option B	LeastC	rem_leastc	70.34255
Genex: Option B	LeastC	rem_leastc	47.25853
Genex: Option B	LeastC	rem_leastc	289.9992
Genex: Option B	LeastC	rem_leastc	2.945584
Genex: Option B	LeastC	rem leastc	0.027143
Genex: Option B	LeastC	rem_leastc	22.94944
Genex: Option B	LeastC	rem_leastc	241.1351
Genex: Option B	LeastC	_	0.037105
	-	rem_leastc	
Genex: Option B	LeastC	rem_leastc	1.018066
Genex: Option B	LeastC	rem_leastc	148.4483
Genex: Option B	LeastC	rem_leastc	1.735207
Genex: Option B	LeastC	rem_leastc	200.3061
Genex: Option B	LeastC	rem_leastc	6.178431
Genex: Option B	LeastC	rem leastc	0.283346
Genex: Option B	LeastC	rem leastc	63.82911
Genex: Option B	LeastC	rem_leastc	9.483077
Genex: Option B	LeastC	rem_leastc	5.112733
Genex: Option B	LeastC	rem_leastc	42.39342
Genex: Option B	LeastC	rem_leastc	217.1046
Genex: Option B	LeastC	rem_leastc	274.7574
Genex: Option B	LeastC	rem_leastc	44.75944
Genex: Option B	LeastC	rem_leastc	34.00366
Genex: Option B	LeastC	rem_leastc	15.84078
Genex: Option B	LeastC	rem leastc	44 64747
Genex: Option B	LeastC	rem_leastc	44.64747
Genex: Option B	LeastC	rem_leastc	122.9537
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	122.9537 566.3628
Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805 543.976
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 1.813457 14.67879 36.05249 345.0211 166.6805 543.976 7.497423
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805 543.976 7.497423 441.0962
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805 543.976 7.497423 441.0962 13.40562
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805 543.976 7.497423 441.0962 1.3.40562 1.641803
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	122.9537 566.3628 14.21165 28.92059 237.4161 82.96334 60.5029 45.99594 87.13361 54.99165 47.59365 1.813457 14.67879 36.05249 345.0211 166.6805 543.976 7.497423 441.0962 13.40562

Genex: Option B	LeastC	rom looste	70.1032
Genex: Option B	LeastC	rem_leastc rem_leastc	46.96682
Genex: Option B	LeastC	rem_leastc	247.8017
Genex: Option B	LeastC	rem_leastc	164.7249
Genex: Option B	LeastC	rem leastc	7.676785
Genex: Option B	LeastC	rem_leastc	497.7703
Genex: Option B	LeastC	rem leastc	73.2098
Genex: Option B	LeastC	rem leastc	15.16272
Genex: Option B	LeastC	rem leastc	30.13856
Genex: Option B	LeastC	rem_leastc	24.79674
Genex: Option B	LeastC	rem_leastc	12.80719
Genex: Option B	LeastC	rem leastc	30.18079
Genex: Option B	LeastC	rem leastc	50.3982
Genex: Option B	LeastC	rem leastc	2.007452
Genex: Option B	LeastC	rem_leastc	5.297635
Genex: Option B	LeastC	rem leastc	4.628709
Genex: Option B	LeastC	rem_leastc	23.61133
Genex: Option B	LeastC	rem_leastc	73.136
Genex: Option B	LeastC	rem_leastc	95.10482
Genex: Option B	LeastC	rem_leastc	875.1531
Genex: Option B	LeastC	rem leastc	5.458226
Genex: Option B	LeastC	rem leastc	5.761792
Genex: Option B	LeastC	rem_leastc	61.16559
Genex: Option B	LeastC	rem_leastc	19.86604
Genex: Option B	LeastC	rem_leastc	144.7855
Genex: Option B	LeastC	rem_leastc	40.26181
Genex: Option B	LeastC	rem_leastc	239.4669
Genex: Option B	LeastC	rem_leastc	135.8535
Genex: Option B	LeastC	rem_leastc	6.56791
Genex: Option B	LeastC	rem_leastc	1.495947
Genex: Option B	LeastC	rem_leastc	42.50296
Genex: Option B	LeastC	rem_leastc	81.64126
Genex: Option B	LeastC	rem_leastc	0.284174
Genex: Option B	LeastC	rem_leastc	342.7593
Genex: Option B	LeastC	rem_leastc	90.20453
Genex: Option B	LeastC	rem_leastc	11.22949
Genex: Option B	LeastC	rem_leastc	370.7128
Genex: Option B	LeastC	rem_leastc	2.448721
Genex: Option B	LeastC	rem_leastc	89.05145
Genex: Option B	LeastC	rem_leastc	87.37309
Genex: Option B	LeastC	rem_leastc	11.05859
Genex: Option B	LeastC	rem_leastc	493.3194
Genex: Option B	LeastC	rem_leastc	53.89835
Genex: Option B	LeastC	rem_leastc	17.69808
Genex: Option B	LeastC	rem_leastc	7.514547
Genex: Option B	LeastC	rem_leastc	7.557274
Genex: Option B	LeastC	rem_leastc	508.631
Genex: Option B	LeastC	rem_leastc	427.6424
Genex: Option B	LeastC	rem_leastc	104.0435
Genex: Option B	LeastC	rem_leastc	13.22293
Genex: Option B	LeastC	rem_leastc	4.416903
Genex: Option B	LeastC	rem_leastc	0.859066
Genex: Option B	LeastC	rem_leastc	44.88932
Genex: Option B	LeastC	rem_leastc	10.31003
Genex: Option B	LeastC	rem_leastc	3.907877
Genex: Option B	LeastC	rem_leastc	0.286307
Genex: Option B	LeastC	rem_leastc	3.536081
Genex: Option B	LeastC	rem_leastc	4.302634
Genex: Option B	LeastC	rem_leastc	40.54574
Genex: Option B	LeastC	rem_leastc	46.35656
Genex: Option B	LeastC	rem_leastc	35.1372
Genex: Option B	LeastC	rem_leastc	21.46795
Genex: Option B	LeastC	rem_leastc	63.82186
Genex: Option B Genex: Option B	LeastC	rem_leastc	6.597762
Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	6.597762 22.59072
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114 17.66116
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114 17.66116 9.664155
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114 17.66116 9.664155 27.0727
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114 17.66116 9.664155 27.0727 47.85526
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	6.597762 22.59072 14.81235 25.26813 44.78436 15.78862 488.7074 91.32114 17.66116 9.664155 27.0727

Genex: Option B         LeastC         rem_leastc         0.426342           Genex: Option B         LeastC         rem_leastc         167.5648           Genex: Option B         LeastC         rem_leastc         0.0236342           Genex: Option B         LeastC         rem_leastc         0.043366           Genex: Option B         LeastC         rem_leastc         0.043366           Genex: Option B         LeastC         rem_leastc         13.3243           Genex: Option B         LeastC         rem_leastc         13.3243           Genex: Option B         LeastC         rem_leastc         13.3243           Genex: Option B         LeastC         rem_leastc         2.83706           Genex: Option B         LeastC         rem_leastc         2.83706           Genex: Option B         LeastC         rem_leastc         17.8677           Genex: Option B         LeastC         rem_leastc         <	Genex: Option B	LeastC	rem_leastc	145.1778
Genex: Option B         LeastC         rem_leastc         0.226342           Genex: Option B         LeastC         rem_leastc         15.25302           Genex: Option B         LeastC         rem_leastc         0.043366           Genex: Option B         LeastC         rem_leastc         0.24366           Genex: Option B         LeastC         rem_leastc         0.24366           Genex: Option B         LeastC         rem_leastc         13.8242           Genex: Option B         LeastC         rem_leastc         7.69207           Genex: Option B         LeastC         rem_leastc         7.69207           Genex: Option B         LeastC         rem_leastc         0.739702           Genex: Option B         LeastC         rem_leastc         0.739702           Genex: Option B         LeastC         rem_leastc         0.739702           Genex: Option B         LeastC         rem_leastc         10.749471           Genex: Option B         LeastC         rem_leastc         13.80472           Genex: Option B         LeastC         rem_leastc         14.744071           Genex: Option B         LeastC         rem_leastc         14.744071           Genex: Option B         LeastC         rem_leastc			_	64.6371
Genex: Option B         LeastC         rem_leastc         115.2530           Genex: Option B         LeastC         rem_leastc         0.043365           Genex: Option B         LeastC         rem_leastc         0.243965           Genex: Option B         LeastC         rem_leastc         121.50385           Genex: Option B         LeastC         rem_leastc         121.50385           Genex: Option B         LeastC         rem_leastc         7.692077           Genex: Option B         LeastC         rem_leastc         7.692077           Genex: Option B         LeastC         rem_leastc         2.839104           Genex: Option B         LeastC         rem_leastc         2.943985           Genex: Option B         LeastC         rem_leastc         2.943985           Genex: Option B         LeastC         rem_leastc         7.05411           Genex: Option B         LeastC         rem_leastc         7.05411           Genex: Option B         LeastC         rem_leastc         7.843586           Genex: Option B         LeastC         rem_leastc         7.843586           Genex: Option B         LeastC         rem_leastc         7.843586           Genex: Option B         LeastC         rem_leastc				
Genex: Option B         LeastC         rem_leastC         0.04336           Genex: Option B         LeastC         rem_leastC         0.24396           Genex: Option B         LeastC         rem_leastC         0.24396           Genex: Option B         LeastC         rem_leastC         13.8243           Genex: Option B         LeastC         rem_leastC         7.69207           Genex: Option B         LeastC         rem_leastC         7.69207           Genex: Option B         LeastC         rem_leastC         0.739912           Genex: Option B         LeastC         rem_leastC         0.74490           Genex: Option B         LeastC         rem_leastC         17.64403           Genex: Option B         LeastC         rem_leastC         13.86712           Genex: Option B         LeastC         rem_leastC         13.86712           Genex: Option B         LeastC         rem_leastC         13.86712           Genex: Option B         LeastC         rem_leastC				
Genex: Option B         LeastC         rem_leastC         0.043366           Genex: Option B         LeastC         rem_leastC         31.21996           Genex: Option B         LeastC         rem_leastC         31.21996           Genex: Option B         LeastC         rem_leastC         13.38243           Genex: Option B         LeastC         rem_leastC         348943           Genex: Option B         LeastC         rem_leastC         348943           Genex: Option B         LeastC         rem_leastC         2.83910           Genex: Option B         LeastC         rem_leastC         2.83940           Genex: Option B         LeastC         rem_leastC         2.83940           Genex: Option B         LeastC         rem_leastC         17.49671           Genex: Option B         LeastC         rem_leastC         17.49671           Genex: Option B         LeastC         rem_leastC         17.49671           Genex: Option B         LeastC         rem_leastC         2.83780           Genex: Option B         LeastC         rem_leastC         2.83641           Genex: Option B         LeastC         rem_leastC         2.83641           Genex: Option B         LeastC         rem_leastC <t< td=""><td></td><td></td><td>_</td><td></td></t<>			_	
Genex: Option B         LeastC         rem_leastc         0.249365           Genex: Option B         LeastC         rem_leastc         21.50388           Genex: Option B         LeastC         rem_leastc         1.50388           Genex: Option B         LeastC         rem_leastc         7.692077           Genex: Option B         LeastC         rem_leastc         7.692077           Genex: Option B         LeastC         rem_leastc         2.994386           Genex: Option B         LeastC         rem_leastc         0.739912           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.64518           Genex: Option B         LeastC         rem_leastc         12.645183           Genex: Option B         LeastC         rem_leastc         22.834388           Genex: Option B         LeastC         rem_leastc         22.834388           Genex: Option B         LeastC         rem_leastc         22.834388           Genex: Option B         LeastC         rem_leastc <td></td> <td></td> <td></td> <td></td>				
Cenex: Option B         LeastC         rem_leastC         31.21998           Genex: Option B         LeastC         rem_leastC         13.38243           Genex: Option B         LeastC         rem_leastC         13.38243           Genex: Option B         LeastC         rem_leastC         3.140981           Genex: Option B         LeastC         rem_leastC         3.140981           Genex: Option B         LeastC         rem_leastC         2.89704           Genex: Option B         LeastC         rem_leastC         2.994385           Genex: Option B         LeastC         rem_leastC         2.994385           Genex: Option B         LeastC         rem_leastC         17.96717           Genex: Option B         LeastC         rem_leastC         17.96717           Genex: Option B         LeastC         rem_leastC         2.83586           Genex: Option B         LeastC         rem_leastC         2.84588           Genex: Option B         LeastC         rem_leastC         2.83641           Genex: Option B         LeastC         rem_leastC         2.83641           Genex: Option B         LeastC         rem_leastC         2.83641           Genex: Option B         LeastC         rem_leastC				
Genex: Option B         LeastC         rem_leastc         21.5038           Genex: Option B         LeastC         rem_leastc         48.91066           Genex: Option B         LeastC         rem_leastc         3.8242           Genex: Option B         LeastC         rem_leastc         2.839104           Genex: Option B         LeastC         rem_leastc         2.839104           Genex: Option B         LeastC         rem_leastc         2.839104           Genex: Option B         LeastC         rem_leastc         0.739912           Genex: Option B         LeastC         rem_leastc         17.40671           Genex: Option B         LeastC         rem_leastc         17.34671           Genex: Option B         LeastC         rem_leastc         17.34672           Genex: Option B         LeastC         rem_leastc         2.84358           Genex: Option B         LeastC         rem_leastc         2.84358           Genex: Option B         LeastC         rem_leastc         2.84358           Genex: Option B         LeastC         rem_leastc         2.03065           Genex: Option B         LeastC         rem_leastc         149.9884           Genex: Option B         LeastC         rem_leastc				
Genex: Option B         LeastC         rem_leastC         13.3824           Genex: Option B         LeastC         rem_leastC         7.67207           Genex: Option B         LeastC         rem_leastC         3.140981           Genex: Option B         LeastC         rem_leastC         2.839104           Genex: Option B         LeastC         rem_leastC         6.5.74696           Genex: Option B         LeastC         rem_leastC         0.739917           Genex: Option B         LeastC         rem_leastC         17.46677           Genex: Option B         LeastC         rem_leastC         17.6677           Genex: Option B         LeastC         rem_leastC         17.6677           Genex: Option B         LeastC         rem_leastC         17.6677           Genex: Option B         LeastC         rem_leastC         17.6122           Genex: Option B         LeastC         rem_leastC         12.73818           Genex: Option B         LeastC         rem_leastC         13.38.616           Genex: Option B         LeastC         rem_leastC         149.9886           Genex: Option B         LeastC         rem_leastC         149.9866           Genex: Option B         LeastC         rem_leastC			_	
Genex: Option B         LeastC         rem_leastc         48.91066           Genex: Option B         LeastC         rem_leastc         3.140981           Genex: Option B         LeastC         rem_leastc         2.83910           Genex: Option B         LeastC         rem_leastc         2.83910           Genex: Option B         LeastC         rem_leastc         2.994388           Genex: Option B         LeastC         rem_leastc         2.84948           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         13.06975           Genex: Option B         LeastC         rem_leastc         13.06975           Genex: Option B         LeastC         rem_leastc         13.06975           Genex: Option B         LeastC         rem_leastc         13.83.012           Genex: Option B         LeastC         rem_leastc         13.83.012           Genex: Option B         LeastC         rem_leastc         13.43.936           Genex: Option B         LeastC         rem_leastc         14.94.9886           Genex: Option B         LeastC         rem_leastc         14.94.9866           Genex: Option B         LeastC         rem_leastc </td <td></td> <td></td> <td></td> <td></td>				
Genex: Option B         LeastC         rem_leastC         7.692077           Genex: Option B         LeastC         rem_leastC         3.140981           Genex: Option B         LeastC         rem_leastC         2.839104           Genex: Option B         LeastC         rem_leastC         2.994386           Genex: Option B         LeastC         rem_leastC         2.97396           Genex: Option B         LeastC         rem_leastC         7.073971           Genex: Option B         LeastC         rem_leastC         7.05411           Genex: Option B         LeastC         rem_leastC         7.46407           Genex: Option B         LeastC         rem_leastC         2.843686           Genex: Option B         LeastC         rem_leastC         2.84386           Genex: Option B         LeastC         rem_leastC         2.70387           Genex: Option B         LeastC         rem_leastC         2.73386           Genex: Option B         LeastC         rem_leastC         2.73387           Genex: Option B         LeastC         rem_leastC         2.749155           Genex: Option B         LeastC         rem_leastC         2.749155           Genex: Option B         LeastC         rem_leastC				
Genex: Option B         LeastC         rem_leastc         3.140981           Genex: Option B         LeastC         rem_leastc         2.83704           Genex: Option B         LeastC         rem_leastc         2.994386           Genex: Option B         LeastC         rem_leastc         2.994386           Genex: Option B         LeastC         rem_leastc         17.94971           Genex: Option B         LeastC         rem_leastc         17.04971           Genex: Option B         LeastC         rem_leastc         17.04971           Genex: Option B         LeastC         rem_leastc         17.45408           Genex: Option B         LeastC         rem_leastc         17.84508           Genex: Option B         LeastC         rem_leastc         17.84508           Genex: Option B         LeastC         rem_leastc         129.73636           Genex: Option B         LeastC         rem_leastc         129.949686           Genex: Option B         LeastC         rem_leastc         129.94764           Genex: Option B         LeastC         rem_leastc         169.1414           Genex: Option B         LeastC         rem_leastc         169.1414           Genex: Option B         LeastC         rem_leastc <td></td> <td></td> <td>_</td> <td></td>			_	
Genex: Option B         LeastC         rem_leastc         2.839100           Genex: Option B         LeastC         rem_leastc         2.994388           Genex: Option B         LeastC         rem_leastc         0.739911           Genex: Option B         LeastC         rem_leastc         1.749671           Genex: Option B         LeastC         rem_leastc         1.749671           Genex: Option B         LeastC         rem_leastc         1.3.06972           Genex: Option B         LeastC         rem_leastc         2.8.3588           Genex: Option B         LeastC         rem_leastc         2.8.43588           Genex: Option B         LeastC         rem_leastc         2.8.43588           Genex: Option B         LeastC         rem_leastc         2.8.455183           Genex: Option B         LeastC         rem_leastc         2.2.7.0381           Genex: Option B         LeastC         rem_leastc         2.2.7.0381           Genex: Option B         LeastC         rem_leastc         2.3.4374           Genex: Option B         LeastC         rem_leastc         1.4.9.9945           Genex: Option B         LeastC         rem_leastc         1.4.9.9945           Genex: Option B         LeastC         rem			_	
Genex: Option B         LeastC         rem_leastc         65.74694           Genex: Option B         LeastC         rem_leastc         2.99338           Genex: Option B         LeastC         rem_leastc         2.387905           Genex: Option B         LeastC         rem_leastc         17.46671           Genex: Option B         LeastC         rem_leastc         17.46671           Genex: Option B         LeastC         rem_leastc         17.4512           Genex: Option B         LeastC         rem_leastc         2.84388           Genex: Option B         LeastC         rem_leastc         2.84384           Genex: Option B         LeastC         rem_leastc         2.70387           Genex: Option B         LeastC         rem_leastc         2.270387           Genex: Option B         LeastC         rem_leastc         1.388.612           Genex: Option B         LeastC         rem_leastc         1.2023.065           Genex: Option B         LeastC         rem_leastc         1.691141           Genex: Option B         LeastC         rem_leastc         1.691414           Genex: Option B         LeastC         rem_leastc         1.691414           Genex: Option B         LeastC         rem_leastc			_	
Cenex: Option B         LeastC         rem_leastc         2.99388           Genex: Option B         LeastC         rem_leastc         0.739912           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         70.5411           Genex: Option B         LeastC         rem_leastc         2.84388           Genex: Option B         LeastC         rem_leastc         2.84388           Genex: Option B         LeastC         rem_leastc         2.84384           Genex: Option B         LeastC         rem_leastc         2.84364           Genex: Option B         LeastC         rem_leastc         2.84364           Genex: Option B         LeastC         rem_leastc         1.849988           Genex: Option B         LeastC         rem_leastc         1.849914           Genex: Option B         LeastC         rem_leastc         1.84944           Genex: Option B         LeastC         rem_leastc				
Genex: Option B         LeastC         rem_leastc         0.739912           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         13.06972           Genex: Option B         LeastC         rem_leastc         13.64522           Genex: Option B         LeastC         rem_leastc         13.64523           Genex: Option B         LeastC         rem_leastc         13.64512           Genex: Option B         LeastC         rem_leastc         13.64523           Genex: Option B         LeastC         rem_leastc         14.99886           Genex: Option B         LeastC         rem_leastc         14.99886           Genex: Option B         LeastC         rem_leastc         14.99886           Genex: Option B         LeastC         rem_leastc         14.99154           Genex: Option B         LeastC         rem_leastc         14.99154           Genex: Option B         LeastC         rem_leastc         14.99154           Genex: Option B         LeastC         rem_leastc			_	
Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         17.49671           Genex: Option B         LeastC         rem_leastc         13.06975           Genex: Option B         LeastC         rem_leastc         2.84358           Genex: Option B         LeastC         rem_leastc         2.84358           Genex: Option B         LeastC         rem_leastc         2.84364           Genex: Option B         LeastC         rem_leastc         2.83643           Genex: Option B         LeastC         rem_leastc         2.83643           Genex: Option B         LeastC         rem_leastc         138.6417           Genex: Option B         LeastC         rem_leastc         2.49.986           Genex: Option B         LeastC         rem_leastc         7.4.99154           Genex: Option B         LeastC         rem_leastc         16.94374           Genex: Option B         LeastC         rem_leastc         17.6447           Genex: Option B         LeastC         rem_leastc         17.6447           Genex: Option B         LeastC         rem_leastc         27.68105           Genex: Option B         LeastC         rem_leastc			_	
Genex: Option B         LeastC         rem_leastc         70.5411           Genex: Option B         LeastC         rem_leastc         70.5411           Genex: Option B         LeastC         rem_leastc         72.843588           Genex: Option B         LeastC         rem_leastc         74.543588           Genex: Option B         LeastC         rem_leastc         74.545518           Genex: Option B         LeastC         rem_leastc         72.645518           Genex: Option B         LeastC         rem_leastc         72.70387           Genex: Option B         LeastC         rem_leastc         72.70387           Genex: Option B         LeastC         rem_leastc         74.99155           Genex: Option B         LeastC         rem_leastc         74.99167           Genex: Option B         LeastC         rem_leastc         74.642           Genex: Option B         LeastC         rem_leastc         74.642           Genex: Option B         LeastC         rem_leastc			_	
Genex: Option B         LeastC         rem_leastc         70.5411           Genex: Option B         LeastC         rem_leastc         13.06972           Genex: Option B         LeastC         rem_leastc         24.3588           Genex: Option B         LeastC         rem_leastc         54.55183           Genex: Option B         LeastC         rem_leastc         22.70387           Genex: Option B         LeastC         rem_leastc         23.477           Genex: Option B         LeastC         rem_leastc         14.97988           Genex: Option B         LeastC         rem_leastc         15.68937           Genex: Option B         LeastC         rem_leastc         14.34424           Genex: Option B         LeastC         rem_leastc         10.19417           Genex: Option B         LeastC         rem_leastc         10.6827           Genex: Option B         LeastC         rem_leastc				
Genex: Option B         LeastC         rem_leastc         13.06975           Genex: Option B         LeastC         rem_leastc         2.843586           Genex: Option B         LeastC         rem_leastc         75.54102           Genex: Option B         LeastC         rem_leastc         73.61523           Genex: Option B         LeastC         rem_leastc         22.038647           Genex: Option B         LeastC         rem_leastc         1388.618           Genex: Option B         LeastC         rem_leastc         22.0386           Genex: Option B         LeastC         rem_leastc         23.4377           Genex: Option B         LeastC         rem_leastc         74.99156           Genex: Option B         LeastC         rem_leastc         16.91414           Genex: Option B         LeastC         rem_leastc         12.893.4297           Genex: Option B         LeastC         rem_leastc         12.893.4297           Genex: Option B         LeastC         rem_leastc         12.84742           Genex: Option B         LeastC         rem_leastc         12.648936           Genex: Option B         LeastC         rem_leastc         12.648936           Genex: Option B         LeastC         rem_least				
Genex: Option B         LeastC         rem_leastc         2.843588           Genex: Option B         LeastC         rem_leastc         47.54403           Genex: Option B         LeastC         rem_leastc         73.61523           Genex: Option B         LeastC         rem_leastc         22.70387           Genex: Option B         LeastC         rem_leastc         22.70387           Genex: Option B         LeastC         rem_leastc         149.9886           Genex: Option B         LeastC         rem_leastc         23.437           Genex: Option B         LeastC         rem_leastc         24.99154           Genex: Option B         LeastC         rem_leastc         16.91414           Genex: Option B         LeastC         rem_leastc         16.91414           Genex: Option B         LeastC         rem_leastc         14.99154           Genex: Option B         LeastC         rem_leastc         13.04297           Genex: Option B         LeastC         rem_leastc         14.93442           Genex: Option B         LeastC         rem_leastc         7.76444           Genex: Option B         LeastC         rem_leastc         10.9944           Genex: Option B         LeastC         rem_leastc			1	
Genex:         Option B         LeastC         rem_leastc         47.54403           Genex:         Option B         LeastC         rem_leastc         54.55183           Genex:         Option B         LeastC         rem_leastc         20.83647           Genex:         Option B         LeastC         rem_leastc         22.70387           Genex:         Option B         LeastC         rem_leastc         1388.616           Genex:         Option B         LeastC         rem_leastc         22.3063           Genex:         Option B         LeastC         rem_leastc         149.9986           Genex:         Option B         LeastC         rem_leastc         149.9196           Genex:         Option B         LeastC         rem_leastc         149.9196           Genex:         Option B         LeastC         rem_leastc         139.4297           Genex:         Option B         LeastC         rem_leastc         149.444           Genex:         Option B         LeastC         rem_leastc         149.444           Genex:         Option B         LeastC         rem_leastc         101.9941           Genex:         Option B         LeastC         rem_leastc         101.9941				
Genex:         Option B         LeastC         rem_leastc         73.61523           Genex:         Option B         LeastC         rem_leastc         20.83647           Genex:         Option B         LeastC         rem_leastc         22.70386           Genex:         Option B         LeastC         rem_leastc         1398.618           Genex:         Option B         LeastC         rem_leastc         22.70367           Genex:         Option B         LeastC         rem_leastc         22.8347           Genex:         Option B         LeastC         rem_leastc         74.99155           Genex:         Option B         LeastC         rem_leastc         74.99156           Genex:         Option B         LeastC         rem_leastc         16.91414           Genex:         Option B         LeastC         rem_leastc         13.89442           Genex:         Option B         LeastC         rem_leastc         13.4422           Genex:         Option B         LeastC         rem_leastc         7.76444           Genex:         Option B         LeastC         rem_leastc         50.0022           Genex:         Option B         LeastC         rem_leastc         50.82704				
Genex: Option B         LeastC         rem_leastc         73.61523           Genex: Option B         LeastC         rem_leastc         22.70387           Genex: Option B         LeastC         rem_leastc         123.86.41           Genex: Option B         LeastC         rem_leastc         149.9886           Genex: Option B         LeastC         rem_leastc         22.70387           Genex: Option B         LeastC         rem_leastc         20.3063           Genex: Option B         LeastC         rem_leastc         74.99154           Genex: Option B         LeastC         rem_leastc         16.91141           Genex: Option B         LeastC         rem_leastc         18.94742           Genex: Option B         LeastC         rem_leastc         18.47442           Genex: Option B         LeastC         rem_leastc         18.47442           Genex: Option B         LeastC         rem_leastc         2.66187           Genex: Option B         LeastC         rem_leastc         2.66187           Genex: Option B         LeastC         rem_leastc         2.66187           Genex: Option B         LeastC         rem_leastc         27.2852           Genex: Option B         LeastC         rem_leastc				
Genex:         Option B         LeastC         rem_leastc         20.83647           Genex:         Option B         LeastC         rem_leastc         1388.616           Genex:         Option B         LeastC         rem_leastc         1388.616           Genex:         Option B         LeastC         rem_leastc         149.988           Genex:         Option B         LeastC         rem_leastc         23.337           Genex:         Option B         LeastC         rem_leastc         16.91414           Genex:         Option B         LeastC         rem_leastc         16.91414           Genex:         Option B         LeastC         rem_leastc         128.9936           Genex:         Option B         LeastC         rem_leastc         138.4422           Genex:         Option B         LeastC         rem_leastc         14.3442           Genex:         Option B         LeastC         rem_leastc         2.66187           Genex:         Option B         LeastC         rem_leastc         33.04297           Genex:         Option B         LeastC         rem_leastc         3.05095           Genex:         Option B         LeastC         rem_leastc         3.04274				
Genex: Option B         LeastC         rem_leastc         122.70387           Genex: Option B         LeastC         rem_leastc         1388.61E           Genex: Option B         LeastC         rem_leastc         2023.065           Genex: Option B         LeastC         rem_leastc         2023.065           Genex: Option B         LeastC         rem_leastc         23.4374           Genex: Option B         LeastC         rem_leastc         74.9915           Genex: Option B         LeastC         rem_leastc         93.94763           Genex: Option B         LeastC         rem_leastc         126.8933           Genex: Option B         LeastC         rem_leastc         130.4297           Genex: Option B         LeastC         rem_leastc         7.6444           Genex: Option B         LeastC         rem_leastc         18.47445           Genex: Option B         LeastC         rem_leastc         10.19941           Genex: Option B         LeastC         rem_leastc         10.9941           Genex: Option B         LeastC         rem_leastc         50.08270           Genex: Option B         LeastC         rem_leastc         10.72.255           Genex: Option B         LeastC         rem_leastc				
Genex: Option B       LeastC       rem_leastc       1388.616         Genex: Option B       LeastC       rem_leastc       149.9886         Genex: Option B       LeastC       rem_leastc       223.063         Genex: Option B       LeastC       rem_leastc       74.99154         Genex: Option B       LeastC       rem_leastc       74.99154         Genex: Option B       LeastC       rem_leastc       126.8933         Genex: Option B       LeastC       rem_leastc       138.4744         Genex: Option B       LeastC       rem_leastc       138.4744         Genex: Option B       LeastC       rem_leastc       17.76446         Genex: Option B       LeastC       rem_leastc       10.9941         Genex: Option B       LeastC       rem_leastc       10.9941         Genex: Option B       LeastC       rem_leastc       10.9941         Genex: Option B       LeastC       rem_leastc       500.0821         Genex: Option B       LeastC       rem_leastc       123.2225         Genex: Option B       LeastC       rem_leastc       123.2225         Genex: Option B       LeastC       rem_leastc       123.2225         Genex: Option B       LeastC       rem_leastc <td></td> <td></td> <td>-</td> <td></td>			-	
Genex: Option B       LeastC       rem_leastc       149.9886         Genex: Option B       LeastC       rem_leastc       2023.063         Genex: Option B       LeastC       rem_leastc       74.99155         Genex: Option B       LeastC       rem_leastc       74.99156         Genex: Option B       LeastC       rem_leastc       74.99156         Genex: Option B       LeastC       rem_leastc       93.94765         Genex: Option B       LeastC       rem_leastc       138.4745         Genex: Option B       LeastC       rem_leastc       138.47445         Genex: Option B       LeastC       rem_leastc       7.6444         Genex: Option B       LeastC       rem_leastc       2.66187         Genex: Option B       LeastC       rem_leastc       30.9957         Genex: Option B       LeastC       rem_leastc       500.08270         Genex: Option B       LeastC       rem_leastc       500.8270         Genex: Option B       LeastC       rem_leastc       12.72855         Genex: Option B       LeastC       rem_leastc       12.72855         Genex: Option B       LeastC       rem_leastc       13.76944         Genex: Option B       LeastC       rem_leastc </td <td></td> <td></td> <td>_</td> <td></td>			_	
Genex: Option B       LeastC       rem_leastc       2023.063         Genex: Option B       LeastC       rem_leastc       23.4374         Genex: Option B       LeastC       rem_leastc       74.9915         Genex: Option B       LeastC       rem_leastc       93.94763         Genex: Option B       LeastC       rem_leastc       93.94763         Genex: Option B       LeastC       rem_leastc       126.8933         Genex: Option B       LeastC       rem_leastc       18.47445         Genex: Option B       LeastC       rem_leastc       7.76444         Genex: Option B       LeastC       rem_leastc       2.66187         Genex: Option B       LeastC       rem_leastc       200.827         Genex: Option B       LeastC       rem_leastc       202.222         Genex: Option B       LeastC       rem_leastc       202.222         Genex: Option B       LeastC       rem_leastc       123.2222         Genex: Option B       LeastC       rem_leastc			_	
Genex: Option B       LeastC       rem_leastc       23.4374         Genex: Option B       LeastC       rem_leastc       74.99154         Genex: Option B       LeastC       rem_leastc       16.91414         Genex: Option B       LeastC       rem_leastc       126.8933         Genex: Option B       LeastC       rem_leastc       130.4297         Genex: Option B       LeastC       rem_leastc       148.47444         Genex: Option B       LeastC       rem_leastc       14.334424         Genex: Option B       LeastC       rem_leastc       7.76446         Genex: Option B       LeastC       rem_leastc       2.66187         Genex: Option B       LeastC       rem_leastc       3.50956         Genex: Option B       LeastC       rem_leastc       500.0827         Genex: Option B       LeastC       rem_leastc       500.82704         Genex: Option B       LeastC       rem_leastc       127.8827         Genex: Option B       LeastC       rem_leastc       127.8222         Genex: Option B       LeastC       rem_leastc       127.8222         Genex: Option B       LeastC       rem_leastc       147.844         Genex: Option B       LeastC       rem_leastc </td <td></td> <td></td> <td></td> <td></td>				
Genex: Option B       LeastC       rem_leastc       74.99154         Genex: Option B       LeastC       rem_leastc       16.91414         Genex: Option B       LeastC       rem_leastc       126.8936         Genex: Option B       LeastC       rem_leastc       130.4297         Genex: Option B       LeastC       rem_leastc       18.47445         Genex: Option B       LeastC       rem_leastc       4.334424         Genex: Option B       LeastC       rem_leastc       7.76444         Genex: Option B       LeastC       rem_leastc       2.66187         Genex: Option B       LeastC       rem_leastc       101.9941         Genex: Option B       LeastC       rem_leastc       350955         Genex: Option B       LeastC       rem_leastc       27.68106         Genex: Option B       LeastC       rem_leastc       27.28552         Genex: Option B       LeastC       rem_leastc       103.27226         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc				
Genex: Option B       LeastC       rem_leastc       16.91414         Genex: Option B       LeastC       rem_leastc       93.94765         Genex: Option B       LeastC       rem_leastc       126.8936         Genex: Option B       LeastC       rem_leastc       13.04297         Genex: Option B       LeastC       rem_leastc       4.334424         Genex: Option B       LeastC       rem_leastc       4.334424         Genex: Option B       LeastC       rem_leastc       2.66187         Genex: Option B       LeastC       rem_leastc       101.9941         Genex: Option B       LeastC       rem_leastc       500.0827         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       123.2226         Genex: Option B       LeastC       rem_leastc       123.2226         Genex: Option B       LeastC       rem_leastc       147.6445         Genex: Option B       LeastC       rem_leastc       147.6446         Genex: Option B       LeastC       rem_leastc       147.2455         Genex: Option B       LeastC       rem_leastc<			_	
Genex: Option B       LeastC       rem_leastc       93.94763         Genex: Option B       LeastC       rem_leastc       126.8933         Genex: Option B       LeastC       rem_leastc       184.7445         Genex: Option B       LeastC       rem_leastc       4.334424         Genex: Option B       LeastC       rem_leastc       7.76446         Genex: Option B       LeastC       rem_leastc       2.66137         Genex: Option B       LeastC       rem_leastc       101.9947         Genex: Option B       LeastC       rem_leastc       500.0821         Genex: Option B       LeastC       rem_leastc       500.82704         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       123.222         Genex: Option B       LeastC       rem_leastc       137.262         Genex: Option B       LeastC       rem_leastc       147.8944         Genex: Option B       LeastC       rem_leastc       191.8055         Genex: Option B       LeastC       rem_leastc       191.8055         Genex: Option B       LeastC       rem_leastc <td></td> <td></td> <td>_</td> <td></td>			_	
Genex: Option B       LeastC       rem_leastc       126.8936         Genex: Option B       LeastC       rem_leastc       330.4297         Genex: Option B       LeastC       rem_leastc       18.47445         Genex: Option B       LeastC       rem_leastc       4.334424         Genex: Option B       LeastC       rem_leastc       7.76446         Genex: Option B       LeastC       rem_leastc       101.9941         Genex: Option B       LeastC       rem_leastc       101.9941         Genex: Option B       LeastC       rem_leastc       500.0821         Genex: Option B       LeastC       rem_leastc       500.82704         Genex: Option B       LeastC       rem_leastc       123.2225         Genex: Option B       LeastC       rem_leastc       123.2225         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc       191.8055         Genex: Option B       LeastC       rem_leastc <td></td> <td></td> <td>_</td> <td></td>			_	
Genex: Option B       LeastC       rem_leastc       330.4297         Genex: Option B       LeastC       rem_leastc       18.47445         Genex: Option B       LeastC       rem_leastc       7.76446         Genex: Option B       LeastC       rem_leastc       7.76446         Genex: Option B       LeastC       rem_leastc       101.9941         Genex: Option B       LeastC       rem_leastc       330.6956         Genex: Option B       LeastC       rem_leastc       500.0821         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       50.82704         Genex: Option B       LeastC       rem_leastc       123.2226         Genex: Option B       LeastC       rem_leastc       34.76944         Genex: Option B       LeastC       rem_leastc       32.86117         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc				
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Genex: Option BLeastCrem_leastc50.82704Genex: Option BLeastCrem_leastc27.28552Genex: Option BLeastCrem_leastc123.2226Genex: Option BLeastCrem_leastc367.1263Genex: Option BLeastCrem_leastc34.76944Genex: Option BLeastCrem_leastc3.286117Genex: Option BLeastCrem_leastc107.844Genex: Option BLeastCrem_leastc107.844Genex: Option BLeastCrem_leastc107.844Genex: Option BLeastCrem_leastc180.466Genex: Option BLeastCrem_leastc180.466Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc12.90265Genex: Option B<				
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Genex: Option B       LeastC       rem_leastc       34.76944         Genex: Option B       LeastC       rem_leastc       3.286117         Genex: Option B       LeastC       rem_leastc       107.844         Genex: Option B       LeastC       rem_leastc       191.8055         Genex: Option B       LeastC       rem_leastc       180.466         Genex: Option B       LeastC       rem_leastc       180.466         Genex: Option B       LeastC       rem_leastc       1.27862         Genex: Option B       LeastC       rem_leastc       0.952919         Genex: Option B       LeastC       rem_leastc       0.952919         Genex: Option B       LeastC       rem_leastc       0.324657         Genex: Option B       LeastC       rem_leastc       0.324657         Genex: Option B       LeastC       rem_leastc       12.90265         Genex: Option B       LeastC       rem_leastc       12.90265         Genex: Option B       LeastC       rem_leastc       12.90265         Genex: Option B       LeastC       rem_leastc       34.31715         Genex: Option B       LeastC       rem_leastc       34.31715         Genex: Option B       LeastC       rem_leastc <td></td> <td>LeastC</td> <td></td> <td></td>		LeastC		
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Genex: Option BLeastCrem_leastc191.8055Genex: Option BLeastCrem_leastc80.41676Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc2.4.74313Genex: Option BLeastCrem_leastc2.4.74313Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc1.2.90265Genex: Option BLeastCrem_leastc3.24657Genex: Option BLeastCrem_leastc3.4.31719Genex: Option BLeastCrem_leastc1.2.0026587Genex: Option BLeastCrem_leastc1.2.06587Genex: Option BLeastCrem_leastc3.785716Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc1.70.6071Genex: Option BLeastCrem_leastc1.70.6071Genex: Option BLeastCrem_leastc1.70.6071Genex: Option BLeastCrem_leastc1.76309Genex: Option BLeastCrem_leastc1.76309Genex: Option BLeastCrem_leastc1.641911Genex: Option BLeastCrem_leastc1.342659Genex: Option BLeastCrem_leastc1.344265Genex: Option BLeastCrem_leastc1.785633Genex: Option BLeastCrem_leastc1.58381Genex: Op			rem_leastc	3.286117
Genex: Option BLeastCrem_leastc80.41676Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc12.90265Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31717Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785716Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option B<	Genex: Option B	LeastC	rem_leastc	107.844
Genex: Option BLeastCrem_leastc180.466Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc12.90265Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31717Genex: Option BLeastCrem_leastc34.31717Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc13.49255Genex: Option BLeastCrem_leastc8.19716Genex: Option BLeastCrem_leastc8.19716Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option B <td>Genex: Option B</td> <td>LeastC</td> <td>rem_leastc</td> <td>191.8055</td>	Genex: Option B	LeastC	rem_leastc	191.8055
Genex: Option BLeastCrem_leastc1.27862Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc12.90265Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31719Genex: Option BLeastCrem_leastc4.292203Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option B	Genex: Option B	LeastC	rem_leastc	80.41676
Genex: Option BLeastCrem_leastc24.74313Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc12.90265Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31716Genex: Option BLeastCrem_leastc34.31717Genex: Option BLeastCrem_leastc4.292203Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc37.85718Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc8.19718Genex: Option BLeastCrem_leastc3.19718Genex: Option BLeastCrem_leastc3.19718Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc5.069492Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option B </td <td></td> <td></td> <td></td> <td>180.466</td>				180.466
Genex: Option BLeastCrem_leastc0.952919Genex: Option BLeastCrem_leastc12.90265Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31716Genex: Option BLeastCrem_leastc34.31716Genex: Option BLeastCrem_leastc12.00587Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc37.85718Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc8.153098Genex: Option BLeastCrem_leastc8.319718Genex: Option BLeastCrem_leastc3.4255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc5.069492Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc15.8383Genex: Option B <td></td> <td>LeastC</td> <td>rem_leastc</td> <td>1.27862</td>		LeastC	rem_leastc	1.27862
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Genex: Option BLeastCrem_leastc0.324657Genex: Option BLeastCrem_leastc34.31719Genex: Option BLeastCrem_leastc34.31719Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785718Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc8.153096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option B			rem_leastc	0.952919
Genex: Option BLeastCrem_leastc34.31719Genex: Option BLeastCrem_leastc4.292203Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785718Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc81.53096Genex: Option BLeastCrem_leastc8.319718Genex: Option BLeastCrem_leastc8.319718Genex: Option BLeastCrem_leastc8.319718Genex: Option BLeastCrem_leastc15.44255Genex: Option BLeastCrem_leastc17.442563Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc59.0305Genex: Option BLeastCrem_leastc59.0305Genex: Option B		LeastC	rem_leastc	12.90265
Genex: Option BLeastCrem_leastc4.292203Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785718Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc44.47043Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc81.53096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc2.661891Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.86432Genex: Option BLeastCrem_leastc17.85632Genex: Option BLeastCrem_leastc17.85632Genex: Option BLeastCrem_leastc17.85632Genex: Option BLeastCrem_leastc17.85632Genex: Option BLeastCrem_leastc13.39533Genex: Option BLeastCrem_leastc13.39533Genex: Option BLeastCrem_leastc13.39533Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc59.0303Genex: Option B <td></td> <td>LeastC</td> <td></td> <td>0.324657</td>		LeastC		0.324657
Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785716Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc44.47043Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc8.153096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc2.661891Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc15.8383Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc59.0305Genex: Option BLeastCrem_leastc59.0305Genex: Option BLeastCrem_leastc50.63732Genex: Option BLeastCrem_leastc50.63732Genex: Option B <td></td> <td>LeastC</td> <td>rem_leastc</td> <td>34.31719</td>		LeastC	rem_leastc	34.31719
Genex: Option BLeastCrem_leastc12.06587Genex: Option BLeastCrem_leastc3.785716Genex: Option BLeastCrem_leastc0.559217Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc170.6071Genex: Option BLeastCrem_leastc44.47043Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc8.153096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc2.661891Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc17.85637Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc15.8383Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc13.39535Genex: Option BLeastCrem_leastc59.0305Genex: Option BLeastCrem_leastc59.0305Genex: Option BLeastCrem_leastc50.63732Genex: Option BLeastCrem_leastc50.63732Genex: Option B <td>Genex: Option B</td> <td>LeastC</td> <td>rem_leastc</td> <td>4.292203</td>	Genex: Option B	LeastC	rem_leastc	4.292203
Genex: Option B       LeastC       rem_leastc       0.559217         Genex: Option B       LeastC       rem_leastc       170.6071         Genex: Option B       LeastC       rem_leastc       44.47043         Genex: Option B       LeastC       rem_leastc       2.756305         Genex: Option B       LeastC       rem_leastc       81.53096         Genex: Option B       LeastC       rem_leastc       8.319716         Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       15.8381         Genex: Option B       LeastC       rem_leastc       13.39533         Genex: Option B       LeastC       rem_leastc       13.39533         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc </td <td></td> <td>LeastC</td> <td>rem_leastc</td> <td>12.06587</td>		LeastC	rem_leastc	12.06587
Genex: Option B       LeastC       rem_leastc       170.6071         Genex: Option B       LeastC       rem_leastc       44.47043         Genex: Option B       LeastC       rem_leastc       2.756305         Genex: Option B       LeastC       rem_leastc       81.53096         Genex: Option B       LeastC       rem_leastc       8.319716         Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       5.069492         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       13.39533         Genex: Option B       LeastC       rem_leastc       13.39533         Genex: Option B       LeastC       rem_leastc       13.39533         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc<	Genex: Option B	LeastC	rem_leastc	3.785718
Genex: Option BLeastCrem_leastc44.47043Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc81.53096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc2.661891Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc13.9533Genex: Option BLeastCrem_leastc13.9533Genex: Option BLeastCrem_leastc59.0300Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc60.63732Genex: Option B		LeastC	rem_leastc	0.559217
Genex: Option BLeastCrem_leastc44.47043Genex: Option BLeastCrem_leastc2.756305Genex: Option BLeastCrem_leastc81.53096Genex: Option BLeastCrem_leastc8.319716Genex: Option BLeastCrem_leastc2.661891Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc13.44255Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc17.85633Genex: Option BLeastCrem_leastc15.8381Genex: Option BLeastCrem_leastc13.9533Genex: Option BLeastCrem_leastc13.9533Genex: Option BLeastCrem_leastc59.0300Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc59.0303Genex: Option BLeastCrem_leastc60.63732Genex: Option B	Genex: Option B	LeastC	rem_leastc	170.6071
Genex: Option B       LeastC       rem_leastc       81.53096         Genex: Option B       LeastC       rem_leastc       8.319716         Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       5.069492         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       13.39535         Genex: Option B       LeastC       rem_leastc       13.39535         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       60.63732         Genex: Option B       LeastC       rem_leastc       60.63732		LeastC	rem_leastc	44.47043
Genex: Option B       LeastC       rem_leastc       81.53096         Genex: Option B       LeastC       rem_leastc       8.319716         Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       5.069492         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       13.39535         Genex: Option B       LeastC       rem_leastc       13.39535         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       59.0300         Genex: Option B       LeastC       rem_leastc       60.63732         Genex: Option B       LeastC       rem_leastc       60.63732	Genex: Option B	LeastC	rem_leastc	2.756305
Genex: Option B       LeastC       rem_leastc       8.319718         Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       13.64255         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       15.8381         Genex: Option B       LeastC       rem_leastc       13.39539         Genex: Option B       LeastC       rem_leastc       13.39539         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       60.63732			_	81.53098
Genex: Option B       LeastC       rem_leastc       2.661891         Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       5.069492         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       15.8381         Genex: Option B       LeastC       rem_leastc       13.39539         Genex: Option B       LeastC       rem_leastc       13.39539         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       60.63732			_	8.319718
Genex: Option B       LeastC       rem_leastc       13.44255         Genex: Option B       LeastC       rem_leastc       5.069492         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       17.85633         Genex: Option B       LeastC       rem_leastc       15.8381         Genex: Option B       LeastC       rem_leastc       13.39539         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       59.0309         Genex: Option B       LeastC       rem_leastc       60.63732			_	2.661891
Genex: Option B     LeastC     rem_leastc     5.069492       Genex: Option B     LeastC     rem_leastc     17.85633       Genex: Option B     LeastC     rem_leastc     15.8381       Genex: Option B     LeastC     rem_leastc     13.39539       Genex: Option B     LeastC     rem_leastc     13.39539       Genex: Option B     LeastC     rem_leastc     59.0309       Genex: Option B     LeastC     rem_leastc     59.0309       Genex: Option B     LeastC     rem_leastc     60.63732			_	13.44255
Genex: Option B         LeastC         rem_leastc         17.85633           Genex: Option B         LeastC         rem_leastc         15.8381           Genex: Option B         LeastC         rem_leastc         13.39539           Genex: Option B         LeastC         rem_leastc         13.39539           Genex: Option B         LeastC         rem_leastc         59.0309           Genex: Option B         LeastC         rem_leastc         59.0309           Genex: Option B         LeastC         rem_leastc         60.63732			_	5.069492
Genex: Option B         LeastC         rem_leastc         15.8381           Genex: Option B         LeastC         rem_leastc         13.39539           Genex: Option B         LeastC         rem_leastc         59.0309           Genex: Option B         LeastC         rem_leastc         59.0309           Genex: Option B         LeastC         rem_leastc         60.63732	Genex: Option B		_	
Genex: Option B         LeastC         rem_leastc         13.39539           Genex: Option B         LeastC         rem_leastc         59.0309           Genex: Option B         LeastC         rem_leastc         60.63732			rem_leastc	17.80033
Genex: Option B         LeastC         rem_leastc         59.0305           Genex: Option B         LeastC         rem_leastc         60.63732	Genex: Option B	LeastC	_	15.8381
Genex: Option B LeastC rem_leastc 60.63732	Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc	
Conovi Option P Locat C rore laste 405.000	Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC	rem_leastc rem_leastc	15.8381 13.39539
Genex: Option B LeastC rem_leastc 135.982	Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	15.8381

Genex: Option B	LeastC	rem_leastc	26.17916
Genex: Option B	LeastC	rem leastc	12.59796
Genex: Option B	LeastC	rem leastc	29.47318
Genex: Option B	LeastC	rem_leastc	137.9151
Genex: Option B	LeastC	rem leastc	109.8415
Genex: Option B	LeastC	rem leastc	149.8079
Genex: Option B	LeastC	rem leastc	462.3725
Genex: Option B	LeastC	rem_leastc	210.8056
Genex: Option B	LeastC	rem leastc	96.2025
Genex: Option B	LeastC	rem leastc	11.50008
Genex: Option B	LeastC	rem_leastc	10.76424
Genex: Option B	LeastC	rem_leastc	183.6566
Genex: Option B	LeastC	rem_leastc	17.68986
Genex: Option B	LeastC	rem leastc	190.5614
	LeastC		190.5614
Genex: Option B		rem_leastc	59.1258
Genex: Option B	LeastC	rem_leastc	
Genex: Option B	LeastC	rem_leastc	8.331762
Genex: Option B	LeastC	rem_leastc	13.08896
Genex: Option B	LeastC	rem_leastc	2.857761
Genex: Option B	LeastC	rem_leastc	8.412393
Genex: Option B	LeastC	rem_leastc	612.04
Genex: Option B	LeastC	rem_leastc	604.5275
Genex: Option B	LeastC	rem_leastc	9.989402
Genex: Option B	LeastC	rem_leastc	26.86946
Genex: Option B	LeastC	rem_leastc	1.759112
Genex: Option B	LeastC	rem_leastc	129.2239
Genex: Option B	LeastC	rem_leastc	13.45173
Genex: Option B	LeastC	rem_leastc	2.664493
Genex: Option B	LeastC	rem_leastc	58.33691
Genex: Option B	LeastC	rem_leastc	17.44888
Genex: Option B	LeastC	rem_leastc	7.161197
Genex: Option B	LeastC	rem_leastc	348.5125
Genex: Option B	LeastC	rem_leastc	9.412106
Genex: Option B	LeastC	rem_leastc	1.264701
Genex: Option B	LeastC	rem_leastc	553.999
Genex: Option B	LeastC	rem_leastc	192.4199
Genex: Option B	LeastC	rem_leastc	9.22467
Genex: Option B	LeastC	rem_leastc	7.926539
Genex: Option B	LeastC	rem_leastc	29.46511
Genex: Option B	LeastC	rem_leastc	324.4124
Genex: Option B	LeastC	rem_leastc	4.364596
Genex: Option B	LeastC	rem_leastc	248.962
Genex: Option B	LeastC	rem_leastc	10.31886
Genex: Option B	LeastC	rem_leastc	215.5128
Genex: Option B	LeastC	rem_leastc	64.58989
Genex: Option B	LeastC	rem_leastc	25.19911
Genex: Option B	LeastC	rem_leastc	404.6123
Genex: Option B	LeastC	rem_leastc	0.053489
Genex: Option B	LeastC	rem_leastc	0.589376
Genex: Option B	LeastC	rem_leastc	25 02660
Genex: Option B	LeastC	rom loosto	25.03669
		rem_leastc	0.34877
Genex: Option B	LeastC	rem_leastc	0.34877 120.4981
Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618
Genex: Option B Genex: Option B	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669
Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615
Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029
Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289
Genex: Option B Genex: Option B	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024 63.09774
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024 63.09774
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024 63.09774 1.551911
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024 63.09774 1.551911 6.945101
Genex: Option B Genex: Option B	LeastC LeastC	rem_leastc rem_leastc	0.34877 120.4981 200.9618 192.5669 22.14615 17.90449 37.14543 100.6029 40.38307 16.41434 9.77248 32.21486 46.3546 0.008122 44.16289 5.923245 3.490973 131.4287 8.152129 0.000275 0.000024 63.09774 1.551911 6.945101 0.000132

	1 10		15 00015	
Genex: Option B	LeastC	rem_leastc	15.80917	
Genex: Option B	LeastC	rem_leastc	192.9281	
Genex: Option B	LeastC	rem_leastc	0.037976	
Genex: Option B	LeastC	rem_leastc	49.50354	
Genex: Option B	LeastC	rem_leastc	125.1382	
Genex: Option B	LeastC	rem_leastc	20.527	
Genex: Option B	LeastC	rem_leastc	2.774712	
Genex: Option B	LeastC	rem_leastc	102.0444	
Genex: Option B	LeastC	rem_leastc	4.059098	
Genex: Option B	LeastC	rem_leastc	0.453452	
Genex: Option B	LeastC	rem_leastc	12.2734	
Genex: Option B	LeastC	rem_leastc	8.720471	
Genex: Option B	LeastC	rem_leastc	9.605091	
Genex: Option B	LeastC	rem_leastc	0.326735	
Genex: Option B	LeastC	rem_leastc	0.727728	
Genex: Option B	LeastC	rem_leastc	0.288437	
Genex: Option B	LeastC	rem_leastc	13.82986	
Genex: Option B	LeastC	rem_leastc	57.22385	
Genex: Option B	LeastC	rem_leastc	4.842688	
Genex: Option B	LeastC	rem_leastc	32.01963	
Genex: Option B	LeastC	rem_leastc	0.78514	
Genex: Option B	LeastC	rem_leastc	31.26611	
Genex: Option B	LeastC	rem_leastc	0.826022	
Genex: Option B	LeastC	rem_leastc	0.584156	
Genex: Option B	LeastC	rem_leastc	0.600281	
Genex: Option B	LeastC	rem_leastc	6.922604	
Genex: Option B	LeastC	rem_leastc	51.43218	
Genex: Option B	LeastC	rem_leastc	133.6165	
Genex: Option B	LeastC	rem_leastc	10.23287	
Genex: Option B	LeastC	rem_leastc	0.001706	
Genex: Option B	O-dom	rem_oc	1.471846	1891 Total OC-dom remnant for Option B
Genex: Option B	O-dom	rem_oc	195.4305	
Genex: Option B	O-dom	rem_oc	153.7515	
Genex: Option B	O-dom	rem_oc	138.478	
Genex: Option B	O-dom	rem_oc	10.5891	
Genex: Option B	O-dom	rem_oc	5.651625	
Genex: Option B	O-dom	rem_oc	0.037097	
Genex: Option B	O-dom	rem_oc	15.35427	
Genex: Option B	O-dom	rem_oc	5.626584	
Genex: Option B	O-dom	rem_oc	0.099917	
Genex: Option B	O-dom	rem_oc	1.166769	
Genex: Option B	O-dom	rem_oc	270.0271	
Genex: Option B	0-dom	rem oc	374.3618	

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Genex: Option B	O-dom	rem_oc	27.99034
Genex: Option B	O-dom	rem_oc	47.06895
Genex: Option B	O-dom	rem_oc	1.946463
Genex: Option B	O-dom	rem_oc	1.771366
Genex: Option B	O-dom	rem_oc	108.0733
Genex: Option B	O-dom	rem_oc	0.590693
Genex: Option B	O-dom	rem_oc	26.872
Genex: Option B	O-dom	rem_oc	0.925363
Genex: Option B	O-dom	rem_oc	0.182819
Genex: Option B	O-dom	rem_oc	0.144678
Genex: Option B	O-dom	rem_oc	9.653946
Genex: Option B	O-dom	rem_oc	0.001118
Genex: Option B	O-dom	rem_oc	5.780518
Genex: Option B	O-dom		2.728088
		rem_oc	
Genex: Option B	O-dom	rem_oc	2.647043
Genex: Option B	O-dom	rem_oc	1.321795
Genex: Option B	O-dom	rem_oc	1.075075
Genex: Option B	O-dom	rem_oc	1.028013
Genex: Option B	O-dom	rem_oc	4.05067
Genex: Option B	O-dom	rem_oc	9.522925
Genex: Option B	O-dom	rem_oc	0.001392
Genex: Option B	O-dom	rem_oc	0.003394
Genex: Option B	O-dom	rem_oc	0.001035
Genex: Option B	0-dom	rem_oc	33.69548
Genex: Option B	O-dom	rem_oc	1.381382
Genex: Option B	O-dom	rem_oc	0.928588
Genex: Option B	O-dom		0.928588
		rem_oc	3.286292
Genex: Option B	O-dom	rem_oc	
Genex: Option B	O-dom	rem_oc	0.813393
Genex: Option B	O-dom	rem_oc	4.770736
Genex: Option B	O-dom	rem_oc	1.561019
Genex: Option B	O-dom	rem_oc	2.309776
Genex: Option B	O-dom	rem_oc	1.058237
Genex: Option B	O-dom	rem_oc	5.369112
Genex: Option B	O-dom	rem_oc	2.128498
Genex: Option B	O-dom	rem_oc	0.502309
Genex: Option B	O-dom	rem_oc	0.053237
Genex: Option B	O-dom	rem_oc	14.87732
Genex: Option B	O-dom	rem_oc	48.60826
-	O-dom	_	0.43457
Genex: Option B		rem_oc	
Genex: Option B	O-dom	rem_oc	58.61665
Genex: Option B	O-dom	rem_oc	62.13515
Genex: Option B	O-dom	rem_oc	4.286207
Genex: Option B	O-dom	rem_oc	8.433967
Genex: Option B	O-dom	rem_oc	4.418017
Genex: Option B	O-dom	rem_oc	0.816551
Genex: Option B	O-dom	rem_oc	1.360325
Genex: Option B	O-dom	rem_oc	0.61658
Genex: Option B	O-dom	rem_oc	0.594068
Genex: Option B	O-dom	rem_oc	9.111942
Genex: Option B	O-dom	rem_oc	8.567498
			5.563262
Genex: Option B	O-dom	rem_oc	
Genex: Option B	O-dom	rem_oc	58.10314
Genex: Option B	O-dom	rem_oc	11.76356
Genex: Option B	O-dom	rem_oc	15.50552
Genex: Option B	O-dom	rem_oc	4.488126
Genex: Option B	O-dom	rem_oc	3.278022
Genex: Option B	O-dom	rem_oc	3.436123
Genex: Option B	O-dom	rem_oc	11.64185
Genex: Option B	O-dom	rem_oc	18.22205
Genex: Option B	O-dom	rem_oc	0.0018
Genex: Option B	O-dom	rem_oc	4.444007
Genex: Option B	O-dom		6.059981
		rem_oc	
Genex: Option B	O-dom	rem_oc	0.338932
Genex: Option B	O-dom	rem_oc	15.98738
		rem_oc	0.645188
Genex: Option B	O-dom		
Genex: Option B	O-dom	rem_oc	3.072902
Genex: Option B Genex: Option B	O-dom O-dom		3.072902 0.000344
Genex: Option B	O-dom	rem_oc	
Genex: Option B Genex: Option B	O-dom O-dom	rem_oc rem_oc	0.000344

Genex: Option B	O-subdom	rem_oc	28.0519
Genex: Option B	O-subdom	rem_oc	46.6716
Genex: Option B	O-subdom	rem_oc	4.034857
Genex: Option B	O-subdom	rem_oc	187.4794
Genex: Option B	O-subdom	rem_oc	6.946681
Genex: Option B	O-subdom	rem_oc	2.117614
Genex: Option B	O-subdom	rem_oc	59.27586
Genex: Option B	O-subdom	rem_oc	21.51577
Genex: Option B	O-subdom	rem_oc	85.42148

442 Total OC-subdom remnant for Option B

### Kidston Corridor Option C - Regional ecosystems v8

Name	VM_POLY	VM_STATUS	Area_ha	
GENEX Option C	LeastC	hvr leastc	0.042944	266 Total LC regrowth for Option C
GENEX Option C	LeastC	hvr_leastc	1.282108	
GENEX Option C	LeastC	hvr_leastc	3.724701	
GENEX Option C	LeastC	hvr_leastc	3.327094	
GENEX Option C	LeastC	_	0.198184	
		hvr_leastc		
GENEX Option C	LeastC	hvr_leastc	0.215019	
GENEX Option C	LeastC	hvr_leastc	0.871264	
GENEX Option C	LeastC	hvr_leastc	0.001538	
GENEX Option C	LeastC	hvr_leastc	0.330203	
GENEX Option C	LeastC	hvr_leastc	20.73632	
GENEX Option C	LeastC	hvr_leastc	5.521453	
GENEX Option C	LeastC	hvr_leastc	3.584613	
GENEX Option C	LeastC	hvr_leastc	0.039661	
GENEX Option C	LeastC	hvr_leastc	0.247062	
GENEX Option C	LeastC	hvr_leastc	0.779189	
GENEX Option C	LeastC	hvr_leastc	4.748415	
GENEX Option C	LeastC	hvr_leastc	1.859195	
GENEX Option C	LeastC	hvr_leastc	10.84232	
GENEX Option C	LeastC	hvr_leastc	0.563293	
GENEX Option C	LeastC	hvr_leastc	0.799334	
GENEX Option C	LeastC	hvr_leastc	0.1578	
GENEX Option C	LeastC	hvr_leastc	0.423847	
GENEX Option C	LeastC	hvr_leastc	0.022131	
GENEX Option C	LeastC	hvr_leastc	1.832039	
GENEX Option C	LeastC	hvr_leastc	2.074049	
GENEX Option C	LeastC	hvr_leastc	11.76166	
GENEX Option C	LeastC	hvr_leastc	0.071373	
GENEX Option C	LeastC	hvr_leastc	0.032936	
GENEX Option C	LeastC	hvr_leastc	0.275677	
		_	4.030559	
GENEX Option C	LeastC	hvr_leastc		
GENEX Option C	LeastC	hvr_leastc	4.565859	
GENEX Option C	LeastC	hvr_leastc	2.294056	
GENEX Option C	LeastC	hvr_leastc	1.055631	
GENEX Option C	LeastC	hvr_leastc	1.583837	
GENEX Option C	LeastC	hvr_leastc	4.886182	
GENEX Option C	LeastC	hvr_leastc	0.483842	
GENEX Option C	LeastC	hvr_leastc	0.022396	
GENEX Option C	LeastC	hvr_leastc	16.55398	
GENEX Option C	LeastC	hvr_leastc	0.847925	
GENEX Option C	LeastC	hvr_leastc	0.452182	
GENEX Option C	LeastC	hvr_leastc	0.355135	
GENEX Option C	LeastC	hvr_leastc	1.329568	
GENEX Option C	LeastC	hvr_leastc	7.449196	
GENEX Option C	LeastC	hvr_leastc	10.2054	
GENEX Option C	LeastC	hvr_leastc	0.252609	
GENEX Option C	LeastC	hvr_leastc	0.004932	
GENEX Option C	LeastC	hvr_leastc	2.193916	
GENEX Option C	LeastC	hvr_leastc	0.628194	
GENEX Option C	LeastC	hvr_leastc	4.798001	
GENEX Option C	LeastC	hvr_leastc	4.892764	
GENEX Option C	LeastC	hvr_leastc	0.857374	
GENEX Option C	LeastC	hvr_leastc	0.016948	
GENEX Option C	LeastC	hvr_leastc	0.026688	
GENEX Option C	LeastC	hvr_leastc	0.489348	
GENEX Option C	LeastC	hvr leastc	0.001268	
GENEX Option C	LeastC	hvr_leastc	0.733646	
GENEX Option C	LeastC	hvr_leastc	0.001018	
GENEX Option C	LeastC	hvr_leastc	2.958507	
GENEX Option C	LeastC	hvr_leastc	2.596304	
GENEX Option C	LeastC	hvr_leastc	2.274126	
GENEX Option C	LeastC	hvr_leastc	0.860982	
GENEX Option C	LeastC	hvr_leastc	0.17804	
		_		
GENEX Option C	LeastC	hvr_leastc	1.002297	
GENEX Option C	LeastC	hvr_leastc	0.00441	
GENEX Option C	LeastC	hvr_leastc	13.17734	
GENEX Option C	LeastC	hvr_leastc	0.224692	
GENEX Option C	LeastC	hvr_leastc	1.836962	
GENEX Option C	LeastC	hvr_leastc	0.523549	
GENEX Option C	LeastC	hvr_leastc	5.202041	
GENEX Option C	LeastC	hvr_leastc	0.41416	
GENEX Option C	LeastC	hvr_leastc	0.004426	
GENEX Option C	LeastC	hvr_leastc	0.175718	
GENEX Option C	LeastC	hvr_leastc	1.371884	
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GENEX Option C	LeastC	hvr_leastc	0.720414	
GENEX Option C	LeastC	hvr_leastc	0.086502	
GENEX Option C	LeastC	hvr_leastc	0.349904	
GENEX Option C	LeastC	hvr_leastc	9.197092	
GENEX Option C	LeastC	hvr_leastc	2.849699	
GENEX Option C	LeastC	hvr_leastc	1.297257	
GENEX Option C	LeastC	hvr_leastc	63.51923	
GENEX Option C	LeastC	hvr_leastc	0.000024	
GENEX Option C	LeastC	hvr_leastc	0.000132	
GENEX Option C	LeastC	hvr_leastc	0.000386	
GENEX Option C	LeastC	hvr_leastc	2.657052	
GENEX Option C	LeastC	hvr_leastc	1.646678	
GENEX Option C	LeastC	hvr leastc	3.808301	
GENEX Option C	LeastC	hvr_leastc	0.001706	
	0-dom	_		2.4 Total OC dam LIV/D regrowth for Option C
GENEX Option C		hvr_oc	0.888013	2.4 Total OC-dom HVR regrowth for Option C
GENEX Option C	O-dom	hvr_oc	0.034724	
GENEX Option C	O-dom	hvr_oc	0.084695	
GENEX Option C	O-dom	hvr_oc	0.476706	
GENEX Option C	O-dom	hvr_oc	0.890819	
GENEX Option C	non-rem	non_remnant	28.55118	2458 Total non-remnant for Option C
GENEX Option C	non-rem	non_remnant	4.598363	
GENEX Option C	non-rem	non_remnant	3.133693	
GENEX Option C	non-rem	non_remnant	19.4655	
GENEX Option C	non-rem	non_remnant	11.13507	
GENEX Option C	non-rem	non_remnant	3.84902	
GENEX Option C	non-rem	non remnant	1.69844	
GENEX Option C	non-rem	non_remnant	29.78413	
GENEX Option C	non-rem	non_remnant	1.037267	
GENEX Option C	non-rem	non_remnant	631.163	
· · · · · · · · · · · · · · · · · · ·	1	_	1.27489	
GENEX Option C	non-rem	non_remnant		
GENEX Option C	non-rem	non_remnant	1.573922	
GENEX Option C	non-rem	non_remnant	0.416775	
GENEX Option C	non-rem	non_remnant	0.284209	
GENEX Option C	non-rem	non_remnant	0.934327	
GENEX Option C	non-rem	non_remnant	32.73321	
GENEX Option C	non-rem	non_remnant	0.21711	
GENEX Option C	non-rem	non_remnant	7.341556	
GENEX Option C	non-rem	non_remnant	1.536304	
GENEX Option C	non-rem	non_remnant	1.624022	
GENEX Option C	non-rem	non_remnant	1.525838	
GENEX Option C	non-rem	non_remnant	6.35254	
GENEX Option C	non-rem	non_remnant	0.311335	
GENEX Option C	non-rem	non_remnant	0.284912	
GENEX Option C	non-rem	non remnant	0.799373	
GENEX Option C	1	_	1.586069	
· · · · · · · · · · · · · · · · · · ·	non-rem	non_remnant		
GENEX Option C	non-rem	non_remnant	106.2974	
GENEX Option C	non-rem	non_remnant	3.895228	
GENEX Option C	non-rem	non_remnant	288.9389	
GENEX Option C	non-rem	non_remnant	4.258125	
GENEX Option C	non-rem	non_remnant	2.125011	
GENEX Option C	non-rem	non_remnant	0.056622	
GENEX Option C	non-rem	non_remnant	0.167595	
GENEX Option C	non-rem	non_remnant	0.535546	
GENEX Option C	non-rem	non_remnant	0.756154	
GENEX Option C	non-rem	non_remnant	8.966574	
GENEX Option C	non-rem	non_remnant	16.25738	
GENEX Option C	non-rem	non_remnant	347.132	
GENEX Option C	non-rem	non_remnant	0.000275	
GENEX Option C	non-rem	non_remnant	0.000386	
GENEX Option C	non-rem	non_remnant	0.234067	
GENEX Option C	non-rem	non_remnant	282.7614	
GENEX Option C			0.000204	
GENEX Option C	non-rem	non_remnant	0.000204	
· · · · · · · · · · · · · · · · · · ·	non-rem	non_remnant		
GENEX Option C	non-rem	non_remnant	8.914747	
GENEX Option C	non-rem	non_remnant	1.286566	
GENEX Option C	non-rem	non_remnant	1.685735	
GENEX Option C	non-rem	non_remnant	0.526879	
GENEX Option C	non-rem	non_remnant	0.121333	
GENEX Option C	non-rem	non_remnant	3.735213	
GENEX Option C	non-rem	non_remnant	117.347	
			468.6799	
GENEX Option C	non-rem	non_remnant	10010777	
GENEX Option C GENEX Option C	non-rem non-rem	non_remnant	0.001706	
· · · · · · · · · · · · · · · · · · ·	1			29 Total water non-remnant for Option C
GENEX Option C	non-rem	non_remnant	0.001706	29 Total water non-remnant for Option C
GENEX Option C GENEX Option C	non-rem water	non_remnant non_remnant	0.001706 0.341308	29 Total water non-remnant for Option C

GENEX Option C	LeastC	rem_leastc	38.01406	52609 Total LC remnant for Option C
GENEX Option C	LeastC	rem_leastc	0.714087	
GENEX Option C	LeastC	rem_leastc	26.02467	
GENEX Option C	LeastC	rem_leastc	170.1283	
GENEX Option C	LeastC	rem_leastc	51.69668	
GENEX Option C	LeastC	rem_leastc	3.551085	
GENEX Option C	LeastC	rem_leastc	7.754542	
GENEX Option C	LeastC	rem_leastc	108.8235	
GENEX Option C	LeastC	rem_leastc	21.73775	
GENEX Option C	LeastC	rem_leastc	418.7269	
GENEX Option C	LeastC	rem_leastc	57.48674	
GENEX Option C	LeastC	rem_leastc	52.76348	
GENEX Option C	LeastC	rem_leastc	7.20723	
GENEX Option C	LeastC	rem_leastc	4.89243	
GENEX Option C	LeastC	rem_leastc	28.01646	
GENEX Option C	LeastC	rem_leastc	4.747283	
GENEX Option C	LeastC	rem_leastc	39.4278	
GENEX Option C	LeastC	rem_leastc	5.172642	
GENEX Option C	LeastC	rem_leastc	137.9748	
GENEX Option C	LeastC	rem_leastc	1.521492	
GENEX Option C	LeastC	rem_leastc	53.50867	
GENEX Option C	LeastC	rem_leastc	46.62655	
GENEX Option C	LeastC	rem_leastc	30.00713	
GENEX Option C	LeastC	rem_leastc	331.3991	
GENEX Option C	LeastC	rem_leastc	332.4453	
GENEX Option C	LeastC	rem_leastc	21.32897	
GENEX Option C	LeastC	rem_leastc	2.386717	
GENEX Option C	LeastC	rem_leastc	102.9591	
GENEX Option C	LeastC	rem_leastc	145.4984	
GENEX Option C	LeastC	rem_leastc	963.089	
GENEX Option C	LeastC	rem_leastc	76.38114	
GENEX Option C	LeastC	rem_leastc	43.23859	
GENEX Option C	LeastC	rem_leastc	47.90637	
GENEX Option C	LeastC	rem_leastc	262.2519	
GENEX Option C	LeastC	rem_leastc	19.36124	
GENEX Option C	LeastC	rem_leastc	38.22447	
GENEX Option C	LeastC	rem_leastc	67.2146	
GENEX Option C	LeastC	rem_leastc	13.1407	
GENEX Option C	LeastC	rem_leastc	319.7845	
GENEX Option C	LeastC	rem_leastc	1.831239	
GENEX Option C	LeastC	rem_leastc	31.41553	
GENEX Option C	LeastC	rem_leastc	1097.731	
GENEX Option C	LeastC	rem_leastc	45.09985	
GENEX Option C	LeastC	rem_leastc	10.77315	
GENEX Option C	LeastC	rem_leastc	11.55033	
GENEX Option C	LeastC	rem_leastc	378.1576	
GENEX Option C	LeastC	rem_leastc	61.16019	
GENEX Option C	LeastC	rem_leastc	8.994313	
GENEX Option C	LeastC	rem_leastc	306.4878	
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc	20.92654	
		rem_leastc	35.97101	
GENEX Option C	LeastC	rem_leastc	6.257186	
GENEX Option C	LeastC	rem_leastc	94.1651	
GENEX Option C	LeastC	rem_leastc	74.39137	
GENEX Option C	LeastC	rem_leastc	1.175948	
GENEX Option C	LeastC	rem_leastc	8.255441	
GENEX Option C	LeastC	rem_leastc	110.7072	
GENEX Option C	LeastC	rem_leastc	21.48688	
GENEX Option C	LeastC LeastC	rem_leastc	0.579317	
GENEX Option C GENEX Option C	LeastC	rem_leastc rem_leastc	22.56373 2.539484	
GENEX Option C	LeastC	rem_leastc	2.539484	
GENEX Option C	LeastC	rem_leastc	224.1086	
GENEX Option C	LeastC	rem_leastc	14.71189	
GENEX Option C	LeastC	rem_leastc	92.26965	
GENEX Option C	LeastC	rem_leastc	92.26965	
GENEX Option C	LeastC	rem_leastc	339.8278	
GENEX Option C	LeastC	rem_leastc	41.16227	
GENEX Option C	LeastC	rem_leastc	21.93453	
GENEX Option C	LeastC	rem_leastc	1.387434	
GENEX Option C	LeastC	rem_leastc	14.70381	
GENEX Option C	LeastC	rem_leastc	29.1314	
GENEX Option C	LeastC	rem_leastc	16.62314	
GENEX Option C	LeastC	rem_leastc	12.50073	
GENEX Option C	LeastC	rem_leastc	7.261978	
GENEX Option C	LeastC	rem_leastc	18.25935	
GENEX Option C	LeastC	rem_leastc	30.77885	
GENEX Option C	LeastC	rem_leastc	29.59063	

GENEX Option C	LeastC	rem_leastc	3.675156
GENEX Option C	LeastC	rem leastc	6.804901
GENEX Option C	LeastC	rem_leastc	325.6778
GENEX Option C	LeastC	rem_leastc	15.98994
GENEX Option C	LeastC	rem_leastc	257.9476
GENEX Option C	LeastC	rem_leastc	4.391148
GENEX Option C	LeastC	rem leastc	5.114837
GENEX Option C	LeastC	rem leastc	9.306435
GENEX Option C	LeastC	rem leastc	103.875
GENEX Option C	LeastC	rem leastc	306.8621
GENEX Option C	LeastC	rem leastc	13.5309
GENEX Option C	LeastC	rem_leastc	23.87092
GENEX Option C	LeastC	rem_leastc	356.2161
GENEX Option C	LeastC	rem leastc	2.204996
GENEX Option C	LeastC	rem leastc	157.3748
GENEX Option C	LeastC	rem leastc	23.59838
GENEX Option C	LeastC	rem_leastc	10.95056
GENEX Option C	LeastC	rem leastc	2.856698
GENEX Option C	LeastC	rem leastc	0.394979
GENEX Option C	LeastC	rem_leastc	12.14453
GENEX Option C			
GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	83.0943 407.5964
GENEX Option C	LeastC	rem_leastc	287.1895
GENEX Option C	LeastC	rem_leastc	3.004676
GENEX Option C	LeastC	rem_leastc	40.44207
GENEX Option C	LeastC	rem_leastc	12.50636
GENEX Option C	LeastC	rem_leastc	28.19659
GENEX Option C	LeastC	rem_leastc	55.5168
GENEX Option C	LeastC	rem_leastc	2.649882
GENEX Option C	LeastC	rem_leastc	1.431883
GENEX Option C	LeastC	rem_leastc	9.169954
GENEX Option C	LeastC	rem_leastc	7.708445
GENEX Option C	LeastC	rem_leastc	92.95
GENEX Option C	LeastC	rem_leastc	94.69141
GENEX Option C	LeastC	rem_leastc	5.433617
GENEX Option C	LeastC	rem_leastc	64.5069
GENEX Option C	LeastC	rem_leastc	2.113
GENEX Option C	LeastC	rem_leastc	3.737052
GENEX Option C	LeastC	rem_leastc	174.1968
GENEX Option C	LeastC	rem_leastc	78.54987
GENEX Option C	LeastC	rem_leastc	44.18646
GENEX Option C	LeastC	rem_leastc	302.4601
GENEX Option C	LeastC	rem_leastc	13.03633
GENEX Option C	LeastC	rem_leastc	0.02649
GENEX Option C	LeastC	rem_leastc	4.129607
GENEX Option C	LeastC	rem_leastc	3.67986
GENEX Option C	LeastC	rem_leastc	41.87516
GENEX Option C	LeastC	rem_leastc	123.6476
GENEX Option C	LeastC	rem_leastc	29.82518
GENEX Option C	LeastC	rem_leastc	709.0444
GENEX Option C	LeastC	rem_leastc	60.01941
GENEX Option C	LeastC	rem_leastc	1.604473
GENEX Option C	LeastC	rem_leastc	53.97776
GENEX Option C	LeastC	rem_leastc	186.7913
GENEX Option C	LeastC	rem_leastc	44.12555
GENEX Option C GENEX Option C	LeastC LeastC		
		rem_leastc	44.12555
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517
GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168 59.34102
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168 59.34102 140.7262
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.70751 32.7168 59.34102 140.7262 49.51686
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168 59.34102 140.7262 49.51686 0.707326
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168 59.34102 140.7262 49.51686 0.707326 7.045497
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	44.12555 134.2578 29.8517 14.57555 74.74253 12.649 121.5994 8.356103 70.14877 85.00014 8.976845 114.0486 5.472839 34.84204 101.7071 48.67651 32.7168 59.34102 140.7262 49.51686 0.707326

GENEX Option C	LeastC	rem_leastc	56.10603
GENEX Option C	LeastC	rem leastc	28.6392
GENEX Option C	LeastC	rem_leastc	26.53059
GENEX Option C	LeastC	rem_leastc	22.65911
GENEX Option C	LeastC	rem_leastc	5.090376
GENEX Option C	LeastC	rem_leastc	7.380356
GENEX Option C	LeastC	rem_leastc	18.65086
GENEX Option C	LeastC	rem leastc	8.082343
GENEX Option C	LeastC	rem_leastc	4.497161
GENEX Option C	LeastC	rem leastc	61.08123
GENEX Option C	LeastC	rem_leastc	475.5428
GENEX Option C	LeastC	rem_leastc	102.9064
GENEX Option C	LeastC	rem_leastc	0.252096
GENEX Option C	LeastC	rem_leastc	41.80824
GENEX Option C	LeastC	rem_leastc	296.3187
GENEX Option C	LeastC	rem_leastc	101.5743
GENEX Option C	LeastC	rem_leastc	73.64092
GENEX Option C	LeastC	rem_leastc	14.84019
GENEX Option C	LeastC	rem_leastc	54.19917
GENEX Option C	LeastC	rem_leastc	14.47524
GENEX Option C	LeastC	rem_leastc	6.594339
GENEX Option C	LeastC	rem_leastc	393.9766
GENEX Option C	LeastC	rem_leastc	101.363
GENEX Option C	LeastC	rem_leastc	0.935278
GENEX Option C	LeastC	rem leastc	49.06945
GENEX Option C	LeastC	rem leastc	57.09075
GENEX Option C	LeastC	rem_leastc	0.126688
GENEX Option C	LeastC	rem_leastc	18.24998
GENEX Option C	LeastC	rem_leastc	25.16174
GENEX Option C	LeastC	rem leastc	34.29543
GENEX Option C	LeastC	rem_leastc	42.10126
GENEX Option C	LeastC	rem leastc	14.19841
GENEX Option C	LeastC	rem_leastc	7.371037
GENEX Option C	LeastC	rem leastc	1.347521
GENEX Option C	LeastC	rem_leastc	183.2169
GENEX Option C	LeastC	rem_leastc	31.9329
GENEX Option C	LeastC	rem_leastc	111.3151
GENEX Option C	LeastC	rem_leastc	27.00255
GENEX Option C	LeastC	rem leastc	9.83722
GENEX Option C	LeastC	rem leastc	7.45148
GENEX Option C	LeastC	rem leastc	11.17567
GENEX Option C	LeastC	rem_leastc	82.99118
GENEX Option C	LeastC	rem leastc	15.84509
GENEX Option C	LeastC	rem_leastc	310.9522
GENEX Option C	LeastC	rem leastc	34.4065
GENEX Option C	LeastC	rem leastc	55.20887
GENEX Option C	LeastC	rem leastc	3.677472
GENEX Option C	LeastC	rem_leastc	371.3333
GENEX Option C	LeastC	rem_leastc	5.444018
GENEX Option C			45.88207
GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	0.245078
GENEX Option C			11.2459
GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	25.08376
	LeastC		
GENEX Option C GENEX Option C		rem_leastc	124.1548 143.1277
GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	5.565565
GENEX Option C	LeastC	rem_leastc	49.50317
GENEX Option C	LeastC		
GENEX Option C		rem_leastc	270.3382
	LeastC	rem_leastc	10.59432
GENEX Option C	LeastC	rem_leastc	17.08195
GENEX Option C	LeastC	rem_leastc	99.73551
GENEX Option C	LeastC	rem_leastc	376.538
GENEX Option C	LeastC	rem_leastc	48.17882
GENEX Option C	LeastC	rem_leastc	41.22485
GENEX Option C	LeastC	rem_leastc	17.63204
GENEX Option C	LeastC	rem_leastc	11.48561
GENEX Option C	LeastC	rem_leastc	46.26185
GENEX Option C	LeastC	rem_leastc	1.445556
GENEX Option C	LeastC	rem_leastc	16.18249
GENEX Option C	LeastC	rem_leastc	2.159236
GENEX Option C	LeastC	rem_leastc	0.637007
GENEX Option C	LeastC	rem_leastc	264.9616
GENEX Option C	LeastC	rem_leastc	926.4869
		Irom loacte	283.8492
GENEX Option C	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	1.547627
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	1.547627 203.148
GENEX Option C	LeastC	rem_leastc	1.547627

GENEX Option C	LeastC	rom logete	1.156638
GENEX Option C	LeastC	rem_leastc rem_leastc	82.71442
GENEX Option C	LeastC	rem leastc	2.022596
GENEX Option C	LeastC	rem_leastc	60.74509
GENEX Option C	LeastC	rem_leastc	41.10972
GENEX Option C	LeastC	rem_leastc	14.57307
GENEX Option C	LeastC	rem_leastc	25.81795
GENEX Option C	LeastC	rem_leastc	3.825454
GENEX Option C	LeastC	rem_leastc	27.65567
GENEX Option C	LeastC	rem_leastc	54.14819
GENEX Option C	LeastC	rem leastc	77.56343
GENEX Option C	LeastC	rem leastc	0.005124
GENEX Option C	LeastC	rem_leastc	0.548982
GENEX Option C			
	LeastC	rem_leastc	83.46924
GENEX Option C	LeastC	rem_leastc	61.9752
GENEX Option C	LeastC	rem_leastc	22.51092
GENEX Option C	LeastC	rem_leastc	78.81351
GENEX Option C	LeastC	rem_leastc	1.1722
GENEX Option C	LeastC	rem_leastc	23.31453
GENEX Option C	LeastC	rem leastc	587.4325
GENEX Option C	LeastC	rem_leastc	204.1666
GENEX Option C	LeastC	rem_leastc	260.2845
GENEX Option C		rem leastc	120.4324
	LeastC		
GENEX Option C	LeastC	rem_leastc	865.5664
GENEX Option C	LeastC	rem_leastc	29.93223
GENEX Option C	LeastC	rem_leastc	12.69863
GENEX Option C	LeastC	rem_leastc	103.5631
GENEX Option C	LeastC	rem_leastc	312.3949
GENEX Option C	LeastC	rem_leastc	99.10889
GENEX Option C	LeastC	rem_leastc	10.33737
GENEX Option C	LeastC	rem_leastc	31.16894
GENEX Option C	LeastC	rem leastc	43.44386
GENEX Option C	LeastC	rem_leastc	193.8193
GENEX Option C			183.4683
	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	105.8241
GENEX Option C	LeastC	rem_leastc	30.87396
GENEX Option C	LeastC	rem_leastc	437.6609
GENEX Option C	LeastC	rem_leastc	26.50336
GENEX Option C	LeastC	rem_leastc	43.01024
GENEX Option C	LeastC	rem_leastc	2.674355
GENEX Option C	LeastC	rem leastc	145.642
GENEX Option C	LeastC	rem_leastc	45.83799
GENEX Option C	LeastC	rem_leastc	1.750816
GENEX Option C	LeastC	rem leastc	7.330887
GENEX Option C	LeastC	rem leastc	80.91222
		_	
GENEX Option C	LeastC	rem_leastc	285.9006
GENEX Option C	LeastC	rem_leastc	1.802566
GENEX Option C	LeastC	rem_leastc	59.32959
GENEX Option C	LeastC	rem_leastc	12.23004
GENEX Option C	LeastC	rem_leastc	8.971527
GENEX Option C	LeastC	rem_leastc	13.00588
GENEX Option C	LeastC	rem_leastc	3.75467
GENEX Option C	LeastC	rem_leastc	178.9832
GENEX Option C	LeastC	rem_leastc	107.5745
GENEX Option C	LeastC	rem_leastc	46.31915
GENEX Option C	LeastC	rem_leastc	61.24909
GENEX Option C	LeastC	rem_leastc	88.93272
	Luait	icii_icasic	00.93272
CENEY Option C	LoootC	rom loosto	0 400004
GENEX Option C	LeastC	rem_leastc	8.600226
GENEX Option C	LeastC	rem_leastc	42.1064
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	42.1064 526.2937
GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9999 2.945584 0.027143 22.94944 241.1351 10.82383
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.99582 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139 1.018066
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.99582 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139 1.018066
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139 1.018066 148.4483
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139 1.018066 148.4483 1.735207
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	42.1064 526.2937 4.800436 12.7743 18.0793 70.34255 47.25853 289.9992 2.945584 0.027143 22.94944 241.1351 10.82383 0.217139 1.018066 148.4483 1.735207 200.3061

CENEX Option C	LoactC	rom logsta	0.283346
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	63.82911
GENEX Option C	LeastC	rem leastc	9.483077
GENEX Option C	LeastC	rem_leastc	5.112733
GENEX Option C	LeastC	rem_leastc	42.39342
GENEX Option C	LeastC	rem_leastc	217.1046
GENEX Option C	LeastC	rem_leastc	274.7574
GENEX Option C	LeastC	rem_leastc	44.75944
GENEX Option C	LeastC	rem_leastc	34.00366
GENEX Option C	LeastC	rem_leastc	15.84078
GENEX Option C	LeastC	rem leastc	44.64747
GENEX Option C	LeastC	rem_leastc	122.9537
GENEX Option C			566.3628
	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	14.21165
GENEX Option C	LeastC	rem_leastc	28.92059
GENEX Option C	LeastC	rem_leastc	237.4161
GENEX Option C	LeastC	rem_leastc	82.96334
GENEX Option C	LeastC	rem_leastc	60.5029
GENEX Option C	LeastC	rem_leastc	45.99594
GENEX Option C	LeastC	rem leastc	87.13361
GENEX Option C	LeastC	rem leastc	54.99165
			47.59365
GENEX Option C	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	1.813457
GENEX Option C	LeastC	rem_leastc	14.67879
GENEX Option C	LeastC	rem_leastc	36.05249
GENEX Option C	LeastC	rem_leastc	345.0211
GENEX Option C	LeastC	rem leastc	166.6805
GENEX Option C	LeastC	rem leastc	55.38093
GENEX Option C	LeastC	rem leastc	543.976
GENEX Option C	LeastC	rem_leastc	7.497423
GENEX Option C	LeastC	rem_leastc	441.0962
GENEX Option C	LeastC	rem_leastc	13.40562
GENEX Option C	LeastC	rem_leastc	1.641803
GENEX Option C	LeastC	rem_leastc	74.30006
GENEX Option C	LeastC	rem_leastc	1.650331
GENEX Option C	LeastC	rem leastc	70.1032
GENEX Option C	LeastC	rem leastc	46.96682
GENEX Option C	LeastC	rem leastc	247.8017
GENEX Option C			
	LeastC	rem_leastc	164.7249
GENEX Option C	LeastC	rem_leastc	7.676785
GENEX Option C	LeastC	rem_leastc	497.7703
GENEX Option C	LeastC	rem_leastc	73.2098
GENEX Option C	LeastC	rem_leastc	15.16272
GENEX Option C	LeastC	rem_leastc	30.13856
GENEX Option C	LeastC	rem leastc	24,79674
GENEX Option C	LeastC	rem_leastc	12.80719
GENEX Option C	LeastC	rem leastc	30.18079
GENEX Option C	LeastC	rem_leastc	50.3982
GENEX Option C	LeastC	rem_leastc	2.007452
GENEX Option C	LeastC	rem_leastc	5.297635
GENEX Option C	LeastC	rem_leastc	4.628709
GENEX Option C	LeastC	rem_leastc	0.075479
GENEX Option C	LeastC		
GENEX Option C	Leasic	rem_leastc	23.61133
GENEX Option C	LeastC		
		rem_leastc	23.61133 73.136
GENEX Option C	LeastC LeastC	rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531
GENEX Option C GENEX Option C	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226
GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855
GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453
GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721 89.05145
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721 89.05145 87.37309
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721 89.05145 87.37309 11.05859
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721 89.05145 87.37309 11.05859 493.3194
GENEX Option C GENEX Option C	LeastC LeastC	rem_leastc rem_leastc	23.61133 73.136 875.1531 5.458226 5.761792 61.16559 19.86604 144.7855 40.26181 239.4669 135.8535 6.56791 1.495947 42.50296 0.284174 342.7593 90.20453 11.22949 370.7128 2.448721 89.05145 87.37309 11.05859

GENEX Option C	LeastC	rem_leastc	7.514547
GENEX Option C	LeastC	rem leastc	508.631
GENEX Option C	LeastC	rem_leastc	427.6424
GENEX Option C	LeastC	rem_leastc	104.0435
GENEX Option C	LeastC	rem leastc	4.416903
GENEX Option C	LeastC	rem leastc	0.859066
GENEX Option C	LeastC	rem leastc	44.88932
GENEX Option C	LeastC	rem_leastc	10.31003
GENEX Option C	LeastC	rem leastc	3.907877
GENEX Option C	LeastC	rem_leastc	0.286307
GENEX Option C	LeastC	rem_leastc	3.536081
GENEX Option C	LeastC	rem_leastc	4.302634
GENEX Option C	LeastC	rem_leastc	40.54574
GENEX Option C	LeastC	rem leastc	46.35656
GENEX Option C	LeastC	rem leastc	35.1372
GENEX Option C	LeastC	rem leastc	21.46795
GENEX Option C	LeastC	rem leastc	63.82186
GENEX Option C	LeastC	rem leastc	6.597762
GENEX Option C	LeastC	rem_leastc	22.59072
GENEX Option C	LeastC	rem_leastc	14.81235
GENEX Option C	LeastC	rem leastc	25.26813
GENEX Option C	LeastC	rem leastc	15.78862
GENEX Option C	LeastC	rem_leastc	488.7074
GENEX Option C			488.7074 91.32114
GENEX Option C	LeastC	rem_leastc rem_leastc	91.32114
GENEX Option C	LeastC		
GENEX Option C	LeastC	rem_leastc	9.664155 27.0727
	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	47.85526
GENEX Option C	LeastC	rem_leastc	76.16868
GENEX Option C	LeastC	rem_leastc	478.2442
GENEX Option C	LeastC	rem_leastc	4.100452
GENEX Option C	LeastC	rem_leastc	145.1778
GENEX Option C	LeastC	rem_leastc	64.6371
GENEX Option C	LeastC	rem_leastc	0.226342
GENEX Option C	LeastC	rem_leastc	15.25306
GENEX Option C	LeastC	rem_leastc	167.5648
GENEX Option C	LeastC	rem_leastc	0.043366
GENEX Option C	LeastC	rem_leastc	0.249369
GENEX Option C	LeastC	rem_leastc	31.21998
GENEX Option C	LeastC	rem_leastc	21.50388
GENEX Option C	LeastC	rem_leastc	13.38243
GENEX Option C	LeastC	rem_leastc	48.91066
GENEX Option C	LeastC	rem_leastc	7.692077
GENEX Option C	LeastC	rem_leastc	3.140981
GENEX Option C	LeastC	rem_leastc	2.839104
GENEX Option C	LeastC	rem_leastc	65.74696
GENEX Option C	LeastC	rem_leastc	2.994388
GENEX Option C	LeastC	rem_leastc	0.739912
GENEX Option C	LeastC	rem_leastc	17.49671
GENEX Option C	LeastC	rem_leastc	70.5411
GENEX Option C	LeastC	rem_leastc	13.06975
GENEX Option C	LeastC	rem_leastc	2.843588
GENEX Option C	LeastC	rem_leastc	47.54403
GENEX Option C	LeastC	rem_leastc	54.55183
GENEX Option C	LeastC	rem_leastc	20.83647
GENEX Option C	LeastC	rem_leastc	22.70387
GENEX Option C	LeastC	rem_leastc	1388.618
GENEX Option C	LeastC	rem_leastc	2023.063
GENEX Option C	LeastC	rem_leastc	23.4374
GENEX Option C	LeastC	rem_leastc	16.91414
GENEX Option C	LeastC	rem_leastc	93.94763
GENEX Option C	LeastC	rem_leastc	126.8936
GENEX Option C	LeastC	rem_leastc	330.4297
GENEX Option C	LeastC	rem_leastc	18.47445
GENEX Option C	LeastC	rem_leastc	4.334424
GENEX Option C	LeastC	rem_leastc	7.76446
GENEX Option C	LeastC	rem_leastc	2.66187
GENEX Option C	LeastC	rem_leastc	101.9941
GENEX Option C	LeastC	rem_leastc	33.50958
GENEX Option C	LeastC	rem_leastc	500.0821
GENEX Option C	LeastC	rem_leastc	27.68105
GENEX Option C	LeastC	rem_leastc	50.82704
GENEX Option C	LeastC	rem_leastc	27.28552
GENEX Option C	LeastC	rem_leastc	123.2228
	LeastC	rem_leastc	367.1263
		. sm_icusic	
GENEX Option C	LeastC	rem leaste	34 76044
GENEX Option C	LeastC	rem_leastc	34.76944 107.844
	LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	34.76944 107.844 80.41676

GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC LeastC LeastC LeastC	rem_leastc rem_leastc rem_leastc	180.466 1.27862 24.74313
GENEX Option C GENEX Option C GENEX Option C GENEX Option C	LeastC	-	
GENEX Option C GENEX Option C GENEX Option C			
GENEX Option C GENEX Option C		rem leastc	0.952919
GENEX Option C	LeastC	rem_leastc	12.90265
	LeastC	rem leastc	0.324657
	LeastC	rem leastc	34.31719
GENEX Option C	LeastC	rem leastc	4.292203
GENEX Option C	LeastC	rem leastc	12.06587
GENEX Option C	LeastC	rem leastc	3.785718
GENEX Option C	LeastC	rem leastc	0.559217
GENEX Option C	LeastC	rem leastc	170.6071
GENEX Option C	LeastC	rem_leastc	44.47043
GENEX Option C	LeastC	rem leastc	2.756305
GENEX Option C	LeastC	rem leastc	81.53098
GENEX Option C	LeastC	rem leastc	8.319718
GENEX Option C	LeastC	rem_leastc	2.661891
GENEX Option C	LeastC	rem leastc	13.44255
GENEX Option C	LeastC	rem leastc	5.069492
GENEX Option C	LeastC	rem leastc	17.85633
GENEX Option C	LeastC	rem leastc	15.8381
GENEX Option C	LeastC	rem leastc	59.0309
GENEX Option C	LeastC	rem leastc	135.982
GENEX Option C	LeastC	rem leastc	26.17916
GENEX Option C	LeastC	rem leastc	12.59796
GENEX Option C	LeastC	rem leastc	29.47318
GENEX Option C	LeastC	rem leastc	109.8415
GENEX Option C	LeastC	rem leastc	149.8079
GENEX Option C	LeastC	rem_leastc	462.3725
GENEX Option C	LeastC	rem_leastc	462.3725
GENEX Option C	LeastC	rem leastc	96.2025
GENEX Option C	LeastC	rem leastc	11.50008
GENEX Option C	LeastC	rem_leastc	10.76424
GENEX Option C		-	183.6566
	LeastC	rem_leastc	
GENEX Option C	LeastC	rem_leastc	17.68986
GENEX Option C	LeastC	rem_leastc	190.5614
GENEX Option C	LeastC	rem_leastc	12.3432
GENEX Option C	LeastC	rem_leastc	59.1258
GENEX Option C	LeastC	rem_leastc	8.331762
GENEX Option C	LeastC	rem_leastc	2.857761
GENEX Option C	LeastC	rem_leastc	8.412393
GENEX Option C	LeastC	rem_leastc	604.5275
GENEX Option C	LeastC	rem_leastc	9.989402
GENEX Option C	LeastC	rem_leastc	61.22569
GENEX Option C	LeastC	rem_leastc	26.86946
GENEX Option C	LeastC	rem_leastc	1.759112
GENEX Option C	LeastC	rem_leastc	129.2239
GENEX Option C	LeastC	rem_leastc	13.45173
GENEX Option C	LeastC	rem_leastc	2.664493
GENEX Option C	LeastC	rem_leastc	58.33691
GENEX Option C	LeastC	rem_leastc	17.44888
GENEX Option C	LeastC	rem_leastc	7.161197
GENEX Option C	LeastC	rem_leastc	348.5125
GENEX Option C	LeastC	rem_leastc	0.257113
GENEX Option C	LeastC	rem_leastc	9.412106
GENEX Option C	LeastC	rem_leastc	1.264701
GENEX Option C	LeastC	rem_leastc	553.999
GENEX Option C	LeastC	rem_leastc	192.4199
GENEX Option C	LeastC	rem_leastc	9.22467
GENEX Option C	LeastC	rem_leastc	7.926539
GENEX Option C	LeastC	rem_leastc	29.46511
GENEX Option C	LeastC	rem_leastc	324.4124
GENEX Option C	LeastC	rem_leastc	4.364596
GENEX Option C	LeastC	rem_leastc	248.962
GENEX Option C	LeastC	rem_leastc	10.31886
GENEX Option C	LeastC	rem_leastc	215.5128
GENEX Option C	LeastC	rem_leastc	64.58989
GENEX Option C	LeastC	rem_leastc	25.19911
GENEX Option C	LeastC	rem_leastc	404.6123
GENEX Option C	LeastC	rem_leastc	0.053489
GENEX Option C	LeastC	rem_leastc	4.220058
GENEX Option C	LeastC	rem_leastc	10.04527
GENEX Option C	LeastC	rem_leastc	2.741389
GENEX Option C	LeastC	rem_leastc	0.589376
GENEX Option C	LeastC	rem_leastc	25.03669
OF NEW O HILL O	LeastC	rem_leastc	0.008478
GENEX Option C	1 10	rem_leastc	0.289362
GENEX Option C GENEX Option C GENEX Option C	LeastC	rem_leastc	25.32193

GENEX Option C	LeastC	rem_leastc	120.4981	
GENEX Option C	LeastC	rem_leastc	192.5669	
GENEX Option C	LeastC	rem_leastc	53.11645	
GENEX Option C	LeastC	rem_leastc	22.14615	
GENEX Option C	LeastC	rem_leastc	17.90449	
GENEX Option C	LeastC	rem_leastc	100.6029	
GENEX Option C	LeastC	rem_leastc	40.38307	
GENEX Option C	LeastC	rem_leastc	16.41434	
GENEX Option C	LeastC	rem_leastc	9.77248	
GENEX Option C	LeastC	rem_leastc	32.21486	
GENEX Option C	LeastC	rem_leastc	46.3546	
GENEX Option C	LeastC	rem_leastc	0.008122	
GENEX Option C	LeastC LeastC	rem_leastc	44.16289 5.923245	
GENEX Option C	LeastC LeastC	rem_leastc		
GENEX Option C		rem_leastc	3.490973	
GENEX Option C	LeastC	rem_leastc	131.4287	
GENEX Option C	LeastC	rem_leastc	8.152129	
GENEX Option C	LeastC	rem_leastc	0.000275	
GENEX Option C	LeastC	rem_leastc	0.000024	
GENEX Option C	LeastC	rem_leastc	63.09774	
GENEX Option C	LeastC	rem_leastc	1.551911	
GENEX Option C	LeastC	rem_leastc	6.945101	
GENEX Option C	LeastC	rem_leastc	0.000132	
GENEX Option C	LeastC	rem_leastc	11.23512	
GENEX Option C	LeastC	rem_leastc	3.025342	
GENEX Option C	LeastC	rem_leastc	96.39073	
GENEX Option C	LeastC	rem_leastc	15.80917	
GENEX Option C	LeastC	rem_leastc	192.9281	
GENEX Option C	LeastC	rem_leastc	0.037976	
GENEX Option C	LeastC	rem leastc	49.50354	
GENEX Option C	LeastC	rem_leastc	125.1382	
GENEX Option C	LeastC	rem_leastc	20.527	
GENEX Option C	LeastC		2.774712	
GENEX Option C	LeastC	rem_leastc	2.774712	
		rem_leastc		
GENEX Option C	LeastC	rem_leastc	4.059098	
GENEX Option C	LeastC	rem_leastc	0.453452	
GENEX Option C	LeastC	rem_leastc	12.2734	
GENEX Option C	LeastC	rem_leastc	8.720471	
GENEX Option C	LeastC	rem_leastc	9.605091	
GENEX Option C	LeastC	rem_leastc	0.326735	
GENEX Option C	LeastC	rem_leastc	0.727728	
GENEX Option C	LeastC	rem_leastc	0.288437	
GENEX Option C	LeastC	rem_leastc	13.82986	
GENEX Option C	LeastC	rem_leastc	57.22385	
GENEX Option C	LeastC	rem_leastc	4.842688	
GENEX Option C	LeastC	rem_leastc	32.01963	
GENEX Option C	LeastC	rem_leastc	0.78514	
GENEX Option C	LeastC	rem_leastc	31.26611	
GENEX Option C	LeastC	rem leastc	0.826022	
GENEX Option C	LeastC	rem_leastc	0.584156	
GENEX Option C	LeastC	rem_leastc	0.584156	
GENEX Option C	LeastC	rem_leastc	6.922604	
GENEX Option C	LeastC	rem_leastc	51.43218	
GENEX Option C	LeastC	rem_leastc	133.6165	
GENEX Option C	LeastC	rem_leastc	10.23287	
GENEX Option C	LeastC	rem_leastc	0.001706	
GENEX Option C	O-dom	rem_oc	36.82174	_
GENEX Option C	O-dom	rem_oc	204.122	
GENEX Option C	O-dom	rem_oc	11.61961	
GENEX Option C	O-dom	rem_oc	1.779747	
GENEX Option C	O-dom	rem_oc	8.058584	
GENEX Option C	O-dom	rem_oc	0.817491	
GENEX Option C	O-dom	rem_oc	2.47765	
GENEX Option C	O-dom	_	3.34409	
GENEX Option C	O-dom O-dom	rem_oc	5.401993	
GENEX Option C		rem_oc	5.250489	
	O-dom	rem_oc		
GENEX Option C	O-dom	rem_oc	4.760309	
GENEX Option C	O-dom	rem_oc	0.746089	
GENEX Option C	O-dom	rem_oc	270.0271	
GENEX Option C	O-dom	rem_oc	374.3618	
GENEX Option C	O-dom	rem_oc	1.946463	
	O-dom	rem_oc	1.771366	
GENEX Option C		rem_oc	108.0733	
GENEX Option C GENEX Option C	O-dom	10111_00		
	O-dom O-dom	rem_oc	0.590693	
GENEX Option C GENEX Option C		rem_oc		
GENEX Option C GENEX Option C GENEX Option C	O-dom O-dom	rem_oc rem_oc	26.872	
GENEX Option C GENEX Option C	O-dom	rem_oc		

GENEX Option C         O-dom         rem_oc         21.5996           GENEX Option C         O-dom         rem_oc         21.60491           GENEX Option C         O-dom         rem_oc         21.60491           GENEX Option C         O-dom         rem_oc         9.712734           GENEX Option C         O-dom         rem_oc         0.046029           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         1.30439           GENEX Option C         O-dom         rem_oc         0.05237           GENEX Option C         O-dom         rem_oc         0.02339           GENEX Option C         O-dom         rem_oc         0.52309           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom	GENEX Option C	0-dom	rom oc	5.615905
GENEX Option C         O-dom         rem_oc         20.59073           GENEX Option C         O-dom         rem_oc         9.712734           GENEX Option C         O-dom         rem_oc         9.712734           GENEX Option C         O-dom         rem_oc         9.712734           GENEX Option C         O-dom         rem_oc         0.046029           GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         1.3043967           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         1.306325           GENEX Option C         O-dom         rem_oc         1.306325           GENEX Option C         O-dom <td></td> <td></td> <td>rem_oc</td> <td></td>			rem_oc	
GENEX Option C         O-dom         rem_oc         21.60491           GENEX Option C         O-dom         rem_oc         5.799722           GENEX Option C         O-dom         rem_oc         0.046029           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         1.30439           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.652309           GENEX Option C         O-dom         rem_oc         6.84166           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.18017           GENEX Option C         O-dom         rem_oc         0.813551           GENEX Option C         O-dom         rem_oc         0.8136551           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom <td></td> <td></td> <td></td> <td></td>				
GENEX Option C         O-dom         rem_oc         9.712734           GENEX Option C         O-dom         rem_oc         5.799722           GENEX Option C         O-dom         rem_oc         0.046029           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.653237           GENEX Option C         O-dom         rem_oc         0.62337           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.438077           GENEX Option C         O-dom         rem_oc         4.438077           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         1.304325           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         9.563262           GENEX Option C         O-dom <td></td> <td></td> <td></td> <td></td>				
GENEX Option C         O-dom         rem_oc         5.799722           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         4.7964713           GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.33967           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         0.553262           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom			-	
GENEX Option C         O-dom         rem_oc         0.046029           GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         1.30439           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.053237           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         68.1465           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         5.53262           GENEX Option C         O-dom         rem_oc         5.53262           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom				
GENEX Option C         O-dom         rem_oc         4.964713           GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.6502309           GENEX Option C         O-dom         rem_oc         58.61665           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.11942           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom </td <td>· · · · · · · · · · · · · · · · · · ·</td> <td></td> <td>_</td> <td></td>	· · · · · · · · · · · · · · · · · · ·		_	
GENEX Option C         O-dom         rem_oc         5.718127           GENEX Option C         O-dom         rem_oc         1.30439           GENEX Option C         O-dom         rem_oc         0.502307           GENEX Option C         O-dom         rem_oc         0.053237           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         8.433967           GENEX Option C         O-dom         rem_oc         8.433967           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.503262           GENEX Option C         O-dom         rem_oc         1.176356           GENEX Option C         O-dom         rem_oc         1.378326           GENEX Option C         O-dom         rem_oc         1.50552           GENEX Option C         O-dom			_	
GENEX Option C         O-dom         rem_oc         1.30439           GENEX Option C         O-dom         rem_oc         0.0502309           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         58.61665           GENEX Option C         O-dom         rem_oc         42.86207           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.418017           GENEX Option C         O-dom         rem_oc         0.418017           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         5.63262           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom				
GENEX Option C         O-dom         rem_oc         0.502309           GENEX Option C         O-dom         rem_oc         0.053237           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.438677           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         1.8654748           GENEX Option C         O-dom         rem_oc         1.76556           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         1.765552           GENEX Option C         O-dom         rem_oc         1.651522           GENEX Option C         O-dom <td></td> <td></td> <td>-</td> <td></td>			-	
GENEX Option C         O-dom         rem_oc         0.053237           GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         58.61665           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         5.663262           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         1.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         1.64185           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom			_	
GENEX Option C         O-dom         rem_oc         0.43457           GENEX Option C         O-dom         rem_oc         58.61665           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.43867           GENEX Option C         O-dom         rem_oc         0.813957           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.50532           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom				
GENEX Option C         O-dom         rem_oc         58.61665           GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.416511           GENEX Option C         O-dom         rem_oc         0.616551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.667498           GENEX Option C         O-dom         rem_oc         5.63262           GENEX Option C         O-dom         rem_oc         1.76356           GENEX Option C         O-dom         rem_oc         1.50552           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom			-	
GENEX Option C         O-dom         rem_oc         62.13515           GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         8.433967           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.1360325           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.50526           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom <td></td> <td></td> <td></td> <td></td>				
GENEX Option C         O-dom         rem_oc         4.286207           GENEX Option C         O-dom         rem_oc         8.433967           GENEX Option C         O-dom         rem_oc         0.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         17.76356           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         11.6185           GENEX Option C         O-dom         rem_oc         18.22025           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom			rem_oc	
GENEX Option C         O-dom         rem_oc         8.433967           GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         17.6356           GENEX Option C         O-dom         rem_oc         15.5052           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         4.488126           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom		O-dom	rem_oc	62.13515
GENEX Option C         O-dom         rem_oc         4.418017           GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         1.76356           GENEX Option C         O-dom         rem_oc         1.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         1.64185           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom	GENEX Option C	O-dom	rem_oc	4.286207
GENEX Option C         O-dom         rem_oc         0.816551           GENEX Option C         O-dom         rem_oc         1.360325           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         9.167498           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.63262           GENEX Option C         O-dom         rem_oc         5.63262           GENEX Option C         O-dom         rem_oc         5.63262           GENEX Option C         O-dom         rem_oc         5.810314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         18.14182           GENEX Option C         O-dom         rem_oc         18.22015           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom	GENEX Option C	O-dom	rem_oc	8.433967
GENEX Option C         O-dom         rem_oc         1.360325           GENEX Option C         O-dom         rem_oc         0.61658           GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom	GENEX Option C	O-dom	rem_oc	4.418017
GENEX Option CO-domrem_oc0.61658GENEX Option CO-domrem_oc0.594068GENEX Option CO-domrem_oc9.111942GENEX Option CO-domrem_oc8.567498GENEX Option CO-domrem_oc5.563262GENEX Option CO-domrem_oc5.563262GENEX Option CO-domrem_oc5.50552GENEX Option CO-domrem_oc11.76356GENEX Option CO-domrem_oc3.278022GENEX Option CO-domrem_oc3.278022GENEX Option CO-domrem_oc3.436123GENEX Option CO-domrem_oc11.64185GENEX Option CO-domrem_oc11.64185GENEX Option CO-domrem_oc0.0018GENEX Option CO-domrem_oc4.444007GENEX Option CO-domrem_oc4.444007GENEX Option CO-domrem_oc0.338932GENEX Option CO-domrem_oc0.3072902GENEX Option CO-domrem_oc3.072902GENEX Option CO-domrem_oc3.072902GENEX Option CO-domrem_oc3.072902GENEX Option CO-domrem_oc4.18188GENEX Option CO-domrem_oc4.045188GENEX Option CO-domrem_oc0.00344GENEX Option CO-domrem_oc0.645188GENEX Option CO-domrem_oc0.6421117GENEX	GENEX Option C	O-dom	rem_oc	0.816551
GENEX Option C         O-dom         rem_oc         0.594068           GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.8.10314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         4.88126           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom	GENEX Option C	O-dom	rem_oc	1.360325
GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.810314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         14.88126           GENEX Option C         O-dom         rem_oc         3.478022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom <td>GENEX Option C</td> <td>O-dom</td> <td>rem_oc</td> <td>0.61658</td>	GENEX Option C	O-dom	rem_oc	0.61658
GENEX Option C         O-dom         rem_oc         9.111942           GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.810314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         14.88126           GENEX Option C         O-dom         rem_oc         3.478022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom <td>GENEX Option C</td> <td>O-dom</td> <td>rem oc</td> <td>0.594068</td>	GENEX Option C	O-dom	rem oc	0.594068
GENEX Option C         O-dom         rem_oc         8.567498           GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         5.810314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.00344           GENEX Option C         O-dom		O-dom		9.111942
GENEX Option C         O-dom         rem_oc         5.563262           GENEX Option C         O-dom         rem_oc         58.10314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         15.50552           GENEX Option C         O-dom         rem_oc         4.488126           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         18.22205           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         0.00344           GENEX Option C         O-dom         rem_oc         0.21117           GENEX Option C         O-dom		O-dom	rem oc	8.567498
GENEX Option C         O-dom         rem_oc         58.10314           GENEX Option C         O-dom         rem_oc         11.76356           GENEX Option C         O-dom         rem_oc         15.50552           GENEX Option C         O-dom         rem_oc         4.488126           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0338932           GENEX Option C         O-dom         rem_oc         0.4444007           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.45188           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom		O-dom		
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GENEX Option C         O-dom         rem_oc         15.50552           GENEX Option C         O-dom         rem_oc         4.488126           GENEX Option C         O-dom         rem_oc         3.278022           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         18.22205           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         4.45188           GENEX Option C         O-dom         rem_oc         4.612949           GENEX Option C         O-dom			-	
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GENEX Option C         O-dom         rem_oc         3.436123           GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         18.22205           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.00344           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         1.81888           GENEX Option C         O-subdom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.309472           GENEX Option C         O-su				
GENEX Option C         O-dom         rem_oc         11.64185           GENEX Option C         O-dom         rem_oc         18.22205           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.45188           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         3.072902           GENEX Option C         O-subdom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.18294           GENEX Option C         O-subdo				
GENEX Option C         O-dom         rem_oc         18.22205           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.45188           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.221117           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         4.94581           GENEX Option C         O-subdom         rem_oc         3.073514           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdo			_	
GENEX Option C         O-dom         rem_oc         0.0018           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         <			_	
GENEX Option C         O-dom         rem_oc         4.444007           GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.100344           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-dom         rem_oc         1.81898           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C			-	
GENEX Option C         O-dom         rem_oc         6.059981           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         15.98738           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.100344           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-dom         rem_oc         1.81898           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C				
GENEX Option C         O-dom         rem_oc         0.338932           GENEX Option C         O-dom         rem_oc         15.98738           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         0.199158           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         68.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C			_	
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GENEX Option C         O-dom         rem_oc         0.645188           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         9.199158           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option			-	
GENEX Option C         O-dom         rem_oc         3.072902           GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         9.199158           GENEX Option C         O-dom         rem_oc         9.199158           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.302472           GENEX Option C         O-subdom         rem_oc         6.309472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-dom         rem_oc         0.000344           GENEX Option C         O-dom         rem_oc         9.199158           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         11.81888           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         6.873514           GENEX Option C         O-subdom         rem_oc         6.309472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-dom         rem_oc         9.199158           GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         11.81888           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         6.873514           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577			_	
GENEX Option C         O-dom         rem_oc         6.182994           GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         11.81888           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577			_	
GENEX Option C         O-dom         rem_oc         6.221117           GENEX Option C         O-subdom         rem_oc         11.81888           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         11.81888           GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         24.94581           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         0.873514           GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         63.09472           GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577			_	
GENEX Option C         O-subdom         rem_oc         58.16346           GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         7.270746           GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577				
GENEX Option C         O-subdom         rem_oc         2.117614           GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577			_	
GENEX Option C         O-subdom         rem_oc         59.27586           GENEX Option C         O-subdom         rem_oc         21.51577			_	
GENEX Option C O-subdom rem_oc 21.51577				
CENEX Option C O subdom rom oc 05 42149			rem_oc	
	GENEX Option C	O-subdom	rem_oc	85.42148

334 Total OC-subdom remnant for Option C

## Kidston Corridor Option A - Woody vegetation assessment

Name	Class_name	VMA_Status	Area_ha_options	
Genex: Option A	AdjustNC_LeastC	Least concern	0.3024611	37602 Total Least concern remnant for Option A
Genex: Option A	NONREMNANT	Non-remnant	0.0865846	2174 Total non-remnant vegetation for Option A
Genex: Option A	EASE	Non-remnant bare earth and water	0.08631	4670 Total non-remnant bare earth and water for Option A
Genex: Option A	O-dom	Of concern - dominant	0.0896216	1535 Total Of concern-dom remnant for Option A
Genex: Option A	grassOC_O-subdom	Of concern - sub dominant	0.09	549 Total Of concern-subdom for Option A
Genex: Option A	ScatNC_LeastC	Potential least concern	0.0945669	9002 Total Potential Least concern remnant for Option A
Genex: Option A	ScatNC_O-dom	Potential of concern - dominant	0.1799997	240 Total Potential Of concern-dom remnant for Option A
Genex: Option A	ScatOC_O-subdom	Potential of concern - subdominant	0.36	364 Total Potential Of concern-subdom remnant vegetation for Option A
Genex: Option A	REGROWTH	Potential regrowth	0.18	235 Total Potential regrowth for Option A
Genex: Option A	Regrowth_LeastC	Regrowth Least concern	1.2600001	48 Total Least concern regrowth for Option A
Genex: Option A	Regrowth_O-dom	Regrowth Of concern dominant	1.2599993	6.9 Total Of concern-dom regrowth for Option A

## Kidston Corridor Option B - Woody vegetation assessment

Name	Class_name	VMA_Status	Area_ha_options	
Genex: Option B	AdjustNC_LeastC	Least concern	0.3024611	37969 Total Least concern remnant for Option B
Genex: Option B	NONREMNANT	Non-remnant	0.0865846	1812 Total non-remnant vegetation for Option B
Genex: Option B	EASE	Non-remnant bare earth and water	0.08631	5428 Total non-remnant bare earth and water for Option B
Genex: Option B	O-dom	Of concern - dominant	0.0896216	1460 Total Of concern-dom remnant for Option B
Genex: Option B	AdjustOC_O-subdom	Of concern - sub dominant	0.09	295 Total Of concern-subdom for Option B
Genex: Option B	ScatNC_LeastC	Potential least concern	0.0945669	8439 Potential Least concern remnant for Option B
Genex: Option B	ScatOC_O-dom	Potential of concern - dominant	1.2775804	344 Total Potential Of concern-dom remnant for Option B
Genex: Option B	dryOC_O-subdom	Potential of concern - subdominant	0.1759937	87 Total Potential Of concern-subdom remnant for Option B
Genex: Option B	REGROWTH	Potential regrowth	0.18	219 Total Potential regrowth for Option B
Genex: Option B	Regrowth_LeastC	Regrowth Least concern	1.2600001	214 Total Least concern regrowth for Option B
Genex: Option B	NONREMNANT	Regrowth Of concern dominant	0.09	6.6 Total Of concern-dom regrowth for Option B

## Kidston Corridor Option C - Woody vegetation assessment

Name	Class_name	VMA_Status	Area_ha_options	
GENEX Option C	AdjustNC_LeastC	Least concern	0.3024611	38808 Total Least concern remnant for Option C
GENEX Option C	NONREMNANT	Non-remnant	0.0865846	1718 Total non-remnant vegetation for Option C
GENEX Option C	EASE	Non-remnant bare earth and water	0.08631	5771 Total non-remnant bare earth and water for Option C
GENEX Option C	O-dom	Of concern - dominant	0.0896216	1223 Total Of concern-dom remnant for Option C
GENEX Option C	AdjustOC_O-subdom	Of concern - sub dominant	0.09	618 Total Of concern-subdom for Option C
GENEX Option C	ScatNC_LeastC	Potential least concern	0.0945669	
GENEX Option C	ScatOC_O-dom	Potential of concern - dominant	1.2775804	431 Total Potential Of concern-dom remnant for Option C
GENEX Option C	dryOC_O-subdom	Potential of concern - subdominant	0.1759937	37 Total Potential Of concern-subdom remnant vegetation for Option C
GENEX Option C	REGROWTH	Potential regrowth	0.18	291 Total Potential regrowth for Option C
GENEX Option C	Regrowth_LeastC	Regrowth Least concern	1.2600001	172 Total Least concern regrowth for Option C
GENEX Option C	Regrowth_O-dom	Regrowth Of concern dominant	1.2599993	28.8 Total Of concern-dom regrowth for Option C

## Kidston Corridor Option A - Regulated remnant watercourse vegetation

Name	Scale	Bioregion_type	SO_group	Buffer_m	Area_ha	
Genex: Option A		Non-coastal	1 or 2	25	2579.422	
Genex: Option A		Non-coastal	1 or 2	25	520.0739	
Genex: Option A		Non-coastal	1 or 2	25	45.31245	
Genex: Option A		Non-coastal	1 or 2	25	21.86686	
Genex: Option A		Non-coastal	1 or 2	25	243.4169	
Genex: Option A		Non-coastal	1 or 2	25	11.34511	
Genex: Option A		Non-coastal	1 or 2	25	4.666213	
Genex: Option A		Non-coastal	1 or 2	25	5.498384	
Genex: Option A		Non-coastal	1 or 2	25	0.118428	
Genex: Option A		Non-coastal	1 or 2	25	0.414786	
Genex: Option A		Non-coastal	1 or 2	25	5.22667	
Genex: Option A		Non-coastal	1 or 2	25	2.777769	
Genex: Option A		Non-coastal	1 or 2	25	173.8444	
Genex: Option A		Non-coastal	1 or 2	25	3.009753	
Genex: Option A		Non-coastal	1 or 2	25	1.857839	3619 Total SO 1 or 2
Genex: Option A		Non-coastal	3 or 4	50		3019 TOTAL SO T OF 2
					815.3475	
Genex: Option A		Non-coastal	3 or 4	50	304.8283	
Genex: Option A		Non-coastal	3 or 4	50	45.31245	
Genex: Option A		Non-coastal	3 or 4	50	13.60399	
Genex: Option A		Non-coastal	3 or 4	50	64.05599	
Genex: Option A		Non-coastal	3 or 4	50	4.666213	
Genex: Option A		Non-coastal	3 or 4	50	15.28811	
Genex: Option A		Non-coastal	3 or 4	50	0.414786	
Genex: Option A		Non-coastal	3 or 4	50	5.22667	
Genex: Option A		Non-coastal	3 or 4	50	1.382051	
Genex: Option A		Non-coastal	3 or 4	50	14.05157	
Genex: Option A		Non-coastal	3 or 4	50	13.67417	
Genex: Option A		Non-coastal	3 or 4	50	3.009753	1301 Total SO 3 or 4
Genex: Option A		Non-coastal	5 or greater	100	569.3731	
Genex: Option A		Non-coastal	5 or greater	100	21.86686	
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A		Non-coastal	5 or greater	100		
Genex: Option A	100000	Non-coastal	5 or greater	100	1.857839	665 Total SO 5 or greater
<u> </u>			•			
Name			SO_group	Buffer_m		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		
Genex: Option A		Coastal	1 or 2	10		65 Total SO 1 or 2
Genex: Option A		Coastal	3 or 4	25		
Genex: Option A		Coastal	3 or 4	25		
Genex: Option A	100000	Coastal	3 or 4	25	5.460993	
Genex: Option A	100000	Coastal	3 or 4	25	7.603141	
Genex: Option A	100000	Coastal	3 or 4	25	0.273101	
Genex: Option A	1	Coastal	3 or 4	25	0.132366	
Genex: Option A	1	Coastal	3 or 4	25	13.3057	
Genex: Option A		Coastal	3 or 4	25		53 Total SO 3 or 4
				_5		

## Kidston Corridor Option B - Regulated remnant watercourse vegetation

Name	Scale	Bioregion_type	SO_group	Buffer_m	Area ha	
Genex: Option B		Non-coastal	1 or 2		1060.103	
Genex: Option B		Non-coastal	1 or 2		177.1531	
Genex: Option B		Non-coastal	1 or 2	25	21.94918	
Genex: Option B		Non-coastal	1 or 2	25	7.799067	
Genex: Option B		Non-coastal	1 or 2	25	243.4169	
Genex: Option B		Non-coastal	1 or 2	25	1370.482	
Genex: Option B		Non-coastal	1 or 2	25	5.498384	
· · · · · · · · · · · · · · · · · · ·		Non-coastal		25		
Genex: Option B		Non-coastal	1 or 2 1 or 2	25	400.4101 5.22667	
Genex: Option B						
Genex: Option B		Non-coastal Non-coastal	1 or 2	25	32.08263	
Genex: Option B			1 or 2	25	2.777769	
Genex: Option B		Non-coastal Non-coastal	1 or 2 1 or 2	25 25	9.573953	
Genex: Option B					173.8444	
Genex: Option B		Non-coastal	1 or 2	25 25	2.500361	
Genex: Option B		Non-coastal	1 or 2	25	0.164711	
Genex: Option B		Non-coastal	1 or 2		3.009753	2510 Total SO 1 or 2
Genex: Option B		Non-coastal	1 or 2	25	1.857839	3518 Total SO 1 or 2
Genex: Option B		Non-coastal	3 or 4	50		
Genex: Option B		Non-coastal	3 or 4	50	60.40728	
Genex: Option B		Non-coastal	3 or 4	50	21.94918	
Genex: Option B		Non-coastal	3 or 4	50	7.825659	
Genex: Option B		Non-coastal	3 or 4	50		
Genex: Option B		Non-coastal	3 or 4		426.1213	
Genex: Option B		Non-coastal	3 or 4	50		
Genex: Option B		Non-coastal	3 or 4	50	5.22667	
Genex: Option B		Non-coastal	3 or 4	50	32.08263	
Genex: Option B		Non-coastal	3 or 4	50		
Genex: Option B		Non-coastal	3 or 4	50	5.80639	
Genex: Option B		Non-coastal	3 or 4	50	14.05157	
Genex: Option B		Non-coastal	3 or 4	50	2.500361	
Genex: Option B		Non-coastal	3 or 4	50	13.67417	
Genex: Option B		Non-coastal	3 or 4	50	0.164711	
Genex: Option B		Non-coastal	3 or 4	50		1325 Total SO 3 or 4
Genex: Option B		Non-coastal	5 or greater	100	313.161	
Genex: Option B		Non-coastal	5 or greater	100		
Genex: Option B	100000	Non-coastal	5 or greater	100		
Genex: Option B		Non-coastal	5 or greater	100	44.12056	
Genex: Option B		Non-coastal	5 or greater	100	300.0754	
Genex: Option B	100000	Non-coastal	5 or greater	100	2.777769	
Genex: Option B		Non-coastal	5 or greater		9.573953	
Genex: Option B		Non-coastal	5 or greater	100	1.382051	
Genex: Option B	100000	Non-coastal	5 or greater	100	5.80639	
Genex: Option B	100000	Non-coastal	5 or greater	100	9.204484	
Genex: Option B	100000	Non-coastal	5 or greater	100	0.164711	
Genex: Option B	100000	Non-coastal	5 or greater	100	1.857839	704 Total SO 5 or greater
Name	Scale	Bioregion_type	SO_group	Buffer_m		
Genex: Option B		Coastal	1 or 2		0.844454	
Genex: Option B		Coastal	1 or 2		9.624821	
Genex: Option B		Coastal	1 or 2		0.502938	
Genex: Option B		Coastal	1 or 2		0.273101	
Genex: Option B		Coastal	1 or 2	10	18.08111	
Genex: Option B	100000	Coastal	1 or 2	10	0.545677	30 Total SO 1 or 2
Genex: Option B	100000	Coastal	3 or 4	25		
Genex: Option B		Coastal	3 or 4	25	0.273101	
Genex: Option B		Coastal	3 or 4	25	13.3057	
Genex: Option B	100000	Coastal	3 or 4	25	0.545677	20 Total SO 3 or 4

## Kidston Corridor Option C - Regulated remnant watercourse vegetation

Name	Scale	Bioregion_type	SO_group	Buffer_m	Area ha	
GENEX Option C		Non-coastal	1 or 2	25		
GENEX Option C		Non-coastal	1 or 2	25		
GENEX Option C		Non-coastal	1 or 2	25		
GENEX Option C		Non-coastal	1 or 2	25	5.424669	
GENEX Option C		Non-coastal	1 or 2	25	11.34511	
GENEX Option C		Non-coastal	1 or 2	25	1370.482	
-				25		
GENEX Option C		Non-coastal	1 or 2		1.467238	
GENEX Option C		Non-coastal	1 or 2	25		
GENEX Option C		Non-coastal	1 or 2	25	400.4101	
GENEX Option C		Non-coastal	1 or 2	25	32.08263	
GENEX Option C		Non-coastal	1 or 2	25	9.573953	
GENEX Option C		Non-coastal	1 or 2	25	173.8444	
GENEX Option C		Non-coastal	1 or 2	25	2.500361	
GENEX Option C		Non-coastal	1 or 2	25		
GENEX Option C		Non-coastal	1 or 2	25	3.009753	
GENEX Option C		Non-coastal	1 or 2	25	1.857839	3511 Total SO 1 or 2
GENEX Option C		Non-coastal	3 or 4	50	608.131	
GENEX Option C		Non-coastal	3 or 4	50		
GENEX Option C		Non-coastal	3 or 4	50	32.64234	
GENEX Option C		Non-coastal	3 or 4	50	8.930733	
GENEX Option C	100000	Non-coastal	3 or 4	50	426.1213	
GENEX Option C	250000	Non-coastal	3 or 4	50	1.467238	
GENEX Option C	250000	Non-coastal	3 or 4	50	15.28811	
GENEX Option C	250000	Non-coastal	3 or 4	50	144.8488	
GENEX Option C	100000	Non-coastal	3 or 4	50	32.08263	
GENEX Option C	100000	Non-coastal	3 or 4	50	5.80639	
GENEX Option C	100000	Non-coastal	3 or 4	50	14.05157	
GENEX Option C	250000	Non-coastal	3 or 4	50	2.500361	
GENEX Option C	250000	Non-coastal	3 or 4	50	13.67417	
GENEX Option C	100000	Non-coastal	3 or 4	50	0.164711	
GENEX Option C	100000	Non-coastal	3 or 4	50	3.009753	1379 Total SO 3 or 4
GENEX Option C		Non-coastal	5 or greater	100		
GENEX Option C		Non-coastal	5 or greater	100		
GENEX Option C		Non-coastal	5 or greater	100		
GENEX Option C		Non-coastal	5 or greater	100	300.0754	
GENEX Option C		Non-coastal	5 or greater	100		
GENEX Option C		Non-coastal	5 or greater	100	5.80639	
GENEX Option C		Non-coastal	5 or greater	100		
GENEX Option C		Non-coastal	5 or greater		0.164711	
GENEX Option C		Non-coastal	5 or greater		1.857839	603 Total SO 5 or greater
denex option o	100000	Non coustai	o or greater	100	1.007007	
Name	Scale	Bioregion_type	SO_group	Buffer_m	Area_ha	
GENEX Option C		Coastal	1 or 2	10		
GENEX Option C		Coastal	1 or 2	10		
GENEX Option C		Coastal	1 or 2	10		
GENEX Option C		Coastal	1 or 2	10		
GENEX Option C		Coastal	1 or 2		18.08111	
GENEX Option C		Coastal	1 or 2	10		23 Total SO 1 or 2
						23/10/01/30/10/2
GENEX Option C		Coastal	3 or 4	25		
GENEX Option C		Coastal	3 or 4	25		
GENEX Option C		Coastal	3 or 4	25		
GENEX Option C	100000	Coastal	3 or 4	25	0.545677	22 Total SO 3 or 4

### Kidston Corridor Options - Regulated remnant wetland vegetation

Name	RE	PERCENT	VM_POLY	Area_ha	
Genex: Option A	9.11.15a	100	LeastC	1.926624	
Genex: Option A	9.3.25	100	LeastC	0.175368	
Genex: Option A	9.3.19a/9.3.3a/9.3.23	70/20/10	O-subdom	2.611455	4.7 Total remnant wetland vegetation for Option A
Genex: Option B	9.3.26	100	LeastC	3.298466	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	7.444698	
Genex: Option B	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	3.873757	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	3.716969	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	3.716969	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	0.039454	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	0.039454	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	6.792229	
Genex: Option B	9.8.1a/9.8.13	80/20	LeastC	6.792229	
Genex: Option B	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	0.692628	
Genex: Option B	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	0.692628	
Genex: Option B	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	2.587155	
Genex: Option B	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	2.587155	42.3 Total remnant wetland vegetation for Option B
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	3.716969	
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	3.716969	
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	0.039454	
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	0.039454	
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	6.792229	
GENEX Option C	9.8.1a/9.8.13	80/20	LeastC	6.792229	
GENEX Option C	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	0.692628	
GENEX Option C	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	0.692628	
GENEX Option C	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	2.587155	
GENEX Option C	9.8.13/9.8.1a/9.8.1b/9.3.10b	60/20/15/5	LeastC	2.587155	27.7 Total remnant wetland vegetation for Option C

### Kidston Corridor Options - MSES Protected areas

Name	M_Table1_1	M1_1	M1_1_Type	M1_1_Name	Area_ha	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Newcastle Range-The Oaks Nature Refuge	439.8409	439.8 Newcastle Range-The Oaks Nature Refuge
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	1.316996	4.7 Liefway Nature Refuge
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	0.323124	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	36.85525	213.7 Girringun National Park
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	0.000382	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	1.478203	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	28.1685	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	4.236403	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	1.381377	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	2.078499	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	0.414249	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	0.548143	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	117.5568	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	18.62776	
Genex: Option A	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	5.381855	658.2 Total hectares for Option A
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	36.85525	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	0.000382	214.4 Girringun National Park
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	1.478203	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	28.1685	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	4.236403	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	1.381377	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	0.719896	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	117.5568	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	18.62776	
Genex: Option B	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	5.381855	214.4 Total hectares for Option B
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	2.078499	
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	0.414249	3.0 Liefway Nature Refuge
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NR	Liefway Nature Refuge	0.548143	
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	0.719896	142.3 Girringun National Park
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	117.5568	
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	18.62776	
GENEX Option C	STATE CONSERVATION AREAS	Protected Areas	NP	Girringun National Park	5.381855	145.3 Total hectares for Option C

#### Kidston Corridor Options - MSES Wildlife habitat

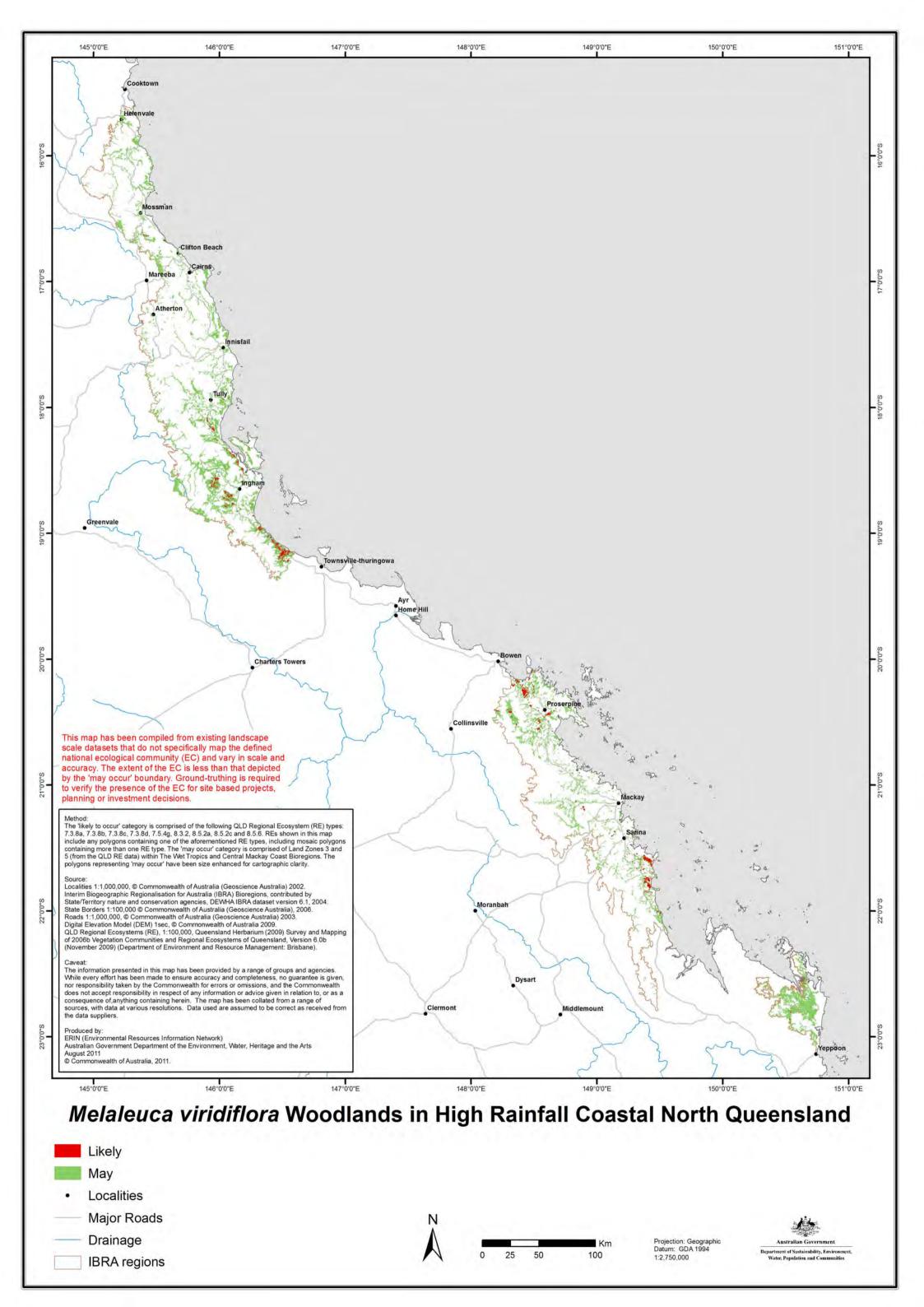
Nomo	M3 1	M2 1 SpocA	MO 1 Faallh	Aroo he
Name Genex: Option A	Threat wildlife & Spec LeastC animals	M3_1_SpecA Acacia tingoorensis, Acacia tingoorensis	<pre>M3_1_EssHb </pre>	Area_ha 0.000635
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	<null></null>	0.000012
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	<null></null>	0.15312
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.187874
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.000084
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.00069
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	34.16968
enex: Option A enex: Option A	Threat wildlife & Spec LeastC animals Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis Acacia tingoorensis, Acacia tingoorensis	Essential <null></null>	163.847 0.016774
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	<null></null>	0.010774
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	<null></null>	0.115602
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	<null></null>	0.00036
Genex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.72007
Genex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	17.55466
Genex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.623486
Genex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.15235
enex: Option A	Threat wildlife & Spec LeastC animals Threat wildlife & Spec LeastC animals	<null></null>	Essential Essential	0.000056
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.02391
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.063902
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.161254
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.23368
enex: Option A	Threat wildlife & Spec LeastC animals	<null></null>	Essential	0.402452
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	7.455134
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	11.393
Senex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential Essential	3.3685
enex: Option A	Threat wildlife & Spec LeastC animals Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis Acacia tingoorensis, Acacia tingoorensis	Essential	1.962138
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	0.280448
enex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	1.301988
Senex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	14.64406
Genex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	6.153846
Genex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	3.000527
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Genex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	9.905173
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Genex: Option A	Threat wildlife & Spec LeastC animals	Acacia tingoorensis, Acacia tingoorensis	Essential	0.72055
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Genex: Option B	Threat wildlife & Spec LeastC animals	<null></null>	Essential	1.70562
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GENEX Option C GENEX Option C	Threat wildlife & Spec LeastC animals Threat wildlife & Spec LeastC animals	<null></null>	Essential Essential	15.50554

Kidston Corridor Options - HES Wetlands

Name	M_Table1_2	M2_1	M2_1_Des	M2_1_Dat	Area_ha	
Genex: Option B	WETLANDS AND WATERWAYS	HES wetlands	WPA wetland	wetland protection area-wetland	1.127498	28 1.1 Total hectares for Option B

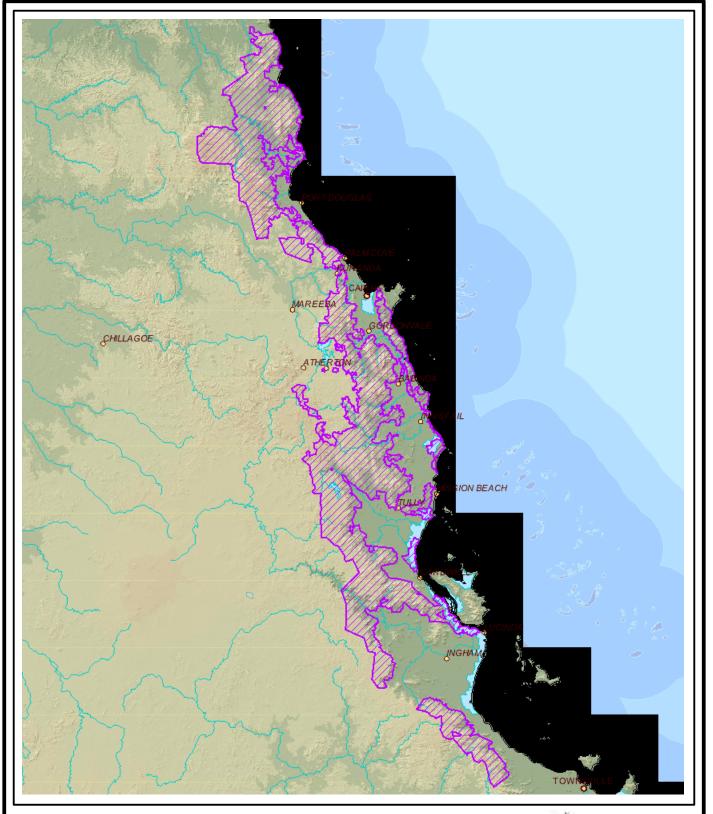
# Appendix H

**TEC DISTRIBUTION MAP** 

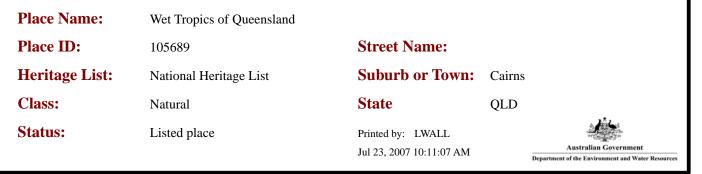


## Appendix I

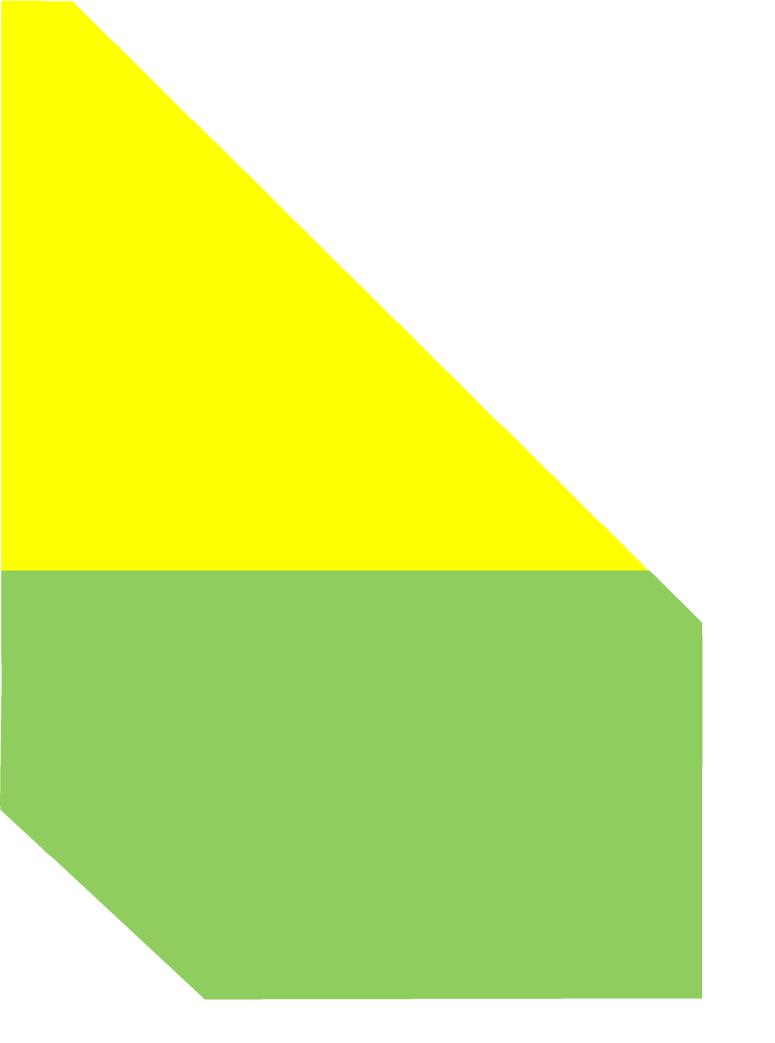
WORLD HERITAGE PROPERTIES



## **Place Details**



Scale 1:2000000

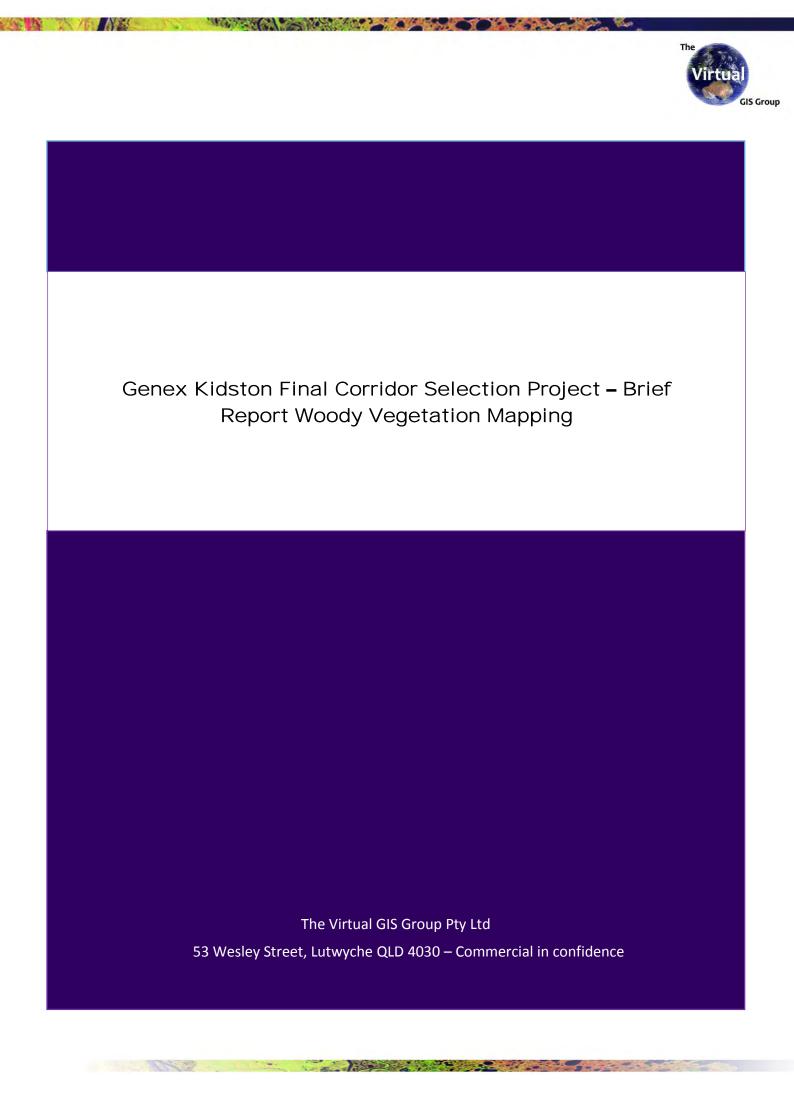






Attachment 2 – Remote Sensing Report

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### Introduction

Powerlink have been commissioned to provide final corridor options for the Genex Kidston upgrade. To assist the assessment of potential ecological and environmental relative to the three options being considered, the extent of woody vegetation needed to be mapped. This report summarises the process to map woody vegetation for the Genex Kidston project.

#### Overview

The purpose of this project was to map the extent of woody vegetation using best available broad scale satellite imagery to assist in subsequent assessment of the ecological value of remnant vegetation near three potential corridor routes.

The work conducted included a review and acquisition of freely available satellite imagery to form the basis of the desktop mapping of woody vegetation. Analysis was conducted using Landsat multispectral satellite imagery at a 30-metre spatial resolution.

The project provides a high-level overview of current woody vegetation cover in relation to existing State mapping, highlighting areas where current mapping may be improved through further investigation. Areas that could be regrowth due to absence of vegetated cover around 15 years ago together with areas where the distribution of canopy could indicate poor condition in existing remnant areas were mapped.

The results of this mapping will be used to assist Parsons Brinckerhoff ecological consultants to produce a desktop assessment of the ecological value of remnant vegetation in and near the proposed easement options and to inform Powerlink of any potential issues in relation to the EPBC Act reporting obligations.

### **Project Extent**

The area of interest is presented in Figure 1. The extent of the project was derived from a one kilometre buffer around the external limits of the provided corridor options, which start in the east near Mount Fox and run westwards for approximately 185 kilometres through the Greenvale and Conjuboy districts in North Queensland. The project extent covers an area of approximately 1,520 km<sup>2</sup>. The project is in a region of relatively dry, scrubby country. The current regional ecosystem mapping indicates that the project extends across bioregions 9 (Einasleigh Uplands) and 7 (Wet Tropics), with most of the area west of Mt Fox falling into the former bioregion. The area is bisected by waterways which tend to run north-south across the project extent and dominant landzones are 3, 5 and 11. Landzone 11 are hills and lowlands on metamorphic rocks. Landzone 5 represents areas of tertiary-early Quaternary loamy and sandy plains and plateaus. Landzone 3 incorporates recent Quaternary alluvial systems along river and creek flats. Landzones 7,8, 10 and 12 are also present across the project extent.

The current regional ecosystem mapping indicates that the project extent, away from the waterways, is dominated by vegetation communities with a scattered canopy. When mapping at a broad scale, this presents as vegetation with low greenness, particularly when the groundcover is very dry grassland. Different images were used to assist in the mapping, particularly to consider changes reflecting the moisture regime at the time of image acquisition.

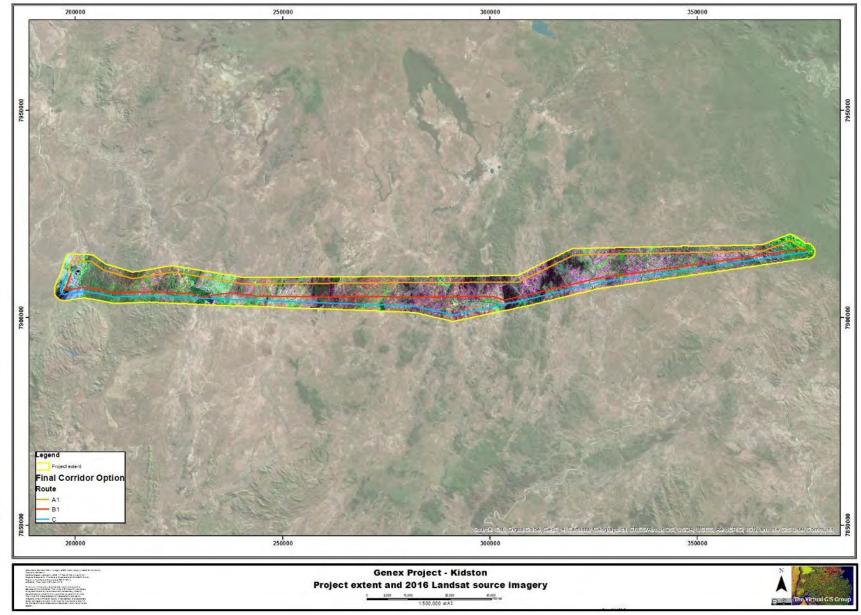


Figure 1: GENEX Kidston project extent and 2016 Landsat source imagery









## Source data

## Remotely sensed data

This project requires the use of remotely sensed imagery to detect and map woody vegetation across the project extent. Remotely sensed data can be sourced from satellite imagery and aerial photography and the suitability for this project is determined by availability of access, frequency of capture, distribution of cloud cover, availability of legacy or archive data for temporal assessment.

As the project extended over an area of approximately 1,520 square kilometres, a remotely sensed solution based on aerial photography was not considered and satellite imagery was used.

## Satellite imagery

Satellite imagery is widely available from a range of international custodians. Most commercial operators charge on an area coverage basis. No budget had been allocated for the purchase of satellite imagery, so available commercial satellites who charge for their data were not considered.

### Availability of access

Access was determined for this project by cost and timeframe for the supply. Due to the extent of the project and the need for the project to commence immediately, multi-temporal Landsat data is free to download from the NASA/USGS website and provides broad scale imagery at 30-metre spatial resolution with multispectral coverage. Landsat imagery was sourced from the United States Geological Society (USGS). Two image scenes cover the project extent from Path/Row 96/73 and 95/73.

## **Frequency of capture**

Both Landsat 8 and 7 are currently operational. Landsat 8 is the preferred satellite and has a repeat cycle over the project extent every 16 days. Landsat 7 provides optional infill coverage and legacy coverage from early 2000s.

## **Distribution of cloud cover**

Although the repeat cycle is 16 days for each satellite, the suitability of the imagery is determined by the presence and extent of cloud over the project extent. To select suitable imagery, each capture was reviewed in relation to the GENEX Kidston corridor options to identify which recent scenes would be suit the assessment of woody vegetation.

### Availability of legacy or archive satellite imagery

As the Landsat programme has been operational since 1972, legacy data from 2000-2002 was acquired to provide an overview on whether clearing or cultivation had occurred in the past 15 years within the project area. Landsat is the only programme which provides free consistent legacy data at the same spatial resolution over such an extended period.

Suitable cloud-free imagery was available during 2016, to provide a current view on the extent and spatial distribution of the woody vegetation. Additional Landsat scenes for each tile were used to compare different moisture conditions and to infill the 96/73 scene which did include a small amount of cloud in the far western region. Only the primary scene across the full corridor was analysed, to keep costs to a minimum. Legacy scenes were acquired to provide a baseline reference for determining if the vegetated cover is likely to be remnant. Although this is an extension to mapping woody vegetation, it does contribute to the overall assessment of the ecological value of the mapped areas.







The images acquired for the GENEX Kidston assessment are listed in Table 1:

Landsat scene (Path/Row)	Primary scene date	Reference scene date	Legacy scene date
96/73 (western)	17 Feb 2016	28 Aug 2016	23 July 2000
95/73 (eastern)	5 August 2016	17 July 2015	5 July 2002

Table 1: Landsat scenes acquired	for the GENEX woody vegetation assessment

Standard ortho-ready image products were used which contain some variability due to terrain distortions but are usually within 1 pixel accuracy in the horizontal. Where available, surface reflectance imagery was acquired to mitigate the impact of differing atmospheric conditions. This was only available for some sourced imagery and not for all required source data. Adjustments in the methodology were deployed to compensate for these differences. The imagery was therefore not pre-processed, relying on the geometric and radiometric accuracies of the source data. The imagery is suitable for analysis to detect and map woody vegetation at a broad scale.

The methodology used to map the woody vegetation can be deployed using any multispectral source imagery if budget or finer resolution imagery becomes available once the final corridor selection has been made.

## Other geospatial datasets

Reference geospatial datasets were sourced to expedite the mapping of woody vegetation. These included:

- Remnant vegetation over the Einasleigh bioregion
- Remnant vegetation over the North-west bioregion
- Regrowth vegetation over the scene extent
- Queensland 1:25,000 drainage
- Queensland Major rivers
- Queensland Statewide landcover and tree study (2000 to 2015) (SLATS)

Other spatial datasets were also generated using finer resolution backdrop only imagery. These data, which were digitised from the screen, included:

- Built structures
- Linear easements, including roads, tracks, powerlines and fences
- Dams
- Waterways

These additional datasets were specifically designed to be aligned with the source imagery and to assist in the assessment of woody vegetation.

SLATS data was aggregated across the 2000 to 2015 assessment period, with all clearance types except damage due to natural disaster being flagged as non-remnant (1,370 sites).

All project-related datasets were prepared and reprojected into GDA 1994 Zone 55.

## Methodology

The methodology deployed captured the extent and spatial distribution of current woody vegetation using an object based image classification process. This methodology has been previously deployed



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at a wide range of scales and using different remotely sensed source datasets for organisations across Australia and internationally.

The scale of the source imagery determined the granularity achievable for mapping woody vegetation. Given the type of vegetation present within the project extent, woody vegetation was mapped using an approach that enabled the separate discrimination of areas presenting a range of greenness indicators.

The methodology was based on an eCognition Geographic Object Based Image Analysis (GEOBIA) approach which isolated undisturbed, woody areas from other landcovers. Using GEOBIA enabled the use of neighbourhood functions to isolate features, which is critical when discriminating between spectrally similar regions which are contextually different.

### 2016 vegetation cover mapping

State Government datasets, both existing remnant and regrowth datasets, were used to inform the process, which identified areas which were consistent with current remnant and regrowth vegetation cover mapping. Areas mapped as cleared in the SLATS mapping program from 2000 to 2015 were also attributed as non-remnant, except where those areas were specifically noted as being subject to natural disaster damage in any given year.

Additionally, those areas which appeared to be very scattered with lower levels of vegetated cover were separately identified as potential woody vegetation. The remnant areas were grouped by their VMA status in State Government regional ecosystem mapping and attributed as either remnant or potentially remnant. This interim dataset was used as part of the process to inform woody vegetation mapping and was not included as one of the final deliverables.

Areas where the imagery pattern appeared to be consistent with other areas that have been verified as non-remnant and were adjacent to those areas were mapped as non-remnant. Areas which did not appear to have consistency in the imagery with other areas mapped as remnant were flagged as potential remnant. Some areas of fire scarring were detected but were not separately mapped.

Objects that were identified as nonvegetated 15 years ago but now mapped as woody vegetation were separated from those which were consistently mapped as woody vegetation, to separate potential regrowth zones from remnant and potential remnant areas.

The methodology adopted thresholds to assign classes into remnant, potential remnant or scattered using the spectral characteristics in the Landsat imagery. This was refined through visual reference to the larger scale reference imagery presented above.

### 2016 woody vegetation mapping

Within the project extent, landcover was mapped to separately identify remnant vegetation cover, from which woody vegetation was aggregated.

The assessment sequence was consistent with the method described by (Neldner, Wilson, E.J.Thompson, & Dillewaard, 2012) for vegetation assessment from imagery only and included an inspection of the legacy scenes across each tile to check for baseline disturbance and/or cultivation in the last 15 years.

Minor variations in vegetation cover were not captured due to the scale of the imagery and the minimum mappable area associated with the spatial resolution of the source Landsat imagery.







The mapping products generated within the GEOBIA environment were exported into vector shapefiles for delivery.

## **Results and discussion**

The Landsat satellite imagery was used to derive landcover classification for the project extent from which indicative regional ecosystem mapping was generated. This was then aggregated into the mapping of woody vegetation for the project.

Although the methodology was consistent with that implemented in other major infrastructure projects in central Queensland, the lack of availability of high resolution multispectral imagery limited the ability to reliably discriminate between non-vegetated and dry, scrubby vegetation.

To highlight these limitations, the following set of examples have been generated to demonstrate the resolvable landcover using Landsat data compared to higher resolution imagery. Table 2 contains a set of screen snapshots over the project area captured at a range of viewing scales. The left-hand column contains the Landsat imagery used in this project, while the right-hand column contains examples obtained from the view-only world imagery available through the ESRI ArcGIS online web service.

The examples used include scales from the recommended publication scale of 1:80,000 for Landsat 7 and 8 at 30-metre resolution (Geoimage, 2016) and demonstrate how the reliability of mapping woody vegetation using Landsat imagery only will deteriorate as the scale of mapping required becomes larger. In the Landsat imagery, the spectral bands displayed are 7,5,2 to RGB. Healthy vegetation appears green whereas areas dominated by exposed soil or very dry grasses appear light pink. Woody vegetation with low chlorophyll levels present as dark magenta. In the alternative, higher spatial resolution true-colour reference data, the heavily vegetated areas appear darker than the surrounding grassland and exposed soils.

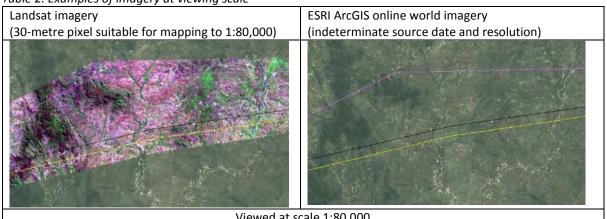
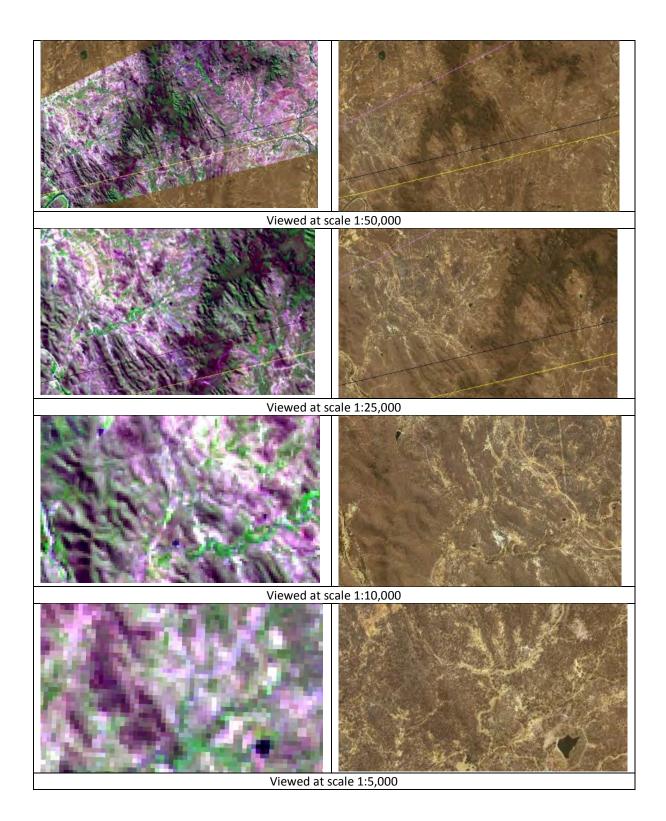


Table 2: Examples of imagery at viewing scale

Viewed at scale 1:80,000







As the viewing scale gets larger, the differences in ability to discriminate woody vegetation from non-vegetated areas becomes harder. The detail available in the ESRI data clearly shows vegetation whereas the Landsat pixels are dominated by the groundcover and vegetation contain low chlorophyll levels rather than highly reflective woody vegetation.





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This is partly due to the spatial resolution but also to the leaf composition of the dominant vegetation communities on all landzones except landzone 3. This is evident in the example at 1:5,000 scale imagery, where only the riparian corridor appears as healthy, green vegetation and the upland areas expected to be dominated by regional ecosystem 9.7.1 and 9.11.5 are not reflecting well in the near infrared part of the spectrum compared to the red channel. This results in low vegetation index levels which usually present across regions which are not vegetated. The two regional ecosystems expected in this small area in the 1:5000 scale example are both sparse structurally. 9.7.1 community contains Eucalyptus persistens woodland on lateritised and deeply weathered surfaces on undulating terrain. 9.11.5 contains Eucalyptus persistens +/- E. crebra woodland on low metamorphic hills. E. crebra is commonly known as narrow leaf ironbark, which goes some way to explaining why the leaves are not reflecting well in the multispectral imagery.

The data presented in Table 3 demonstrates how Landsat is fit for purpose when mapping woody vegetation at a broad scale and how the quality of the result reduces as the viewing scale is increased. Landsat supports mapping typically to 1:80,000 so even the examples shown in this table exceed the suitability of the data for discriminating vegetation cover. In saying that, it is evident that the results are differentiating between areas that are dominated by vegetation and areas which experience higher levels of soil erosion gullies and other disturbances.

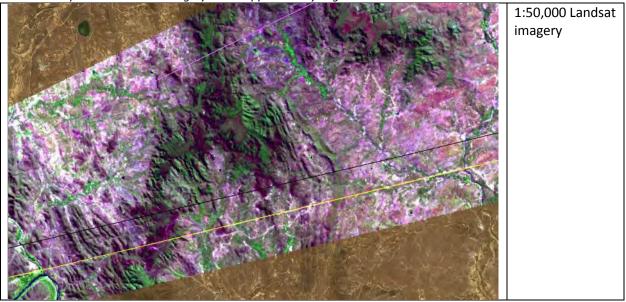


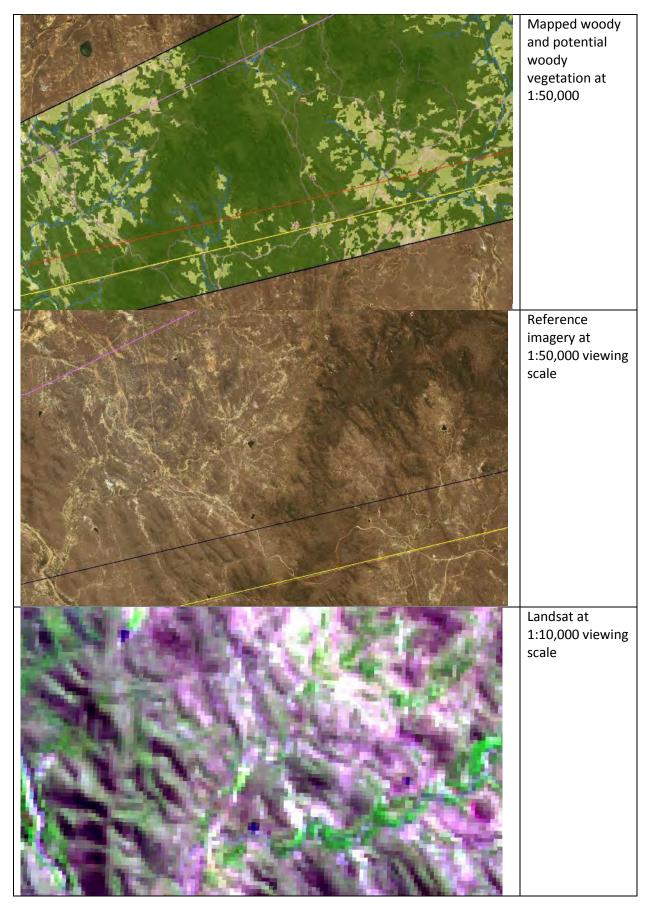
Table 3: Comparison between imagery and mapped woody vegetation at scale







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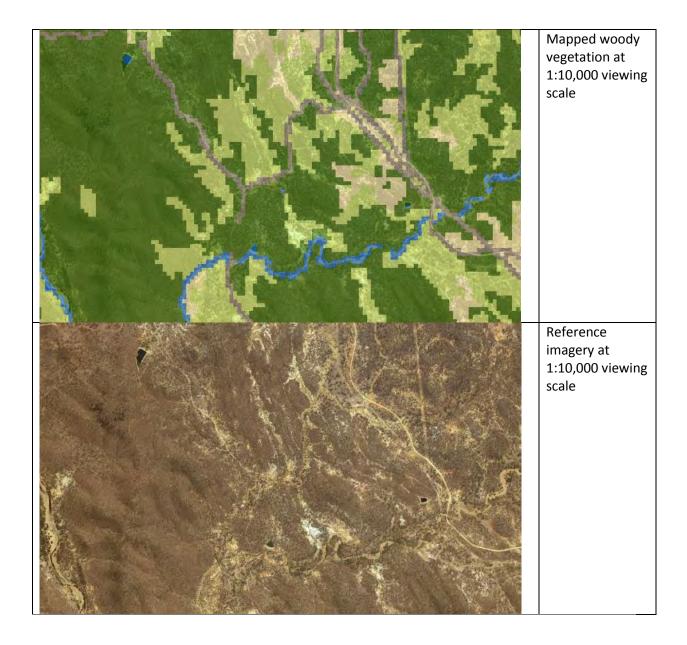
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The results generated for this study must be applied at a scale which is fit for purpose. This is 1:80,000 or at best 1:50,000 when considering woody vegetation only.

Please note that the data captured from the ESRI web services cannot be used for analytical purposes and therefore can only support the project through visual verification.

Landcover mapping was undertaken initially to map the distribution of vegetation cover across the project extent. This was conducted in two steps: 2016 vegetation cover mapping followed by 2016 woody vegetation mapping. These steps were performed independently of the SLATS clearance mapping, presented in Figure 2. A large area in the central-western section of the project extent was impacted by natural disaster damage during the 2010-20111 assessment period covering an area of 1,178 hectares. This is presented as cyan in Figure 2. This was not assigned as non-remnant due to the type of clearance mapped.



Of the remaining area mapped by the SLATS program as being cleared between 2000 and 2015, only 678 hectares were identified. Where these areas were mapped as remnant in the vegetation cover and the woody vegetation mapping, they were recorded as areas of non-remnant in the metrics and mapping, except where they fell below the minimum mapping area in size.

## 2016 vegetation cover mapping

Due to the nature of the distributed and scattered vegetation across most of the project extent, the aggregation of vegetation with very low vegetation greenness indices was incrementally applied to bring dry, scrubby areas into the mapped remnant category. Given the characteristics of the vegetation, some areas remained classified as potential remnant due to the scattered, dry or grassland dominated nature of those communities. These differences in vegetation cover density were provided to assist the PB ecologists review the remnant vegetation status and were generated without further field verification or input from an ecological perspective. Results are shown in Figure 3 and metrics reported in Table 4.

The SLATS data shown in Figure 2 was used to reassign the areas mapped as potential remnant to adjusted remnant where they had been impacted by disturbance caused by natural disaster damage.

Some minor areas of disturbance were observed in the 2000-2002 images which were initially mapped as remnant. These were reassigned potential regrowth accordingly. No evidence of cultivation was detected other than plantations already mapped as non-remnant in the eastern section of the project extent.

The classes of vegetation mapped are listed in Table 4, which were used to aggregate up into woody vegetation.

## 2016 woody vegetation mapping

Using the derived regional ecosystem mapping, the woody vegetation was aggregated into the class groupings described in Table 4 and the results are mapped in Figure 4. The groupings combine all remnant classes into the Woody vegetation class and the Scattered vegetation classes described in the table.

Woody veg class grouping	Woody veg area (ha) within project extent (after MMU removal)	RE class grouping	Class description	RE area (ha) within project extent (before MMU removal)
Woody vegetation	108503	Remnant O-dom	Woody vegetation spatially coincident with Of concern dominant REs including lower and higher vegetation indices	2336
		O-subdom	Woody vegetation spatially coincident with Of concern sub-dominant REs including lower and higher vegetation indices	1531
		LeastC	Woody vegetation spatially coincident with Least Concern REs including lower and higher vegetation indices	101550

#### Table 4: Landcover classes used to map woody vegetation







Scattered vegetation	23818	Scattered or dry O-dom	Scattered or dry dominated areas spatially coincident with Of concern dominant REs	769
		Scattered or dry O-subdom	Scattered or dry dominated areas spatially coincident with Of concern sub-dominant REs	731
		Scattered or dry LeastC	Scattered or dry dominated areas spatially coincident with Least concern dominant REs	24163
Regrowth	500	Regrowth	Areas mapped by State Government as regrowth (2014)	476
Potential regrowth	245	Potential regrowth	Areas mapped a remnant but which did not appear to contain woody vegetation in 2000 and 2002 depending on the image scene referenced	240
Non- remnant	3888	Non-remnant	Areas mapped by State Government as non-remnant	3836
Non- vegetated	12129	Bare	Areas with no or very low greenness considered to comprise very dry grassland, bare, exposed soil, cleared or eroded areas	4117
		Built structures	Buildings, sheds, other structures	8
		Easements	Including roads, tracks, power easements and fence lines	5927
		Water	Dams and waterways	3399
Total	149084			149084

The results of this mapping are presented in Figure 3 and Figure 4.

Minimum mapping area tolerances were applied to retain the integrity of the results, consistent with the source data used to generate the outcomes. Mapping products have been generated at 1:500,000 at A3 to cover the full proposed project extent and support the digital data supplied as a deliverable.

Regional ecosystem RE\_Label attribution has been retained on the resultant digital datasets, with merging of class groups being limited by the RE extents.

Field verification will support the refinement and improvement of outcomes but does need to be aligned with the source imagery. This requires field-based observations to be aggregated based on dominant cover over a sample extent equivalent to the MMU for the imagery. Field-based observations were not available in support of this desktop-only project.







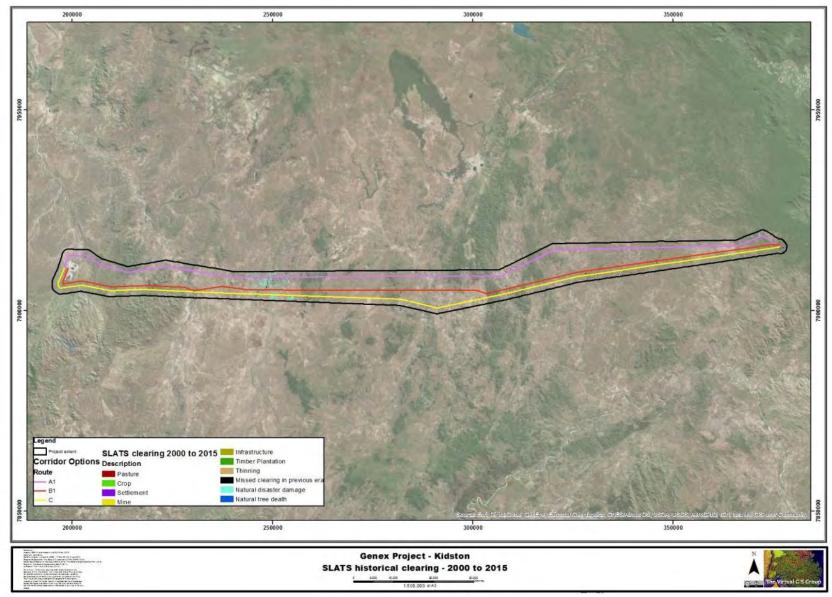


Figure 2: Statewide Landcover and Tree Study clearance 2000 to 2015









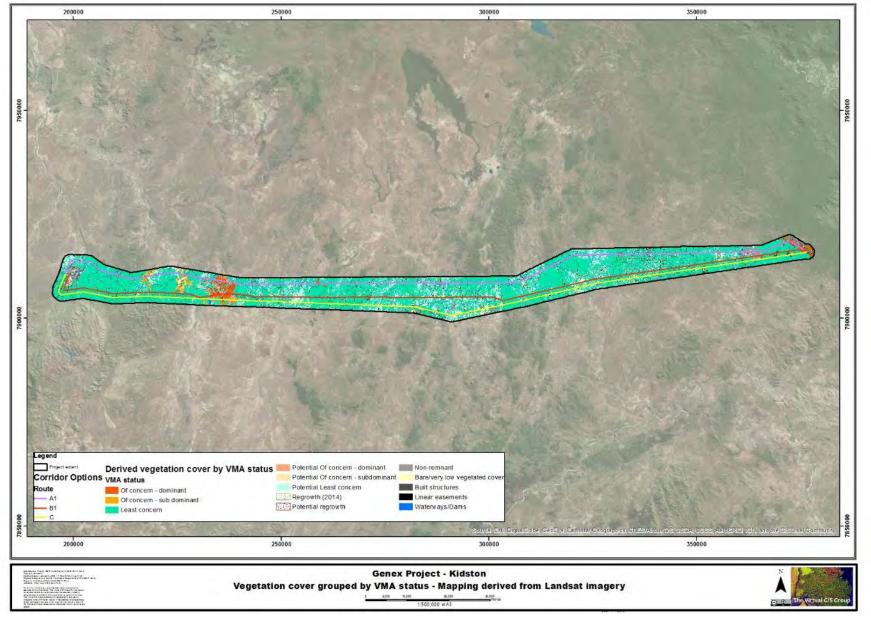


Figure 3: Derived vegetation cover symbolised by VMA status









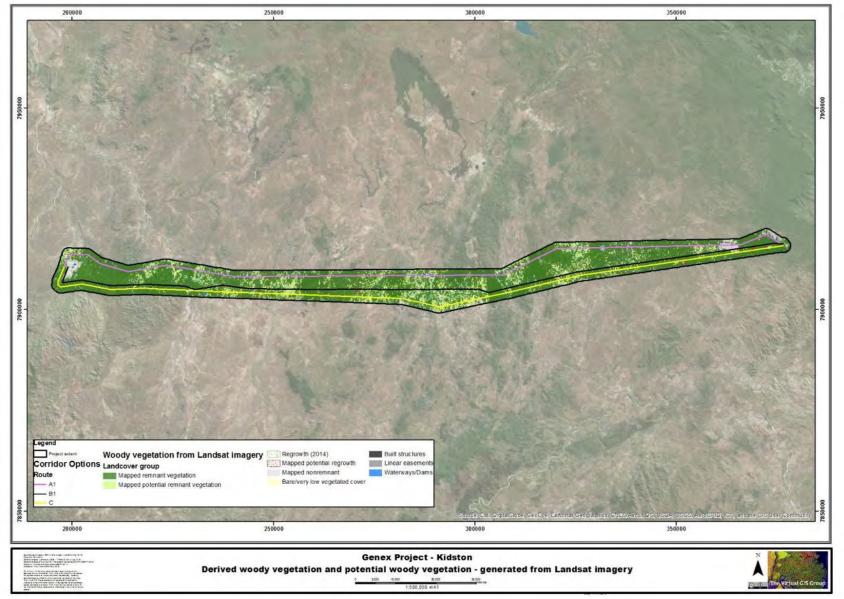


Figure 4: Derived woody vegetation and potential woody vegetation











## **Quality Assurance**

As with most remotely sensed approaches to mapping landcover, accuracy of results increases with careful review of the results, and where necessary, the manual improvement of outcomes in areas of mixed pixels and/or mixed landcovers. Although the process does not require extensive manual intervention, some of the targets in this project are expected to be below the minimum mapping unit in size and so will be eliminated through quality control. Where the integrity of the data reduces with the removal of MMU areas, such as around waterways, easements and adjacent to areas mapped from State Government data, these areas may be retained. Dam, buildings, easements and waterways are not dissolved by MMU due to being point or linear features.

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Formal accuracy assessment was not possible due to lack of field-based documentation but the results should be suitable to assist the assessment of potential ecological and environmental relative to the three options being considered.

Quality Assurance will be performed throughout the project. These functions will include geometric and attribute tests to ensure the data is clean and suitable for delivery. Standard checks for geometric errors such as undershoots, overshoots, closed polygons were performed as well as checking for duplicate objects.

To meet the timeframe and budgetary limitations associated with this project, a series of limitations, assumptions and exclusions were identified at the proposal stage and these were consistent with the project outcomes. Please refer to the proposal for a summary of these.

## **Future options**

Further benefit could be gained by conducting analytics using both the current Landsat imagery and legacy imagery from 2000 and 2002, to be able to map change in vegetation cover.

Additionally, finer resolution satellite imagery would enable more detailed mapping of areas which could be considered non-remnant based on vegetation cover and density which may be inconsistent with the expectations of the regional ecosystems mapped by the State Government.

As with any desktop solution, particularly one which has relied on broad scale imagery without the benefit of field verification data, any results could be improved with the acquisition of field data confirming the vegetation cover, dominant regional ecosystem and woody vegetation status.

## REFERENCES

Geoimage. (2016, November 11). *Landsat 7 specifications*. Retrieved from Geoimage website: http://www.geoimage.com.au/satellite/landsat7

Neldner, V., Wilson, B., E.J.Thompson, & Dillewaard, H. (2012). *Methodology for Survey and Mapping of Regional Ecosystems and Vegetation Communities in Queensland.* Brisbane: State of Queensland.



Version: 6.0

Attachment 3 – Likelihood of Occurrence Assessment

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POWERLINK

## Preliminary Desktop-Based Likelihood of Occurrence Assessment

KIDSTON POWER TRANSMISSION LINE

PUBLIC

FEBRUARY 2017



## Preliminary Desktop-Based Likelihood of Occurrence Assessment

**KIDSTON POWER TRANSMISSION LINE** 

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#### Public

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## 1 INTRODUCTION

Powerlink is investigating corridor options for a powerline to connect a renewable energy generator (Genex) at Kidston in Far North Queensland, to the electricity transmission grid at Mount Fox, approximately 200 km east of Kidston (the Project). The site at Kidston will contain a mix of solar and pumped storage (hydro) technology.

The connection point to Kidston is at Mount Fox, approximately 200 km east of Kidston. It is most likely that a 275 kV transmission line will be required to provide adequate capacity for the connection distance. Powerlink was commissioned by Genex to produce a Corridor Selection Report, to identify a preferred corridor option from a broad study area, and a preliminary alignment for the connection. As part of this work an Initial Desktop Assessment and Preliminary Ecological Constraints Advice report was prepared for Powerlink. It identified the Matters of National Environmental Significance (MNES) and Matters of State Environmental Significance (MSES) of relevance to each corridor option.

The Project is located in a region of Queensland that is relatively remote and where very little to no ecological studies have been conducted. This was apparent by the low number of threatened species records revealed by Wildlife Online database searches. The *Environment Protection and Biodiversity Conservation Act 1999* (EPBC Act) Protected Matters Search Tool also identified a number of threatened species predicted with potential to occur in the corridor options.

Subsequent to this work, the Department of Environment and Heritage Protection (DEHP) suggested the provision of a likelihood of occurrence assessment of threatened flora and fauna species listed under the *Nature Conservation Act 1994* (NC Act), to identify those species that maybe of relevance to the corridor options. This Preliminary Desktop-based Likelihood of Occurrence Assessment has been prepared for Powerlink in response to this request. To ensure completeness, this assessment also includes a likelihood of occurrence assessment for threatened flora and fauna species and threatened ecological communities listed under the EPBC Act.

## 1.1 Purpose

The study area for the Project is located in a region where little to no ecological studies have been undertaken and as a result records for threatened species are limited.

In support of the Initial Desktop Assessment and Preliminary Ecological Constraints Advice, this Preliminary Desktop-based Likelihood of Occurrence Assessment identifies the threatened species and ecological communities listed under the NC Act and/or EPBC Act that may potentially occur in the corridor options, for which future ecological surveys can be targeted. Essentially, it helps fill the gap in information resulting from the lack of species records in the study area.

The findings of this Preliminary Desktop-based Likelihood of Occurrence Assessment will also be useful in terms of identifying the threatened species and ecological communities that may be subject to project related impacts.

## 1.2 Study limitations

The likelihood of occurrence assessment presented herein is based upon a desktop assessment only, no field surveys have been undertaken to verify the presence and extent of the broad vegetation groups, regional ecosystems and/or fauna habitats within the corridor options.

It is a high level assessment, and on that basis the assessment has been conducted using a conservative approach. Therefore, the subsequent number of species with a moderate or high likelihood of occurrence is higher than the number that would likely result if field verification surveys of vegetation and habitats had been performed.

This assessment has relied on database searches conducted to the extent of local government area (LGA) boundaries, and State published broad vegetation group (BVG) mapping to predict the extent of potential habitats that may be supporting threatened species.

This likelihood of occurrence assessment will be updated at the impact assessment stage of the Project as field studies occur, with the potential that species may be both removed and added to the assessment.

## 2 METHODS

The methods used to inform the desktop-based likelihood of occurrence assessment have involved database searches and an assessment against publically available information to inform the habitat types that may potentially support threatened species.

The database searches included of the Protected Matters Search Tool and Wildlife Online covering the local government area (LGA) searches for the following LGAs:

- → Hinchinbrook Shire Council
- → Charters Towers Regional Council
- → Etheridge Shire Council.

Because of the lack of records obtained from the search areas used for the Initial Desktop Assessment and Preliminary Ecological Constraints Advice, LGA wide database searches have been conducted to identify the species that may occur within the vegetation and habitats associated with the corridor options.

The publically available information included:

- → State Department of Natural Resources and Mines regional ecosystem mapping (Version 8) to identify Broad Vegetation Groups (BVGs) for assessing the likelihood of occurrence.
- > Species or ecological community information (i.e. habitat requirements and distribution) from:
  - State Department of Science, Information Technology and Innovation's Species profile search
  - Commonwealth Government administered Species Profile (SPRAT)
  - Commonwealth Government administered Atlas of Living Australia
  - Royal Botanic Gardens NSW administered PlantNET.

## 2.1 Determination of potential habitats within the corridor options

The regional ecosystem mapping associated with the corridor options was interrogated to identify the BVGs of relevance to the corridor options, as presented in Appendix A. Given the current scale of the project and associated desktop assessment, the BVGs are an effective way to identify the type of habitats available for threatened flora and fauna species.

The assessment for threatened flora species habitats was determined in reference to the 1: 1,000,000 scale BVG groups and sub-types associated with the corridor options. This approach accounts for the vegetation community associations and geology that is required for the threatened flora species to potentially occur.

The assessment for threatened fauna species is also determined in reference to the 1: 1,000,000 scale BVG groups with the corridor options, but only in the context of vegetation structural types (e.g. open forest, tussock grasslands, woodlands etc.). BVG sub-types are only used if the species is a specialist and very habitat specific (e.g. Green-eyed Tree Frog (*Litoria serrata*) only found in rainforest).

## 2.2 Assessment criteria

The likelihood of occurrence for threatened species and ecological communities listed under the EPBC Act and/or NC Act, has been assessed against the criteria outlined below in Table 2.1.

<u>Note:</u> In the absence of publically available species and habitat information, a conservative ranking of '*Moderate*' is applied. This is only relevant to some NC Act listed flora species revealed by database searches and does not apply to EPBC Act listed flora or fauna. 3

#### Table 2.1 Likelihood of occurrence assessment criteria

LIKELIHOOD OF OCCURRENCE	ASSESSMENT CRITERIA
Low	<ul> <li>One or more of the following criteria is met:</li> <li>→ suitable habitat unlikely to be present in the corridor options</li> <li>→ species considered extinct in the wild.</li> </ul>
Moderate	<ul> <li>The following criteria is met:</li> <li>→ suitable habitat potentially present in the corridor options.</li> </ul>
High	<ul> <li>The following criteria is met:</li> <li>→ suitable habitat likely to be present in the corridor options area</li> <li>→ previous records in study area.</li> </ul>

# 3 SUMMARY OF LIKELIHOOD OF OCCURRENCE

The threatened flora and fauna species and migratory fauna species that are of moderate or high likelihood of occurrence in the corridor options are presented in the following sections.

None of the threatened ecological communities revealed by database searches are likely to occur in the corridor options. This low likelihood of occurrence is primarily on the basis of the bioregions they are restricted to under the EPBC Act, or because they are only found in near coastal environments:

- Brigalow (Acacia harpophylla dominant and codominant) restricted to Brigalow Belt and Mulga Lands bioregions
- Broad leaf tea-tree (*Melaleuca viridiflora*) woodlands in high rainfall coastal north Queensland restricted to coastal floodplains of the wet tropics bioregion
- Semi-evergreen vine thickets of the Brigalow Belt (North and South) and Nandewar Bioregions restricted to the Brigalow Belt and Nandewar bioregions
- → Littoral Rainforest and Coastal Vine Thickets of Eastern Australia restricted to coastline.

## 3.1 Threatened flora species

The outcome of the likelihood of occurrence for threatened flora species of moderate or high likelihood in the corridor options, is presented in Table 3.1. The complete likelihood of occurrence for threatened flora species is presented in Appendix B.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Acacia armitii	-	_	NT	Moderate
Acacia crombiei	-	V	V	Moderate
Acacia guymeri	-	_	V	Moderate
Acacia longipedunculata	-	_	NT	Moderate
Acacia tingoorensis	-	-	V	High
Ammannia robertsii	-	-	E	Moderate
Aristida thompsonii	-	-	E	Moderate
Arytera dictyoneura	-	-	NT	Moderate
*Borya inopinata	-	-	E	Moderate
Cajanus mareebensis	-	E	E	Moderate
Canarium acutifolium var. acutifolium	-	V	V	Moderate
Cerbera dumicola	-	-	NT	Moderate
*Corchorus subargenteus	-	_	V	Moderate
Corybas cerasinus	-	-	NT	Moderate

## Table 3.1Threatened flora species assessed as having of moderate or high likelihood of occurrence in the<br/>corridor options

SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Corymbia leptoloma	-	V	V	Moderate
Croton magneticus	-	-	V	Moderate
Cycas cairnsiana	-	V	V	Moderate
Cycas couttsiana	-	-	NT	Moderate
Cycas cupida	-	-	V	Moderate
Cycas desolata	-	-	V	Moderate
Cycas platyphylla	-	V	V	Moderate
Dichanthium queenslandicum	-	E	V	Moderate
Dichanthium setosum	-	V	NT	High
Diuris oporina	-	-	NT	Moderate
Dodonaea uncinata	-	-	NT	Moderate
Drosera adelae	-	-	NT	Moderate
*Drummondita calida	-	-	V	Moderate
Elaeocarpus coorangooloo	-	-	NT	Moderate
Eucalyptus paedoglauca	-	V	V	Moderate
Eucalyptus raveretiana	-	V	V	Moderate
Euphorbia carissoides	-	V	V	Moderate
*Gastrodia urceolata	-	-	V	Moderate
Genoplesium tectum	-	E	E	Moderate
*Glossocardia orthochaeta	-	-	E	Moderate
Graptophyllum excelsum	-	-	NT	Moderate
Grevillea glossadenia	-	V	V	Moderate
Habenaria rumphii	-	-	NT	Moderate
Homoranthus porteri	-	V	V	Moderate
Ipomoea saintronanensis	-	-	V	Moderate
Kardomia squarrulosa	-	-	V	Moderate
*Kunzea calida	-	_	Е	Moderate
Labichea brassii	-	-	NT	Moderate
*Lawrencia buchananensis	-	_	V	Moderate
*Leptospermum pallidum	-	-	NT	Moderate
*Lepturus minutus	-	_	V	Moderate
Lindsaea pulchella var. blanda	-	V	PE	Moderate

SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Livistona drudei			V	Moderate
Livistona lanuginosa	-	V	V	Moderate
Macropteranthes leiocaulis	-	-	NT	Moderate
Macropteranthes montana	-	V	V	Moderate
Marsdenia brevifolia	-	V	V	Moderate
*Micromyrtus rotundifolia	-	-	V	Moderate
*Oenanthe javanica	-	-	NT	Moderate
Oldenlandia polyclada	-	-	NT	Moderate
Omphalea celata	-	V	V	Moderate
Parsonsia largiflorens	-	-	Е	Moderate
Paspalidium udum	-	-	V	Moderate
Peristylus banfieldii	-	-	Е	Moderate
*Pluchea punctata	-	-	Е	Moderate
*Solanum angustum	-	-	Е	Moderate
Solanum carduiforme	-	-	V	Moderate
Solanum graniticum	-	_	Е	Moderate
Solanum sporadotrichum	-	_	NT	Moderate
Tephrosia leveillei	-	V	V	Moderate

Key: \* indicates where species and habitat information is not publically available and a conservative ranking of '*Moderate*' is applied to the species in the likelihood of occurrence assessment.

## 3.2 Threatened and migratory fauna species

The outcome of the likelihood of occurrence for threatened and/or migratory fauna species of moderate or high likelihood in the corridor options, is presented in Table 3.2. The complete likelihood of occurrence for threatened and/or migratory fauna species is presented in Appendix C.

## Table 3.2Threatened and/or migratory fauna species assessed as having of moderate or high likelihood of<br/>occurrence in the corridor options

SCIENTIFIC NAME	COMMON NAME			LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Amphibians				
Litoria nannotis	Torrent Tree Frog	E	Е	Moderate
Litoria nyakalensis	Nyakala Frog	CE	Е	Moderate
Litoria rheocola	Common Mistfrog	E	Е	Moderate

SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS	NC ACT STATUS						
Birds (threatened under EPBC Act and/or NC Act)									
Calidris ferruginea	Curlew Sandpiper	М	Е	Moderate					
Calyptorhynchus lathami erebus	Glossy Black-Cockatoo	-	V	Moderate					
Casuarius casuarius johnsonii (southern population)	Southern Cassowary	E	E	Moderate					
Cyclopsitta diophthalma macleayana	Macleay's Fig-Parrot	-	V	Moderate					
Epthianura crocea macgregori	Yellow Chat (Dawson)	CE	Е	Moderate					
Erythrotriorchis radiatus	Red Goshawk	V (M)	Е	Moderate					
Erythrura gouldiae	Gouldian Finch	E	Е	Moderate					
Erythrura trichroa	Blue-faced Parrot-finch	_	NT	Moderate					
Falco hypoleucos	Grey Falcon	-	V	Moderate					
Geophaps scripta scripta	Squatter Pigeon (southern subspecies)	V	V	High					
Grantiella picta	Painted Honeyeater	V	V	High					
Neochmia ruficauda ruficauda	Star Finch (eastern subspecies)	E	E	Moderate					
Poephila cincta cincta	Black-Throated Finch (White-Rumped subspecies)	E	Е	High					
Rostratula australis (syn. R. benghalensis)	Australian Painted Snipe	V	V	High					
Turnix olivii	Buff-breasted Button-quail	Е	Е	Moderate					
Tyto novaehollandiae kimberli	Masked Owl (northern)	V	V	Moderate					
Birds (migratory under EPBC Act and s	pecial least concern under N	C Act)							
Actitis hypoleucos	Common Sandpiper	М	SLC	Moderate					
Apus pacificus	Fork-tailed Swift	М	SLC	High					
Calidris acuminata	Sharp-tailed Sandpiper	М	SLC	Moderate					
Calidris ruficollis	Red-necked Stint	М	SLC	Moderate					
Charadrius ruficapillus	Red-capped Plover	М	SLC	Moderate					
Charadrius veredus	Oriental Plover	М	SLC	Moderate					
Chlidonias leucopterus	White-winged Black Tern	М	SLC	Moderate					
Cuculus opatus (syn. Cuculus saturatus)	Oriental Cuckoo	М	SLC	High					
Gallinago hardwickii	Latham's Snipe	М	SLC	High					
Gallinago megala	Swinhoe's Snipe	М	SLC	High					
Gallinago stenura	Pintail Snipe	М	SLC	High					

SCIENTIFIC NAME	COMMON NAME EPBC STA		NC ACT STATUS	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS	
Gelochelidon nilotica (syn. Sterna nilotica)	Gull-billed Tern	М	SLC	High	
Glareola maldivarum	Oriental Pratincole	М	SLC	Moderate	
Hirundapus caudacutus	White-throated Needletail	М	SLC	High	
Hirundo rustica	Barn Swallow	М	SLC	High	
Hydroprogne caspia (syn. Sternia caspia)	Caspian Tern	М	SLC	Moderate	
Monarcha frater	Black-winged Monarch	М	SLC	Moderate	
Monarcha melanopsis	Black-faced Monarch	М	SLC	Moderate	
Motacilla cinerea	Grey Wagtail	М	SLC	Moderate	
Myiagra cyanoleuca	Satin Flycatcher	М	SLC	High	
Numenius minutus	Little Curlew	М	SLC	Moderate	
Plegadis falcinellus	Glossy Ibis	М	SLC	High	
Rhipidura rufifrons	Rufous Fantail	М	SLC	High	
Tringa glareola	Wood Sandpiper	М	SLC	Moderate	
Tringa nebularia	Common Greenshank	М	SLC	Moderate	
Tringa stagnatilis	Marsh Sandpiper	М	SLC	Moderate	
Mammals					
Dasyurus hallucatus	Northern Quoll	E	-	High	
Dasyurus maculatus gracilis	Spotted-Tailed Quoll (Northern Subspecies)	E	E	Moderate	
Hipposideros diadema reginae	Diadem Leaf-Nosed Bat	-	NT	Moderate	
Hipposideros semoni	Semon's Leafnosed-bat	E	E	Moderate	
Macroderma gigas	Ghost Bat	V	V	Moderate	
Macrotis lagotis	Greater Bilby	V	E	High	
Onychogalea fraenata	Bridled Nailtail Wallaby	E	E	Moderate	
Ornithorhynchus anatinus	Platypus	-	SLC	High	
Petauroides volans	Greater Glider	V	V	High	
Petaurus gracilis	Mahogany Glider	E	E	High	
Phascolarctos cinereus	Koala	V	V	High	
Pteropus conspicillatus	Spectacled Flying-fox	V	V	Moderate	
Rhinolophus robertsi	Large-eared Horseshoe-bat	E	E	Moderate	
Sminthopsis archeri	Chestnut Dunnart	-	NT	Moderate	
Tachyglossus aculeatus	Short-beaked Echidna	-	SLC	High	

SCIENTIFIC NAME	COMMON NAME	EPBC ACT STATUS		LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Reptiles				
Acanthophis antarcticus	Common Death Adder	-	V	Moderate
Delma mitella	-	V	NT	Moderate
Egernia rugosa	Yakka Skink	V	V	Moderate
Lampropholis mirabilis	-	-	NT	Moderate
Lerista cinerea	Vine-thicket Fine-lined Slider	_	V	Moderate
Lerista hobsoni	Hobson's fine-liner slider	_	V	Moderate
Lerista vanderduysi	Leaden-bellied fine-line slider	V	V	Moderate
Lerista vittata	-	V	V	Moderate

The numbers of threatened and/or migratory species listed under the EPBC Act and/or NC Act that are of moderate or high likelihood of occurring in habitats potentially supported by the corridor options are presented below:

## EPBC ACT

Threatened flora species:

- → 1 threatened flora species as having a high likelihood of occurrence, including:
  - 1 vulnerable
- → **19** threatened flora species as having a moderate likelihood of occurrence, including:
  - 3 endangered
  - 16 vulnerable

Threatened fauna species:

- → 9 threatened fauna species as having a high likelihood of occurrence, including:
  - 3 endangered
  - 6 vulnerable
- → 20 threatened fauna species as having a moderate likelihood of occurrence, including:
  - 2 critically endangered
  - 10 endangered
  - 8 vulnerable
- → Migratory fauna species:
  - 11 migratory fauna species as having a high likelihood of occurrence
  - 15 migratory fauna species as having a moderate likelihood of occurrence

## NC ACT

Threatened flora species:

- → 2 threatened flora species as having a high likelihood of occurrence, including:
  - 1 vulnerable
  - 1 near threatened
- → 62 threatened flora species as having a moderate likelihood of occurrence, including:
  - 1 presumed extinct
  - 12 endangered
  - 31 vulnerable
  - 18 near threatened

Threatened fauna species (including special least concern):

- → 21 threatened fauna species as having a high likelihood of occurrence, including:
  - 3 endangered
  - 5 vulnerable
  - 13 special least concern
- → 46 threatened fauna species as having a moderate likelihood of occurrence, including:
  - 14 endangered
  - 12 vulnerable
  - 5 near threatened
  - 15 special least concern.

## 4 CONCLUSION

This preliminary desktop-based likelihood of occurrence has identified the threatened flora and fauna species, and migratory fauna species, listed under the EPBC Act and/or NC Act that have a moderate to high likelihood of occurring in the Project's study area.

It has identified the species of conservation significance that should be considered when designing field surveys. It also identifies those species with a high ranking that may be at risk of project related impacts.

The findings of this assessment will be beneficial in planning the next phase of the Project.

# Appendix A

**BROAD VEGETATION GROUPS (1:1,000,000 SCALE)** 

## 1:1,000,000 scale - Broad vegetation groups (BVG) used for likelihood of occurrence a

BVG_1M	BVG DESCRIPTION
:1 Million	
7	Semi-evergreen to deciduous microphyll vine thicket
7a	Semi-evergreen vine thickets on wide range of substrates. (land zones 8, 9, 11, 12, 5, 4, 3, 10, [7]) (BRB, EIU, SEQ, CQC, [WET, GUP]) (Tracey 1982 11)
9	EUCALYPT WOODLANDS TO OPEN FORESTS (mainly eastern Qld) Moist to dry eucalypt open forests to woodlands usually on coastal lowlands and ranges
9 9c	Open forests of <i>Corymbia clarksoniana</i> (grey bloodwood) (or <i>C. intermedia</i> (pink bloodwood) or <i>C. novoguinensis</i> ), <i>C.</i>
30	tessellaris (carbeen) ± Eucalyptus tereticornis (blue gum) predominantly on coastal ranges. Other frequent tree species include Eucalyptus drepanophylla (grey ironbark), E. pellita (large-fruited red mahogany), E. brassiana (Cape York red gum) and Lophostemon suaveolens (swamp box). (land zones 12, 11, 8, 5). (WET, CQC, CYP, BRB, [SEQ])
9d	Moist to dry open forest to woodland dominated by <i>Eucalyptus portuensis, Corymbia intermedia</i> (pink bloodwood), <i>E. drepanophylla, E. resinifera</i> or <i>E. reducta</i> +/- <i>Syncarpia glomulifera</i> (turpentine) or <i>E. cloeziana</i> (Gympie messmate) on range Also includes mixed forests with <i>Eucalyptus pellita</i> or <i>C. torelliana</i> (cadaghi) emergents and rainforest understories (land zone 12, 11, 3, 5, 9, 8). (WET, CQC, EIU)
9e	Open forests, woodlands and open woodlands dominated by <i>Corymbia clarksoniana</i> (grey bloodwood) (or <i>C. novoguinensis</i> of <i>C. intermedia</i> (pink bloodwood) or <i>C. polycarpa</i> (long-fruited bloodwood)) frequently with <i>Erythrophleum chlorostachys</i> (red ironwood) or <i>Eucalyptus platyphylla</i> (poplar gum) predominantly on coastal sandplains and alluvia. (land zones 3, 5, 2) (CYP, BRB, CQC, WET, EIU)
11	Moist to dry eucalypt open forests to woodlands mainly on basalt areas
11b	Moist to dry open forests to woodlands dominated by <i>Eucalyptus crebra</i> (narrow-leaved red ironbark) or <i>E. tereticornis</i> (blue gum), frequently with <i>Corymbia</i> species or <i>E. microneura</i> (Gilbert River box) on red krasnozems on undulating terrain. (land zone 8) (EIU)
11c	Moist woodlands dominated by <i>Eucalyptus leptophleba</i> (Molloy red box) ± <i>Corymbia papuana</i> (ghost gum) ± <i>C. tessellaris</i> (carbeen). (land zones 8, 11) (EIU, CYP)
12	Dry eucalypt woodlands to open woodlands, mostly on shallow soils in hilly terrain (mainly on sandstone and weathered rocks)
12b	Woodlands and open woodlands dominated by <i>Eucalyptus crebra</i> (sens. lat) (narrow-leaved red ironbark) and/or <i>Corymbia</i> spp. such as <i>C. clarksoniana</i> (grey bloodwood), <i>C. stockeri, C. setosa</i> (rough leaved bloodwood) or <i>C. peltata</i> (yellowjacket) hilly terrain. (land zones 7, 10, 11) (GUP, EIU, DEU, CYP)
13	Dry to moist eucalypt woodlands and open forests, mainly on undulating to hilly terrain of mainly metamorphic and acid igneous rocks
13a	Woodlands and open woodlands dominated by ironbarks such <i>Eucalyptus cullenii</i> (Cullen's ironbark), <i>E. staigeriana</i> (lemon- scented ironbark) or <i>E. melanophloia</i> (silver-leaved ironbark) and bloodwoods such as <i>Corymbia stockeri</i> subsp. <i>peninsularis</i> , <i>C. clarksoniana</i> (grey bloodwood) or <i>C. leichhardtii</i> (rustyjacket). (land zones 11, 12, 7, 5) (EIU, CYP, GUP)
13b	Woodlands to open woodlands dominated by <i>Eucalyptus microneura</i> (Gilbert River box) on shallow soils on rolling hills. (land zones 12, 11, 9) (EIU, GUP)
13c	Woodlands of <i>Eucalyptus crebra</i> (sens. lat.) (narrow-leaved red ironbark), <i>E. drepanophylla</i> (grey ironbark), <i>E. fibrosa</i> (dusky leaved ironbark), <i>E. shirleyi</i> (Shirley's silver-leaved ironbark) on granitic and metamorphic ranges (land zones 12, 11, 9, [5]) (BRB, EIU, SEQ, NET, CQC)
	EUCALYPT OPEN FORESTS TO WOODLANDS ON FLOODPLAINS
16	Eucalyptus spp. dominated open forest and woodlands drainage lines and alluvial plains
16a	Open forest and woodlands dominated by <i>Eucalyptus camaldulensis</i> (river red gum) (or <i>E. tereticornis</i> (blue gum)) and/or <i>E. coolabah</i> (coolabah) (or <i>E. microtheca</i> (coolabah)) fringing drainage lines. Associated species may include <i>Melaleuca</i> spp., <i>Corymbia tessellaris</i> (carbeen), <i>Angophora</i> spp., <i>Casuarina cunninghamiana</i> (riveroak). Does not include alluvial areas dominated by herb and grasslands or alluvial plains that are not flooded. (land zone 3) (MGD, BRB, GUP, CHC, MUL, DEU, EIU, NWH, SEQ, [NET, WET]) (All bioregions except CYP and CQC)
16b	Woodlands dominated by <i>Eucalyptus leptophleba</i> (Molloy red box), with <i>Corymbia tessellaris</i> (carbeen) or <i>C. clarksoniana</i> (grey bloodwood) or <i>C. dallachiana</i> . On sandy levees. (land zones 3, 5) (GUP, EIU, CYP)
16c	Woodlands and open woodlands dominated by <i>Eucalyptus coolabah</i> (coolabah) or <i>E. microtheca</i> (coolabah) or <i>E. largiflorem</i> (black box) or <i>E. tereticornis</i> (blue gum) or <i>E. chlorophylla</i> on floodplains. Does not include alluvial areas dominated by herb and grasslands or alluvial plains that are not flooded. (land zone 3) (All bioregions except WET, principally GUP, BRB, MUL, SEQ)
16d	River beds, open water or sand, or rock, frequently unvegetated. (land zone 3) (GUP, EIU, BRB, CYP, WET, SEQ, DEU, [CQ0 MUL])
	EUCALYPT DRY WOODLANDS ON INLAND DEPOSITIONAL PLAINS
17	<i>Eucalyptus populnea</i> (poplar box) or <i>E. melanophloia</i> (silver-leaved ironbark) (or <i>E. whitei</i> (White's ironbark)) dry woodlands to open woodlands on sandplains or depositional plains
17a	Woodlands dominated by <i>Eucalyptus populnea</i> (poplar box) (or <i>E. brownii</i> (Reid River box)) on alluvium, sand plains and footslopes of hills and ranges. (land zones 3, 5, 10, 9, 4, 11, 12, [8]) (BRB, MUL, DEU, EIU)
17b	Woodlands to open woodlands dominated by Eucalyptus melanophloia (silver-leaved ironbark) (or E. shirleyi (shirley's silver- leaved ironbark)) on sand plains and footslopes of hills and ranges. (land zones 5, 12, 3, 11, 9, 7) (BRB, DEU, EIU, SEQ, NET
	GUP, NWH)

# Appendix B

FLORA LIKELIHOOD OF OCCURRENCE

## Flora likelihood of occurrence

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Acacia armitii	-	-	NT	Only known from the Einasleigh R. area in central-northern Qld and on a sandstone plateau S of Goomadeer, as well as Coopers Ck at Nabarlek, N.T. Locally common in sandy or shallow, rocky soils, on creek banks and river flats {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Charters Towers LGA, Etheridge LGA	Moderate (Potential to occur in BVGs 16 associated subtypes)
Acacia crombiei	-	V	V	Occurs in small isolated subpopulations in central Queensland. It occurs in the Burke, Mitchell, and North Kennedy pastoral regions. It inhabits wooded downs in woodland and open woodland often associated with gidgee (Acacia cambagei) and whitewood (Atalaya hemiglauca), on alluvial, sandstone and basalt derived soils {Department of the Environment and Energy, 2017 #7538}.	PMST - Charters Towers LGA, Etheridge LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 24 and 27 and associated subtypes)
Acacia guymeri	-	-	V	The species grows in skeletal soils on rocky ridges in disturbed areas and eucalypt woodland. Commonly associated plants include Eucalyptus cullenii, Lamprolobium fruticosum, and a range of different Eucalyptus and Corymbia species {Department of the Environment and Energy, 2017 #7539}.	Wildlife Online - Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 12 and associated subtypes)
Acacia longipedunculata	-	-	NT	Restricted an area near Herberton and in the Paluma Ra., north-eastern Qld. Grows on shallow sandy and rocky soils, in open forest {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 9, 12 and 13 and associated subtypes)
Acacia tingoorensis	-	-	V	Grows in eucalypt woodland or forest, on deep red loam, shallow loamy or sandy soils.	Wildlife Online - Charters Towers LGA	High (Potential to occur in BVGs 9,11,12,13 and associated subtypes)
Ammannia robertsii	-	-	E	Low open woodland of Acacia cambagei on alluvial clay plain.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 16, 24 , 27 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Aristida thompsonii	-	-	E	Restricted distribution around North Kennedy in Queensland. Inhabits patchy heath of Kunzea callida on rocky areas.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 12 and associated subtypes)
Arytera dictyoneura	-	-	NT	Recorded from scattered locations in northern and southern Queensland. In northern Queensland A. dictyoneura occurs in Girringun National Park, on the Cardwell Range, Cardwell Forest Reserve, near Meunga Creek, Clement State Forest and Paluma Range. There is also one record of the species from Strathdickie, near Proserpine. In southern Queensland, there are numerous records of A. dictyoneura in Bulburin National Park, two records in Mount Colosseum National Park, one population near 'shirley' Miriam Vale and one population in Eurimbula National Park. Found in semi-evergreen rainforest, on granite boulder slopes/ soil derived from granite {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Borya inopinata	-	-	E	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> Conservative ranking applied, due to lack of species and habitat information.
Bulbophyllum globuliforme	Hoop Pine Orchid	V	NT	Restricted to the MacPherson Range and north to Gladstone, e.g. Grady's Creek (Border Ranges). In NSW occurs at altitudes between 300 - 600 m where it grows on Hoop Pines (Araucaria cunninghamii) in upland subtropical rainforest. Usually found on the upper trunk and branches of emergent trees {NSW National Parks and Wildlife Service, 2002 #93; Harden, 1993 #4}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA, Wildlife Online - Charters Towers LGA	Low (Suitable habitat is unlikely to occur)
Cajanus mareebensis	-	E	E	Occurs in grassy woodlands of Melaleuca-Acacia, Eucalyptus-Callitris and Eucalyptus-Corymbia woodlands on sandy soils derived from granite with a lower horizon of impeded drainage {Department of the Environment Water Heritage and the Arts, 2008 #7540}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> (Potential to occur in BVG 13 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Canarium acutifolium var. acutifolium	_	V	V	Its distribution occurs between Mossman and Tully in Queensland. Altitudinal range very small, from sea level to 100 m. almost confined to creek and river banks in lowland rain forest {Department of the Environment Water Heritage and the Arts, 2008 #7541}.	PMST - Charters Towers LGA	Moderate (Potential to occur in BVG 16 and associated subtypes)
Cerbera dumicola	_	-	NT	Occurs across a range of habitats in central and southern Queensland. Associated vegetation and species include: sandstone hills in open E. umbra subsp. carnea; on plateaus, in woodland of Acacia shirleyi with Corymbia dolichocarpa; acidic soils in mine rehabilitation area; woodland of A. catenulata and A. shirleyi with E. thozetiana on a slope of sand/clay soil; semi-deciduous notophyll-microphyll vine forest of Brachychiton australis, Gyrocarpus americanus, Flindersia australis, Pleiogynium timorense, Drypetes deplanchei and Sterculia quadrifida on rhyolite hillslopes; open- woodland of E. melanophloia with occasional Acacia shirleyi, E. populnea and E. brownii; semi-evergreen vine thicket with Corymbia citriodora and Corymbia aureola emergents; woodland of A. rhodoxylon on brown, sandy loam; and in Corymbia tessellaris - Acacia aneura open woodland {Queensland Herbarium, 2011 #7542}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 7, 9, and 12 and associated subtypes)
Corchorus subargenteus	_	-	V	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Corybas cerasinus	_	-	NT	Occurs in north-eastern Queensland with a distribution from Cooktown to the Herbert River near Ingham and also on Dunk Island. Occurs in moist to wet forests on exposed ridges and in drier forests. It grows in well-drained sand and gravelly loam{Department of Sustainability Environment water Population and Communities, 2010 #7543}.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Moderate (Potential to occur in BVGs 9 and 11 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Corymbia leptoloma	-	V		Environment Water Heritage and the Arts, 2008 #7544}.	PMST - Charters Towers LGA, Hinchinbrook LGA, Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> (Potential to occur in BVGs 9 and 11 and associated subtypes)
Croton magneticus	-	-		between Townsville and Proserpine, but is known to extend inland to	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Cyathea celebica	-	-		Cyathea celebica occurs on steep slopes and river banks in rainforest, vine forest and forest margins, above 100m altitude. It grows on granitic soil, reddish soils and over metamorphic rock {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Cycas cairnsiana	-	V	V	shallow to skeletal gritty soils over siliceous granites. Plants from the population near Forsayth have slightly wider leaflets than those from near Mt Surprise {Royal Botanic Gardens, 2010 #7525}.	PMST - Charters Towers LGA, Etheridge LGA, Wildlife Online - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 12b, 13b,)
Cycas couttsiana	_	-	NT	in open grassy woodlands in red sandy loams derived from basalt or dolerite {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Charters Towers LGA, Etheridge LGA	Moderate (Potential to occur in BVGs 11b and 11c)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Cycas cupida	_	-	V	This species is known from a single, quite extensive population in the Terrace Range south of Charters Towers, South Kennedy District, Queensland. Found widely scattered in open woodland on low sandstone hills.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs and 12 and associated subtypes)
Cycas desolata	-	-	V	Known only from a single stand near Charters Towers, eastern Qld. Grows on Eucalypt woodland on shallow skeletal soils or low rocky outcrops in flat country of open ironbark woodland.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs and 12 and associated subtypes)
Cycas platyphylla	_	V	V	The main population of C. platyphylla is known from the Petford district, west of the Atherton Tableland, Queensland. There are three smaller quite disjunct populations recorded from Taravale, Wandovale, and at White Mountains, north of Torrens Creek. Occurs in sparse Eucalyptus sideroxylon woodland with a grassy understorey, often on rocky slopes in shallow red stony loams.	PMST - Charters Towers LGA, Etheridge LGA, Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 9 and 11 and associated subtypes)
Cyperus cephalotes	_	E	E	Known from Rockingham Bay, near Cardwell and Trebonne Creek south-east of Ingham in Queensland. It occurs on floating islands in rivers, with the roots entangled in a mass of decaying vegetation.	PMST - Hinchinbrook LGA, Wildlife Online - Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Dendrobium bigibbum	_	V	V	Subpopulations of the Cooktown orchid group are epiphytic (growing on branches) and epilithic (growing on rocks), and occurs at sites with moderate light intensity. The area where it occurs has moderate to high rainfall that is seasonal (more rainfall in summer and autumn). Collections have been made from sea level (including on branches overhanging salt water) to altitudes of at least 250 m above sea level. Many collections have been made along creeks or on rocky hillsides where fire cannot penetrate. Vegetation associations where it occurs include closed forest (low deciduous scrub, coastal dunes, gallery forest), open monsoon forest, mangrove, heath and inland dry scrub	PMST - Etheridge LGA	Low (Suitable habitat is unlikely to occur)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Dendrobium lithocola	-	E	-	Subpopulations of the Cooktown orchid group are epiphytic (growing on branches) and epilithic (growing on rocks), and occurs at sites with moderate light intensity. The area where it occurs has moderate to high rainfall that is seasonal (more rainfall in summer and autumn). Collections have been made from sea level (including on branches overhanging salt water) to altitudes of at least 250 m above sea level. Many collections have been made along creeks or on rocky hillsides where fire cannot penetrate. Vegetation associations where it occurs include closed forest (low deciduous scrub, coastal dunes, gallery forest), open monsoon forest, mangrove, heath and inland dry scrub	PMST - Etheridge LGA	Low (Suitable habitat is unlikely to occur)
Dichanthium queenslandicum	-	E	V	Occurs within the South Eastern Queensland, Brigalow Belt South, Brigalow Belt North, Central Mackay Coast, Desert Uplands, Mitchell Grass Downs and Einasleigh Uplands Bioregions; and the South East Queensland, Condamine, Border Rivers Maranoa-Balonne, Burnett Mary, Fitzroy, Burdekin, Mackay Whitsunday, Southern Gulf and Desert Channels Natural Resource Management Regions.	PMST - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 30b and 32a)
Dichanthium setosum	-	V	NT	Grows in woodland and grassland {Harden, 1993 #4}. On the New England Tablelands and North West Slopes it grows on stony red-brown hard-setting soils over basalt, or on black soil {Department of Environment and Conservation, 2006 #1093}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	High (Potential to occur in BVGs 11,16,17,30,32 and associated subtypes)
Diuris oporina	-	-	NT	Found in Queensland on ridges and slopes in sparse grassy forests at elevations of 700 to 1200 meters as a miniature to small sized, warm to cool growing terrestrial species.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 11,12 and 13 and associated subtypes)
Dodonaea uncinata	-	-	NT	Confined to the Mt Spec area NW of Townsville, Qld. Grows in open forest or woodland, usually in sandstone soils {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 12 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Drosera adelae	-	-	NT	An endemic species that occurs in a restricted area in NEQ from Tully southwards to Hitchenbrook Island. Altitudinal range from 50-800 m. Occurs on creek beds and on moss-covered rocks along creeks in rainforest, open forest, mesophyll vine forest and in Eucalypt forest.	Wildlife Online - Hinchinbrook LGA	Moderate (Potential to occur in BVGs 16 and associated subtypes)
Drummondita calida	-	-	V	Species and habitat information is not publically available.	Wildlife Online - Etheridge LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Elaeocarpus coorangooloo	-	-	NT	Endemic to NEQ, known only from the Windsor Tableland, Atherton Tableland and another site southwest of Paluma. Altitudinal range from 700- 1000 m. Grows in drier rain forest often associated with Kauri Pine (Agathis robusta).	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Eriocaulon carsonii		E	Ε	Grows in running water and forms dense mats in wet soil around shallow springs. The species is an endemic of active or flowing artesian mound springs on the margins of the Great Artesian Basin. Mound springs are natural outlets of the Basin, associated with fractures and fault lines, often having mounds of various sizes. Accumulated evaporite and mud deposits form mounds 1 to 10 metres high and 2 to 100+ metres in diameter. The faults provide direct access for the artesian water to reach the surface. These landforms are probably one of the rarest habitats in Australia. Originally restricted to a single mound at Peery Lake in NSW, in an area of many mounds. More recently the plant has spread to adjacent mounds, indicating that the plant can survive for long periods as a small population and then spontaneously expand. The population structure of Eriocaulon carsonii changed after fencing at Elizabeth Springs in Qld, with large numbers of small immature plants replaced by a smaller number of larger plants. Observations of density of kangaroo scats at Peery Lake suggest that kangaroos heavily graze the mounds. Kangaroo grazing apparently limits the growth of sedges on the mounds, reducing competition, thus benefiting Eriocaulon carsonii. The species is often recorded growing in dense mats of numerous individuals {Harden, 1993 #4} {Royal Botanic Gardens, 2006 #990}.	PMST Etheridge LGA Wildlife Online - Etheridge LGA	Low (Suitable habitat is unlikely to occur)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Eriocaulon carsonii subsp. orientale	-	E	E	Entirely restricted to vents and tails of mound spring wetlands, particularly springs with shallow standing water with slow flow. It is generally associated with vegetated mounds that, over considerable time, have formed organic fen soils (alkaline equivalent of the acidic peat bog).	Wildlife Online - Etheridge LGA	Low (Suitable habitat is unlikely to occur)
Eryngium fontanum	-	E	E	Known from two spring wetland complexes (Moses (Doongmabulla) and Edgbaston-Myross) in central Queensland in the Barcaldine spring super- group (larger regional groups of springs). Occurs in very restricted habitat in shallow permanent ponded water of wetlands associated with artesian springs {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA	Low (Suitable habitat is unlikely to occur)
Eucalyptus paedoglauca	-	V	V	Occurs only in the Townsville area of north-east Queensland. Situated on ridges or hill slopes on shallow sandy-loam soil.	PMST - Charters Towers LGA	Moderate (Potential to occur in BVGs 11,12 and 13 and associated subtypes)
Eucalyptus raveretiana	-	V	V	Wide distribution in coastal and sub-coastal areas of Queensland, from south of Townsville to Nebo, around Rockhampton and areas 100km west of the city. Usually grows along watercourses, and sometimes on river flats or open woodland. Soil varies from sand through to heavy clay.	PMST - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 16 and associated subtypes)
Euphorbia carissoides	-	V	V	Chamaesyce carissoides is restricted to north-east Queensland where it is known from near Georgetown and east to Stannary Hills, with a disjunct occurrence near Hopevale. The species grows on clifflines, among rocky outcrops and hillsides in shrubland and eucalypt low open woodland communities in generally shallow soils derived from sandstone, granite and rhyolite substrates.	PMST - Etheridge LGA Wildlife Online - Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 13 and associated subtypes)
Gastrodia urceolata	-	-	V	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Genoplesium tectum	-	E	E	Known only from a small area south of Cardwell in north-eastern Queensland. Occurs in dense scrub not far from a creek among dense sedges and low shrubs in woodland dominated by Melaleuca viridiflora. Soils are seasonally inundated sandy loams {Department of the Environment and Energy, 2017 #7539}.	PMST - Hinchinbrook LGA	<i>Moderate</i> (Potential to occur in BVG 21a)
Glossocardia orthochaeta	-	-	E	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> Conservative ranking applied, due to lack of species and habitat information.
Graptophyllum excelsum	-	-	NT	Occurs in NEQ and coastal central Queensland. Altitudinal range in NEQ from 350-600 m. Often grows on limestone in monsoon forest and vine thickets.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Grevillea glossadenia	-	V	V	Occurs in north-east Queensland mainly in the Einasleigh Uplands(IBRA) bioregion, with a few collections from the Wet Tropics Bioregion. Grows in eucalypt woodland or low open forest, in shallow to skeletal granitic soils on rolling hills, gravel terraces near stream beds, and along roadsides and mining tracks {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 12 and 13 and associated subtypes)
Habenaria rumphii	_	-	NT	Grows on sandy soils in seasonally-flooded grassy areas in open forest and woodland. Found both in Queensland and the Northern Territory.	Wildlife Online - Hinchinbrook LGA	Moderate (Potential to occur in BVGs 16 and 17 and associated subtypes)
Homoranthus porteri	-	V	V	Restricted to north-east Queensland from near Mareeba southwards to near Ravenshoe. Occurs in shallow soils on a variety of rock types (including rhyolite), usually in woodland or heath. It has been recorded on sandstone pavement, rock outcrops and scree slopes, on the edge of rocky escarpments and rocky hillsides {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA, Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 12 and 13 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
lpomoea saintronanensis	-	-	V	Endemic to NEQ. Altitudinal range from 600-900 m. Grows in vine thicket and monsoon forest.	Wildlife Online - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Kardomia squarrulosa	-	-	V	On sandstone outcrops, with Acacia shirleyi.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 12 and associated subtypes)
Kunzea calida	-	-	E	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Labichea brassii	-	-	NT	Confined to northern Qld. Known from Mt Mulligan in the north south-west to Agate Creek. Recorded growing along creek and river beds {Department of the Environment and Energy, 2017 #7537}.	Wildlife Online - Etheridge LGA	Moderate (Potential to occur in BVGs 16 and 17 and associated subtypes)
Lawrencia buchananensis	-	-	V	Species and habitat information is not publically available.	PMST - Charters Towers LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> Conservative ranking applied, due to lack of species and habitat information.
Leptospermum pallidum	-	-	NT	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA, Etheridge LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Lepturus minutus	-	-	V	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> Conservative ranking applied, due to lack of species and habitat information.
Lindsaea pulchella var. blanda	-	V	PE	Known in Australia from a single specimen collected at 'Rockingham, 4000ft alt This is a very poorly known taxon in Australia with only one record from an uncertain locality, but possibly in the Rockingham Bay Range, Queensland. This species occurs within the Wet Tropics (Queensland) Natural Resource Management Region.	PMST - Charters Towers LGA, Hinchinbrook LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Livistona drudei	-	-	V	Occurs in NEQ from Tully to Conway Beach. Altitudinal range from near sea level to 300 m. Grows along stream banks and coastal plains in Melaleuca swamp forest to fringes of gallery rainforest and rainforest to eucalypt forest.	Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> (Potential to occur in BVGs 21 and 22 and associated subtypes)
Livistona Ianuginosa	-	V	V	Endemic to the Burdekin-Ravenswood-Cape River area inland from Ayr, where it is found on the tributaries of the Burdekin River. Forms colonies along streambanks and gullies well inland from the coast and is found at altitudes of 150–300m above sea level. Its habitat is open woodland on sandy river and creek channels which flow for part of the year, with permanent pools or soaks {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA, Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 16 and 17 and associated subtypes)
Macropteranthes leiocaulis	-	-	NT	Occur in the semi-evergreen vine thickets.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Macropteranthes montana	_	V	V	Known from a small area just south of Cape York Peninsula, Queensland. As far north as Hutchinson (inland from Cape Tribulation) and from localities near Mount Mulligan, Chillagoe, Dimbulah, Petford, Irvinebank, Elizabeth Creek Gorge, Bulleringa National Park and south west as far as the northern Newcastle Range. Occurs in shallow soil in low woodland or vine thicket and occurs within the Northern Gulf and Wet Tropics (Queensland) Natural Resource Management Regions {Department of the Environment and Energy, 2017 #7539}.	PMST - Etheridge LGA Wildlife Online - Etheridge LGA	<i>Moderate</i> (Potential to occur in BVG 7a)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Marsdenia araujacea	-	Х	PE	Was endemic to far northern Queensland between Cooktown and Ingham, south of Cairns. Grew in lowland rainforest	Wildlife Online - Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Marsdenia brevifolia	-	V	V	Occurs in north and central Queensland where it is known from near Townsville, Springsure and north of Rockhampton. Grows on serpentine rock outcrops or crumbly black soils derived from serpentine in eucalypt woodland, often with Broad-leaved Ironbark (Eucalyptus fibrosa) and Corymbia xanthope {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVGs 11 and associated subtypes)
Micromyrtus rotundifolia	-	-	V	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Myrmecodia beccarii	-	V	V	Occurs in coastal woodland and mangrove between Cooktown and Ingham in Queensland. It is a unique epiphyte that has a special association with the golden ant which lives in the chambers of the tuber, and the Apollo Jewel Butterfly which lays its eggs on the plant.	PMST - Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Oenanthe javanica	-	-	NT	Species and habitat information is not publically available.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Oldenlandia polyclada	-	-	NT	Occurs in NEQ with a restricted distribution from Cooktown to Townsville. Altitudinal range from 160-300 m. Grows along drainage lines in forest and vine thicket, occasionally in woodland.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 16 and 17 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Omphalea celata	-	V		Known from three sites in central east Queensland. Locations include Hazlewood Gorge, near Eungella; Gloucester Island, near Bowen; and Cooper Creek in the Homevale Station area, north-west of Nebo. Grows in fragmented semi-evergreen vine thicket along a watercourse on weathered metamorphics in a steep-sided gorge at an altitude of 560m {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Parsonsia largiflorens	-	-	E	Grows in rainforest, including drier types; from the Cairns area south to the Tweed River, now apparently extinct in NSW {Royal Botanic Gardens, 2017 #3885}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Paspalidium udum	-	-	V	Endemic to tropical Australia. Rare in NT. Occurs on black soil plains.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 11 and associated subtypes)
Peristylus banfieldii	-	-	E	Grows in open forest and on swamp margins in seasonally wet (monsoon season) soils.	Wildlife Online - Hinchinbrook LGA	Moderate (Potential to occur in BVGs 16, 21 and 22 and associated subtypes)
Phaius australis	Southern Swamp Orchid	E	E	Previously occurred as far south as Port Macquarie but is now thought to only occur north of Coffs Harbour. Grows in coastal areas in swampy grassland or forest including rainforest, eucalypt o paperbark forest. Flowers sept-oct (Harden 1993; NPWS 2002).		Low (Suitable habitat is unlikely to occur)
Phaius pictus	_	V		Occurs in north-east Queensland, sporadically from the McIlwraith Range, Bloomfield River to Kirrama Range. It is highly localised, restricted to rainforests from 0–600m altitude, and usually occurs in sheltered humid sites close to streams and seepage among forest litter on boulders {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA, PMST Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Phalaenopsis amabilis subsp. rosenstromii	-	E	E	The Native Moth Orchid is found in humid rainforest areas, close to waterfalls or streams, in deep gorges, sheltered slopes or gullies in notophyll vine thickets, deciduous vine thickets and in open forest. The Native Moth Orchid grows in shaded or partially shaded positions, on trees and less commonly on rocks. The species is found at altitudes from 200–500 m above sea level {Department of the Environment, 2015 #7233}	PMST - Charters Towers LGA, Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Phlegmariurus phlegmarioides	-	-	V	The Layered Tassel Fern occurs in the lowland tropical rainforest of north eastern Queensland, Australia.	Wildlife Online - Charters Towers LGA	Low (Suitable habitat is unlikely to occur)
Pluchea punctata	-	-	E	Species and habitat information is not publically available.	Wildlife Online - Etheridge LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Solanum angustum	-	-	E	Species and habitat information is not publically available.	Wildlife Online - Etheridge LGA	Moderate Conservative ranking applied, due to lack of species and habitat information.
Solanum carduiforme	-	-	V	Known from 15 populations in a disjunct distribution from Queensland, the Northern Territory and Western Australia. In Queensland, the distributional range is 600 km and it occurs in two main areas: Bowthorne, Boodjamulla (Lawn Hill) National Park, and nearby 'Musselbrook' (north-west Queensland) and Cobbold Gorge, Forsayth and Richmond (southern Gulf). Known to occur on conglomerate rock formations. Other locations are on sandstone or deeper sandy soil adjacent to sandstone outcrops {Department of the Environment and Energy, 2017 #7539}.	Wildlife Online - Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 11, 12 and 13 and associated subtypes)
Solanum graniticum	-	-	E	Found in Qld on the mainland in the Bowen area and on adjacent Gloucester Island and also in the Eungulla Dam area. Occurs in Eucalypt woodland on soils derived from granite or granodiorite.	Wildlife Online - Charters Towers LGA	Moderate (Potential to occur in BVGs 13 and associated subtypes)

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Solanum sporadotrichum	-	-	NT	Grows in association with semi-evergreen vine thicket, notophyll rainforest or littoral rainforest with Brachychiton australis, Gyrocarpus americanus, Flindersia collina, Araucaria cunninghamii, Acacia fasciculifera and Drypetes deplanchei, or in eucalypt open forest or woodland. Soils are moderately to very fertile.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> (Potential to occur in BVG 7a)
Tephrosia leveillei	_	V	V	Known from the area between Chillagoe and Forty Mile Scrub with one specimen further south, near Ravenswood. Recorded growing on alluvial plains in Eucalyptus cullenii woodland with Corymbia erythrophloia, Erythrophleum chlorostachys and Grevillea glauca, and in tall open forest of Eucalyptus and Corymbia species over dense Heteropogon contortus on red sand.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA, Wildlife Online - Etheridge LGA	<i>Moderate</i> (Potential to occur in BVGs 11, 12 and 13 and associated subtypes)
Tropilis callitrophilis	-	V	V	Distributed in north-eastern Queensland from Mount Finnigan to the Evelyn Tableland. Occurs in rainforests and rainforest margins at high altitudes. It favours Stringybark Cypress Pine (Callitris macleayana) as a host, but also grows on various shrubby myrtles, such as Rhodamnia and Austromyrtus. Flowering between August-September. {Department of the Environment and Energy, 2017 #7521}	PMST - Charters Towers LGA, Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)
Zeuxine polygonoides	-	V	V	In Australia, the Velvet Jewel Orchid grows in moist shady sites in rainforests (mesophyll vine forests and simple notophyll vine forests) in leaf litter on the ground or on large boulders adjacent to streams. Altitudinal range is 450–820 m above sea level. Found mostly from moist, cloudy or very wet rainfall zones on metamorphic substrates, granite or rhyolite. The species can be found in humus on flat topped rocks in association with Anoectochilus yatesiae, Goodyera viridiflora and Liparis simmondsii {Department of the Environment and Energy, 2017 #7526}	PMST - Hinchinbrook LGA	Low (Suitable habitat is unlikely to occur)

Key: X = Extinct, PE= Presumed Extinct, CE = Critically Endangered, E = Endangered, V= Vulnerable, NT = Near Threatened

# Appendix C

FAUNA LIKELIHOOD OF OCCURRENCE

#### Fauna likelihood of occurrence

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Amphibians						
Litoria dayi (syn. Nyctimystes dayi)	Australian Lacelid, Day's Frog	E	E	The Lace-eyed Tree Frog is a rainforest species, endemic to the Wet Tropics Bioregion. The Lace-eyed Tree Frog occurred throughout the Wet Tropics Bioregion from Paluma to Cooktown, northern Queensland, at altitudes between 0 and 1200 m. It is associated with rainforests and rainforest margins. In montane areas the species prefers fast-flowing rocky streams although they also frequent slower watercourses where ample vegetation exists along the margins. At low elevations, the Lace-eyed Tree Frog favours rock soaks, narrow ephemeral streams and rock outcrops in larger watercourses. It may also be found on rocks, boulders and vegetation in or adjacent to streams {Department of the Environment and Energy, 2017 #7520}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Rainforest not present.
Litoria nannotis	Torrent Tree Frog	E	E	The Torrent Tree Frog occurs throughout the Wet Tropics Bioregion, North Queensland, from Paluma to Cooktown, but only has stable populations at lowland sites (180-400 m). It is restricted to rocky stream habitats in rainforest or wet sclerophyll forest where there is fast flowing water, waterfalls and cascades {Department of the Environment and Energy, 2017 #7527}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in open forest riparian habitats (BVG 16 and subsets).
Litoria nyakalensis	Nyakala Frog	CE	E	Endemic to the Wet Tropics Bioregion in Far North Queensland. Adults occur in rainforest and wet sclerophyll forest, next to white water sections of fast flowing streams. Often found on rocks adjacent to the stream or overhanging vegetation (Department of the Environment and Water Resources, 2007).	PMST - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in open forest riparian habitats (BVG 16 and subsets).

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Litoria rheocola	Common Mistfrog	E	E	Endemic to the Wet Tropics Bioregion (Williams & Hero 1998, 2001). The species is restricted to fast flowing rocky creeks and streams in rainforest as well as wet sclerophyll forest (Liem 1974; McDonald 1992). Within these streams this species are often found in the slower more open sections, away from waterfalls (Hodgkison & Hero 2002). Individuals can be found on rocks, logs and vegetation in or adjacent to streams (Hero & Fickling 1994).	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur in open forest riparian habitats (BVG 16 and subsets).
Litoria serrata	Green-eyed Tree Frog	-	V	Occurs from Paluma to Big Tableland in north-eastern Queensland in rainforest habitats.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Rainforest not present.
Birds						
Actitis hypoleucos	Common Sandpiper	Μ	SLC	The Common Sandpiper frequents a wide range of coastal wetlands and some inland wetlands, with varying levels of salinity. It is mostly encountered along muddy margins or rocky shores and rarely on mudflats. It has been recorded in estuaries and deltas of streams, banks farther upstream; around lakes, pools, billabongs, reservoirs, dams and claypans, and occasionally piers and jetties. The muddy margins utilised by the species are often narrow, and may be steep. The species is often associated with mangroves, and sometimes found in areas of mud littered with rocks or snags {Geering, 2007 #3752}{Higgins, 1996 #648}. Roost sites are typically mangroves. The species is known to perch on posts, jetties, moored boats and other artificial structures, and to sometimes rest on mud or 'loaf' on rocks {Higgins, 1996 #648}.	PMST - Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Apus pacificus	Fork-tailed Swift	Μ	SLC	Breeds in the northern hemisphere, wintering south to Australia. It is almost exclusively aerial, flying from less than 1 m to at least 300 m above ground. It mostly occurs over inland plains but sometimes above foothills or in coastal areas over cliffs, beaches, islands and well out to sea. It also occurs over towns and cities. It mostly occurs over dry and/or open habitats, including riparian woodland and tea-tree swamps, low scrub, heathland or saltmarsh, grassland, spinifex sandplains, farmland and sand-dunes. It sometimes occurs above forests. It probably roosts aerially, but has occasionally been observed to land {Higgins, 1999 #531}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<b>High</b> Likely to occur periodically in or above all habitats.
Calidris (Crocethia) alba	Sanderling	М	SLC	A coastal species found on low and open sand beaches exposed to open sea-swells. A migratory species, it has been recorded in NSW from September to May {Pizzey, 1997 #24}.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present.
Calidris acuminata	Sharp-tailed Sandpiper	М	SLC	Occurs in a variety of habitats: tidal mudflat, mangrove swamps, saltmarshes, shallow fresh, brackish, salt inland swamps and lakes; flooded and irrigated paddocks, sewage farms and commercial saltfields {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur periodically in wetland habitats.
Calidris canutus	Red Knot	E (M)	SLC	In Australasia the Red Knot mainly inhabit intertidal mudflats, sandflats and sandy beaches of sheltered coasts, in estuaries, bays, inlets, lagoons and harbours; sometimes on sandy ocean beaches or shallow pools on exposed wave-cut rock platforms or coral reefs. They are occasionally seen on terrestrial saline wetlands near the coast, such as lakes, lagoons, pools and pans, and recorded on sewage ponds and saltworks, but rarely use freshwater swamps. They rarely use inland lakes or swamps {Higgins, 1996 #648}.	PMST - Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Calidris ferruginea	Curlew Sandpiper	М	E	Occurs in inter-tidal mudflats of estuaries, lagoons, mangrove channels and also around lakes, dams, floodwaters and flooded saltbush surrounding inland lakes {Morcombe, 2003 #992}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur periodically in wetland habitats.
Calidris ruficollis	Red-necked Stint	Μ	SLC	Mostly found in coastal areas, including sheltered inlets, bays lagoons and estuaries. They also occur in shallow wetlands near the coast or inland, including lakes, waterholes and dams {Higgins, 1996 #648}. They forage in mudflats, shallow water, sandy open beaches, flooded paddocks and in samphire feeding along the edges. The species roosts on sheltered beaches, spits, banks or islets, of sand, mud, coral or shingle. Occasionally they roost on exposed reefs or shoals {Higgins, 1996 #648} and amongst seaweed, mud and cow-pats {Hobbs, 1961 #3753}. During high tides they may also use sand dunes and claypans.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur periodically in wetland habitats.
Calidris tenuirostris	Great Knot	CE (M)	SLC	Generally a coastal species found on tidal mudflats and sandy ocean shores. A migratory species visiting Australian waters between September and March {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Calyptorhynchus lathami erebus	Glossy Black- Cockatoo	-	V	Calyptorhynchus lathami erebus is occurs in the north and central east coast of Queensland. This subspecies ranges from the Dawson-Mackenzie-Isaac Rivers basin, north to the Connors-Clarke Ranges, south to Dawes and Many Peaks Ranges, and inland to the Expedition, Peak and Denham Ranges, including the Blackdown Tableland. It prefers woodland areas dominated by she-oak Allocasuarina, or open sclerophyll forests and woodlands with a stratum of Allocasuarina beneath Eucalyptus, Corymbia or Angophora. An obligate hollow nester, glossy black-cockatoos require large old trees (living or dead), usually eucalypts, for breeding. As such, nesting sites are mainly in areas containing large old trees {C.Hourigan, 2012 #7528}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in habitats supporting Casuarina and Allocasuarina flora species.
Casuarius casuarius johnsonii (southern population)	Southern Cassowary	E	Е	Although occurring primarily in rainforest and associated vegetation, the cassowary also uses woodland, swamp and disturbed habitats for a year-round supply of fleshy fruits. It occurs in three broad populations. In the Wet Tropics it is distributed widely from Cooktown to just north of Townsville. Core habitat is coastal lowlands between Ingham and Mossman, and uplands in the southern Atherton Tablelands and other ranges. On Cape York, it occurs as two disjunct populations in vine-forest communities: one in MacIlwraith and Iron Ranges, the other in Shelburne Bay {P.Latch, 2007 #7529}.	PMST Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in SEVT, open forest and woodland habitats.
Charadrius leschenaultii	Greater Sand Plover	V (M)	SLC	Entirely coastal in NSW foraging on intertidal sand and mudflats in estuaries, and roosting during high tide on sand beaches or rocky shores. A migratory species it is found in New South Wales generally during the summer months {Pizzey, 2007 #24}.	PMST Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Charadrius mongolus	Lesser Sand Plover	E (M)	SLC	Migratory bird that migrates from the northern hemisphere to coastal areas of northern and east coast of Australia {Garnett, 2000 #21}. The species is almost strictly coastal during the non-breeding season, preferring sandy beaches, mudflats of coastal bays and estuaries, sand-flats and dunes near the coast, occasionally frequenting mangrove mudflats (IUCN Redlist entry).	PMST Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present.
Charadrius ruficapillus	Red-capped Plover	М	SLC	A shoreline species inhabiting both coastal and inland wetland habitats; including broad sandy and shelly beaches, intertidal flats, broad flat margins of saline and freshwater lakes and rivers, saltmarsh, dues and occasionally shallow coastal fresh or saline wetlands {Pizzey, 2012 #3921}. It is distributed widely throughout Australia and Tasmania where suitable habitats occur {Pizzey, 2012 #3921}.	LGA	<i>Moderate</i> May occur periodically in wetland habitats.
Charadrius veredus	Oriental Plover	Μ	SLC	Oriental Plovers are found in coastal habitats, including estuarine mudflats and sandbanks, on sandy or rocky ocean beaches, nearby reefs, or near-coastal grasslands. They also disperse further inland inhabiting flat, open, semi-arid or arid grasslands, where the grass is short and sparse, and interspersed with hard, bare ground, such as claypans, dry paddocks, playing fields, lawns and cattle camps or open areas that have been recently burnt. Oriental Plovers may move to lightly wooded grasslands in wet season and sometimes roost on soft wet mud or in shallow waters of ocean or mudflats and also occasionally in dry, open habitats, such as saltmarsh or paddocks {McCrie, 1984 #3674}{Park, 1983 #3676}{Patterson, 1983 #3677}.	PMST Etheridge LGA Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur periodically in wetland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Chlidonias leucopterus	White-winged Black Tern	Μ	SLC	In Australia, and elsewhere in their non-breeding range, the species mostly inhabits fresh, brackish or saline, and coastal or subcoastal wetlands. It frequents tidal wetlands, such as harbours, bays, estuaries and lagoons, and their associated tidal sandflats and mudflats. Terrestrial wetlands, including swamps, lakes, billabongs, rivers, floodplains, reservoirs, saltworks, sewage ponds and outfalls are also inhabited. They rarely occur on inland wetlands in Australia {Chan, 2007 #3655}{Chan, 2008 #3656}{Chatto, 2006 #3657}{Cramp, 1985 #3658}{Dening, 2003 #3659}{Gochfeld, 1996 #3660}{Higgins, 1996 #648}{Johnstone, 1998 #3661}{Urban, 1986 #3662}.	Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur periodically in wetland habitats.
Cuculus opatus (syn. Cuculus saturatus)	Oriental Cuckoo, Himalayan Cuckoo	М	SLC	A non-breeding migrant to Australia, it often inhabits rainforest, vine thickets, wet sclerophyll forest and open woodland and sometimes occurs in mangroves, wooded swamps and as vagrants in gardens {Higgins, 1999 #531}. The population trend appears to be stable {BirdLife International, 2009 #3694}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	High Likely to occur in open forest to woodland habitats.
Cyclopsitta diophthalma macleayana	Macleay's Fig- Parrot	-	V	Found in upland and lowland rainforests, usually containing fig trees (Ficus); including dry cool subtropical and littoral rainforests, tropical semi-deciduous vine forests and gallery forests. Usually in large tracts of forest, particularly near edges, and rarely in partly cleared and fragmented rainforest {Higgins, 1999 #531}.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in SEVT along the great dividing range.
Epthianura crocea macgregori	Yellow Chat (Dawson)	CE	E	Inhabits marine plain wetlands that experience inundations of both fresh and saltwater tidal influences (Barnard 1913; Houston et al. 2004a, 2004b; Jaensch et al. 2004a). Specifically occupies areas that contain networks of shallow drainage channels and nearby grassland depressions. It breeds, shelters and forages in grasslands and dense beds of rush and sedge, but it also forages in more open habitats nearby, especially more sparse grasslands and Samphire vegetation (Houston et al. 2004a, 2004b; Jaensch et al. 2004a).	Wildlife Online - Etheridge LGA	<i>Moderate</i> May occur in wetland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Erythrotriorchis radiatus	Red Goshawk	V (M)	E	Lives in coastal and sub-coastal tall open forests and woodlands, tropical savannas traversed by wooded or forested rivers and along edges of rainforest. Nests are only built in trees taller than 20 meters which occur within 1 kilometre of a watercourse or wetland. Has a home range of 200 square kilometres and hunts for medium to large birds in open forests and gallery forest {Garnett, 2000 #21}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in open forest riparian habitats (BVG 16 and subsets)
Erythrura gouldiae	Gouldian Finch	E	E	The Gouldian Finch inhabits open woodlands that are dominated by Eucalyptus trees and support a ground cover of Sorghum and other grasses. It has also been recorded in undescribed thickets of vegetation along streams and gorges, and at the margins of stands of mangroves. It sometimes occurs around homesteads and townships. The Gouldian Finch drinks regularly and thus is often seen at watering points and associated habitat such as beds of grass and grass- covered banks around shallow waterholes, watercourses, soaks and springs {Department of the Environment and Energy, 2017 #7530}.	PMST - Charters Towers LGA, Etheridge LGA Wildlife Online - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> May occur in grassland and open grassy woodland habitats near water.
Erythrura trichroa	Blue-faced Parrot- finch	-	NT	The Blue-faced Parrot Finch inhabits Hill forest and montane forest and forest edge, rainforest, and dense secondary growth often observed eating the seeds of introduced grasses such as Signal Grass ( <i>Brachiaria decumbens</i> ) and Guinea or Hamil Grass ( <i>Megathyrsus maximus</i> ).	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in SEVT toward range.
Esacus neglectus	Beach Stone- curlew	CE	-	Found on beaches within species range, including short stretches of muddy sand among mangroves, coralline sands on atolls and prime surf beaches. Does not occupy long stretches of continuous mangroves or cliffs though beaches associated with estuaries or near mangroves are favoured {Garnett, 2000 #21}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Falco hypoleucos	Grey Falcon	-	V	Generally centred on inland drainage systems where the average rainfall is less than 500 millimetres. It is found in timbered lowland plains that are crossed by tree-lined water courses. Nests in the old nests of other birds, particularly raptors {Garnett, 2000 #21}.	Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in open forest riparian habitats (BVG 16 and subsets)
Gallinago hardwickii	Latham's Snipe	М	SLC	Occurs in freshwater or brackish wetlands generally near protective vegetation cover. This species feeds on small invertebrates, seeds and vegetation. It migrates to the northern hemisphere to breed {Garnett, 2000 #21}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<b>High</b> Likely to occur in wetland habitats.
Gallinago megala	Swinhoe's Snipe	Μ	SLC	During the non-breeding season Swinhoe's Snipe occurs at the edges of wetlands, such as wet paddy fields, swamps and freshwater streams. The species is also known to occur in grasslands, drier cultivated areas (including crops of rapeseed and wheat) and market gardens. Habitat specific to Australia includes the dense clumps of grass and rushes round the edges of fresh and brackish wetlands. This includes swamps, billabongs, river pools, small streams and sewage ponds. They are also found in drying claypans and inundated plains pitted with crab holes {Environment, 2015 #3579}.	PMST - Hinchinbrook LGA	<b>High</b> Likely to occur in wetland habitats.
Gallinago stenura	Pintail Snipe	М	SLC	During non-breeding period the Pin-tailed Snipe occurs most often in or at the edges of shallow freshwater swamps, ponds and lakes with emergent, sparse to dense cover of grass/sedge or other vegetation. The species is also found in drier, more open wetlands such as claypans in more arid parts of species' range. It is also commonly seen at sewage ponds; not normally in saline or inter-tidal wetlands {Department of the Environment, 2015 #3580}.	PMST - Hinchinbrook LGA	High Likely to occur in wetland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Gelochelidon nilotica (syn. Sterna nilotica)	Gull-billed Tern	Μ	SLC	Prefer shallow, often ephemeral, terrestrial wetlands, either fresh or saline, especially lakes, swamps and lagoons, particularly those with mudflats; sometimes on inundated ground, including saltpans, claypans and saltmarsh or watercourses and associated floodplains. Also occur in sheltered coastal embayments, estuaries and river deltas with tidal sandflats, mudflats or beaches. Inland, often occur well away from water, on dry samphire, grassy plains or even gibber. Usually breed on large, often ephemeral, inland lakes and swamps, on low exposed islands, banks, flats or spits of dry mud, sand or, occasionally, rocks; either bare or vegetated with sparse dry grass, reeds and rushes or scattered samphire {Higgins, 1997 #2274}.	Wildlife Online - Etheridge LGA, Hinchinbrook LGA	High Likely to occur in wetland habitats.
Geophaps scripta scripta	Squatter Pigeon (Southern Subspecies)	V	V	The Squatter Pigeon (southern) occurs mainly in grassy woodlands and open forests that are dominated by eucalypts. It has also been recorded in sown grasslands with scattered remnant trees, disturbed habitats (i.e. around stockyards, along roads and railways, and around settlements), in scrub and acacia growth, and remains common in heavily-grazed country north of the Tropic of Capricorn. The species is commonly observed in habitats that are located close to bodies of water {Department of Sustainability Environment Water Population and Communities, 2011 #3366}.	PMST - Charters Towers LGA, Wildlife Online - Charters Towers LGA	High Likely to occur in grassland and open grassy woodland habitats.

# C-11

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Glareola maldivarum	Oriental Pratincole	Μ	SLC	Within Australia the Oriental Pratincole is widespread in northern areas, especially along the coasts of the Pilbara Region and the Kimberley Division in Western Australia, the Top End of the Northern Territory, and parts of the Gulf of Carpentaria. It is also widespread but scattered inland, mostly north of 20° S. In non-breeding grounds in Australia, the Oriental Pratincole usually inhabits open plains, floodplains or short grassland (including farmland or airstrips), often with extensive bare areas. They often occur near terrestrial wetlands, such as billabongs, lakes or creeks, and artificial wetlands such as reservoirs, saltworks and sewage farms, especially around the margins. The species also occurs along the coast, inhabiting beaches, mudflats and islands, or around coastal lagoons {Department of the Environment and Energy, 2017 #7531}.	PMST - Etheridge LGA	<i>Moderate</i> Likely to occur in grassland, grazing land and open grassy woodland habitats.
Grantiella picta	Painted Honeyeater	V	V	Lives in dry forests and woodlands. Primary food is the mistletoes in the genus Amyema, though it will take some nectar and insects. Its breeding distribution is dictated by presence of mistletoes which are largely restricted to older trees. Less likely to be found in in strips of remnant box- ironbark woodlands, such as occur along roadsides and in windbreaks, than in wider blocks {Garnett, 2000 #21}.	PMST - Charters Towers LGA, Wildlife Online - Etheridge LGA	High Likely to occur in forests to open grassy woodland habitats.
Hirundapus caudacutus	White-throated Needletail	М	SLC	Occurs in airspace over forests, woodlands, farmlands, plains, lakes, coasts and towns. Breeds in the northern hemisphere and migrates to Australia in October-April {Pizzey, 2007 #24}.	PMST - Hinchinbrook LGA, Charters Towers LGA Wildlife Online - Hinchinbrook LGA, Etheridge LGA	<b>High</b> Likely to occur periodically in or above all habitats.
Hirundo rustica	Barn Swallow	М	SLC	Usually found in airspace over open grassland and wetland habitats such as ponds, freshwater wetlands swimming pools, coastal lagoons and tidal pools. This species is a regular visitor to northern Australia in Qld, NT and WA. This species has been occasional records in NSW at Newcastle, Mullumbimby and Nowra {Higgins, 2006 #2278}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	High Likely to occur periodically in or above all habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Hydroprogne caspia (syn. Sternia caspia)	Caspian Tern	Μ	SLC	The Caspian Tern is found in sheltered coastal embayments preferring sandy or muddy margins. Also found in near-coastal or inland terrestrial wetlands. It forages in open wetlands, preferring sheltered shallow water near the margins. It usually breeds in low islands, cays, spits, banks, ridges, beaches of sand or shell, terrestrial wetlands and stony or rocky islets or banks and occasionally among beach-cast debris above the high-water mark or at artificial sites, including islands in reservoirs, or on dredge-spoil. Generally roosting occurs on bare exposed sand or shell spits, banks or shores. {Higgins, 1996 #648}.	Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.
Limosa lapponica	Bar-tailed Godwit	М	SLC	Occurs in coastal mudflats, sandbars, shores of estuaries, salt marsh and sewage ponds {Morcombe, 2003 #992}.	PMST - Hinchinbrook LGA, Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present.
Limosa limosa	Black-tailed Godwit	М	SLC	A coastal species found on tidal mudflats, swamps, shallow river margins and sewage farms. Also found inland on larger shallow fresh or brackish waters. A migratory species visiting Australia between September and May {Pizzey, 2007 #24}.	PMST - Charters Towers LGA Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	Low Coastal habitats not present.
Monarcha frater	Black-winged Monarch	Μ	SLC	Found along the coast of eastern Australia, becoming less common further south. It is found in rainforests, eucalypt woodlands, coastal scrub and damp gullies. It may be found in more open woodland when migrating.	Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur in SEVT toward range.
Monarcha melanopsis	Black-faced Monarch	М	SLC	Occurs in rainforests, eucalypt woodlands, coastal scrubs, damp gullies in rainforest, eucalypt forest and in more open woodland when migrating {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur periodically in open forest to woodland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Motacilla cinerea	Grey Wagtail	Μ	SLC	A scarce but regular visitor to northern Australia, generally arriving in October and departing around March. The species has a strong association with water. In their normal breeding range, Grey Wagtails are found across a variety of wetlands, especially water courses, but also on the banks of lakes and marshes, as well as artificial wetlands such as sewage farms, reservoirs and fishponds. This association with water extends into non-breeding habitats with all confirmed Australian records being associated with water; especially creeks, rivers and waterfalls. On migration they may forage on rocky tidal flats.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland and riverine habitats.
Motacilla flava	Yellow Wagtail	Μ	SLC	This species occurs in a range of habitats including estuarine habitats such as sand dunes, mangrove forests and coastal saltmarshes. This species also occurs in open grassy areas including disturbed sites such as sports grounds and has been recorded on the edges of wetlands, swamps, lakes and farm dams. This species migrates from Asia to Australia in spring- summer. It has been recorded in the estuarine areas of the Hunter River in Newcastle NSW and in QLD and the north of NT and WA {Higgins, 2006 #2278}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present.
Myiagra cyanoleuca	Satin Flycatcher	М	SLC	Widespread in eastern Australia. In Queensland, it is widespread but scattered in the east. In NSW, they are widespread on and east of the Great Divide and sparsely scattered on the western slopes, with very occasional records on the western plains. In Victoria, the species is widespread in the south and east. Inhabit heavily vegetated gullies in eucalypt-dominated forests and taller woodlands, and on migration, occur in coastal forests, woodlands, mangroves and drier woodlands and open forests. Satin Flycatchers mainly inhabit eucalypt forests, often near wetlands or watercourses. They generally occur in moister, taller forests, often occurring in gullies. They also occur in eucalypt woodlands with open understorey and grass ground cover, and are generally absent from rainforest {Department of the Environment, 2016 #7313}{Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	High Likely to occur in riparian forests, open forests to woodlands.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Neochmia ruficauda ruficauda	Star Finch (Eastern Subspecies)	E	E	The Star Finch (eastern) occurs mainly in grasslands and grassy woodlands that are located close to bodies of fresh water. It also occurs in cleared or suburban areas such as along roadsides and in towns {Department of Sustainability Environment Water Population and Communities, 2011 #3375}.	PMST - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> May occur in grassland and open grassy woodland habitats near water.
Numenius madagascariensis	Eastern Curlew	CE (M)	V	Inhabits coastal estuaries, mangroves, mud flats and sand pits. It is a migratory shorebird which generally inhabits sea and lake shore mud flats, deltas and similar areas, where it forages for crabs and other crustaceans, clam worms and other annelids, molluscs, insects and other invertebrates. Its migration route ranges from its wintering grounds in Australia to its breeding grounds in northern China, Korea and Russia {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present.
Numenius minutus	Little Curlew	М	SLC	On passage the species shows a preference for foraging and resting in swampy meadows near lakes and along river valleys. It overwinters on dry inland grassland, bare cultivation, dry mudflats and coastal plains of black soil with scattered shallow pools of freshwater, swamps, lakes or flooded ground. It shows a preference for short grass swards of less than 20 cm tall, and occasionally occurs in dry saltmarshes, coastal swamps, mudflats or sandflats in estuaries, or on the beaches of sheltered coasts {BirdLife International, 2009 #3757}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.
Numenius phaeopus	Whimbrel	Μ	SLC	Migrates to Taiwan, Philippines, PNG, and a race breeding in NE Siberia is found on the north and south-eastern coastlines of Australia. Juveniles arrive to Australia from spring to early summer. Usually only juveniles remain in Australia but very occasionally adults in breeding plumage may be seen in Australian winters {Pizzey, 2007 #24}.	PMST - Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Pandion cristatus (syn. P. haliaetus)	Eastern Osprey	М	SLC	Generally a coastal species, occurring in estuaries, bays, inlets, islands and surrounding waters, coral atolls, reefs, lagoons, rock cliffs and stacks. Sometimes ascends larger rivers to far inland. Builds nests high in tree, on pylon or on ground on islands. Feeds on fish {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Plegadis falcinellus	Glossy Ibis	М	SLC	It feeds in very shallow water and nests in freshwater or brackish wetlands with tall dense stands of emergent vegetation (e.g. reeds or rushes) and low trees or bushes. It shows a preference for marshes at the edges of lakes and rivers, as well as lagoons, flood-plains, wet meadows, swamps, reservoirs, sewage ponds, rice-fields and irrigated cultivation. It less often occurs in coastal locations such as estuaries, deltas, saltmarshes and coastal lagoons. Roosting sites are often large trees that may be far from water. The nest is a platform of twigs and vegetation usually positioned less than 1 m above water in tall dense stands of emergent vegetation (e.g. reeds or rushes), low trees or bushes over water {BirdLife International, 2009 #3757}.	Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<b>High</b> Likely to occur in wetland habitats.
Pluvialis fulva	Pacific Golden Plover	М	SLC	Prefers sandy, muddy or rocky shores, estuaries and lagoons, reefs, saltmarsh, and or short grass in paddocks and crops. The species is usually coastal, including offshore islands; rarely far inland. Often observed on beaches and mudflats, sandflats and occasionally rock shelves, or where these substrates intermingle; harbours, estuaries and lagoons {Higgins, 1993 #534}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Pluvialis squatarola	Grey Plover	Μ	SLC	In non-breeding grounds in Australia, Grey Plovers occur almost entirely in coastal areas, where they usually inhabit sheltered embayments, estuaries and lagoons with mudflats and sandflats, and occasionally on rocky coasts with wave-cut platforms or reef-flats, or on reefs within muddy lagoons. They also occur around terrestrial wetlands such as near-coastal lakes and swamps, or salt-lakes. The species is also very occasionally recorded further inland, where they occur around wetlands or salt-lakes {Marchant, 1993 #534}. They usually forage on large areas of exposed mudflats and beaches and occasionally in pasture and on muddy margins of inland wetlands {Marchant, 1993 #534}. They usually roost in sandy areas, such as on unvegetated sandbanks or sand-spits on sheltered beaches or other sheltered environments {Jaensch, 1988 #3754}{Pegler, 1983 #3755}.	PMST - Hinchinbrook LGA Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Poephila cincta cincta	Black-Throated Finch (White- Rumped Subspecies)	E	E	Occurs in grassy woodland dominated by eucalypts, paperbarks or acacias where there are seeding grasses and water. Nests in tree hollows or builds domed nests in trees {Garnett, 2000 #21}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	High Likely to occur in grassy open woodland habitats near water. Previous records in corridor options.
Rhipidura rufifrons	Rufous Fantail	М	SLC	Occurs in a range of habitats including the undergrowth of rainforests/wetter eucalypt forests/gullies, monsoon forests paperbarks, sub-inland and coastal scrubs, mangroves, watercourses, parks and gardens. When migrating they may also be recorded on farms, streets and buildings. Migrates to SE Australia in October-April to breed, mostly in or on the coastal side of the Great Dividing Range {Pizzey, 2007 #24}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA	<b>High</b> Likely to occur in riparian and moist forest habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Rostratula australis (syn. R. benghalensis)	Australian Painted Snipe (Painted Snipe)	V	V	Inhabits shallow, vegetated, temporary or infrequently filled wetlands, including where there are trees such as Eucalyptus camaldulensis (River Red Gum), E. populnea (Poplar Box) or shrubs such as Muehlenbeckia florulenta (Lignum) or Sarcocornia quinqueflora (Samphire). Feeds at the water's edge and on mudlflats on seeds and invertebrates, including insects, worms, molluscs and crustaceans. Males incubate eggs in a shallow scrape nest {Garnett, 2000 #21}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Etheridge LGA	<b>High</b> Likely to occur in wetland habitats.
Sterna albifrons	Little Tern	М	Е	Little Terns inhabit sheltered coastal environments, including lagoons, estuaries, river mouths and deltas, lakes, bays,	PMST - Hinchinbrook LGA,	Low
				harbours and inlets. They nest on sand-spits, sandbanks, ridges or islets in these habitats or gently sloping sandy ocean beaches and occasionally in sand-dunes {Garnett, 2000 #21}.	Wildlife Online - Hinchinbrook LGA	Coastal habitats not present
Sterna dougallii	Roseate Tern	М	SLC	In Queensland, scattered records occur in the south-east Gulf of Carpentaria and west Cape York Peninsula, but birds are possibly more widespread, with large numbers nesting on south-east Bountiful Island. Birds are widespread along the east coast of Australia, south to about Hervey Bay. They are more sparsely distributed, further south, occasionally reaching north Fraser Island. It occurs in coastal and marine areas in subtropical and tropical seas. The species inhabits rocky and sandy beaches, coral reefs, sand cays and offshore islands. Birds rarely occur in inshore waters or near the mainland, usually venturing into these areas only accidentally, when nesting islands are nearby {Higgins, 1997 #2274}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Sterna hirundo	Common Tern	М	SLC	A non-breeding migrant to Australia, occurring mainly on the east coast and inhabiting marine, pelagic and coastal habitats. Mostly oceanic but often recorded in bays, harbours and estuaries and occasionally in coastal wetlands. Roosting occurs on unvegetated intertidal sandy ocean beaches, shores of estuaries, lagoons and sand bars {Higgins, 1996 #648}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Sterna sumatrana	Black-naped Tern	Μ	SLC	In Australia, Black-naped Terns are found mainly in the central north and north-east of the country, in central and eastern Northern Territory, the Gulf of Carpentaria and Torres Strait and through the islands and waters of the Great Barrier Reef and Coral Sea. The species is rarely found in inshore waters except when breeding. Black-naped Terns breed and roost on islands, which are very occasionally close to or attached to the mainland at low tides, and forage in seas surrounding colonies. Black-naped Terns are mainly associated with small, offshore sand and coral cays, coral reefs and lagoons, and sandy and rocky islands and islets, and in the surrounding seas. The species is only occasionally recorded in inshore waters away from their breeding colonies or coastal mainland Australia, such as harbours or bays, with their occurrence inshore probably being influenced by climatic conditions {Higgins, 1997 #2274}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Tringa brevipes (syn. Heteroscelus brevipes)	Grey-tailed Tattler	Μ	SLC	It is often found on sheltered coasts with reefs, rock platforms or with intertidal mudflats. It is also found at intertidal rocky, coral or stony reefs, platforms and islets that are exposed at low tide. It has also been found in embayments, estuaries and coastal lagoons, especially fringed with mangroves. It is rarely seen on open beaches and occasionally found around near- coastal wetlands, such as lagoons, lakes and ponds in sewage farms and saltworks. Inland records for the species are rare {Higgins, 1996 #648}. The species forages in shallow water, hard intertidal substrates, rock pools, intertidal mudflats, mangroves, banks of seaweed and among rocks and coral rubble, over which water may surge. The species roosts in mangroves, dense stands of shrubs, snags, rocks, beaches, reefs, artificial structures (sea walls, oyster racks), occasionally in near-coastal saltworks and sewage ponds and rarely on sandy beaches or sand banks {Higgins, 1996 #648}{Rogers, 1999 #3758}.	PMST - Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Tringa glareola	Wood Sandpiper	Μ	SLC	Found in well-vegetated, shallow, freshwater wetlands, such as swamps, billabongs, lakes, pools and waterholes. They are typically associated with emergent, aquatic plants or grass, and dominated by taller fringing vegetation, such as dense stands of rushes or reeds, shrubs, or dead or live trees and often with fallen timber. They also inhabit inundated grasslands, short herbage or wooded floodplains, where floodwaters are temporary or receding, and irrigated crops. This species uses artificial wetlands, including open sewage ponds, reservoirs, large farm dams, and bore drains and occasionally found in stony wetlands. The species forages on mud at the edges of wetlands, either along shores, among open scattered aquatic vegetation, or in clear shallow water {Higgins, 1996 #648}.	PMST - Charters Towers LGA, Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.
Tringa incana	Wandering Tattler	CE (M)	SLC	Generally found on rocky coasts with reefs and platforms, points, spits, piers, offshore islands and shingle beaches or beds. Occasionally seen on coral reefs or beaches, and tends to avoid mudflats Foraging habitat is among rocks or shingle, or in shallow pools at edges of reefs or beaches, mainly along the tideline. Wandering Tattlers have been recorded roosting or perching on top of boulders surrounded by or close to water {Higgins, 1996 #648}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Tringa nebularia	Common Greenshank	М	SLC	Occurs in a range of inland and coastal environments. Inland, it occurs in both permanent and temporary wetlands, billabongs, swamps, lakes floodplains, sewage farms, saltworks ponds, flooded irrigated crops. On the coast, it occurs in sheltered estuaries and bays with extensive mudflats, mangrove swamps, muddy shallows of harbours and lagoons, occasionally rocky tidal ledges. It generally prefers wet and flooded mud and clay rather than sand {Morcombe, 2003 #992}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Tringa stagnatilis	Marsh Sandpiper	М	SLC	Occurs in coastal and inland wetlands (salt or fresh water), estuarine and mangrove mudflats, beaches, shallow or swamps, lakes, billabongs, temporary floodwaters, sewage farms and saltworks ponds {Morcombe, 2003 #992}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in wetland habitats.
Turnix olivii	Buff-breasted Button-quail	E	E	Poorly recorded species occurring in Northeast Queensland, from Iron Range and near Coen, Cooktown, Musgrave, Mount Molloy, Mareeba, Chillagoe and Ingham, although most recent records are form the Mt Molloy area. Said to prefer sparse short grassy areas on stony ground, and avoiding densely grassed areas (DoE 2015)	Wildlife Online - Hinchinbrook LGA	<i>Moderate</i> May occur in rocky open grassy habitats.
Tyto novaehollandiae kimberli	Masked Owl (northern)	V	V	In Queensland, there are historical records from the Normanton region, and from Pascoe, Archer, Chester and Watson Rivers on Cape York Peninsula. It occurs along the southern rim of the Gulf of Carpentaria, Cape York Peninsula and south to Atherton Tablelands and the Einasleigh-Burdekin divide. Individuals have been recorded from riparian forest, rainforest, open forest, Melaleuca swamps and the edges of mangroves, as well as along the margins of sugar cane fields {Higgins, 1999 #531}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in riparian forest and open forest to woodland habitats.
Xenus cinereus	Terek Sandpiper	М	SLC	Found on tidal mudflats and estuaries and on shores and reefs of offshore islands {Pizzey, 2007 #24}.	PMST - Hinchinbrook LGA, Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Coastal habitats not present

## C-21

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Fish						
Pristis pristis ( syn. Pristis microdon)	Freshwater Sawfish	V	-	The Freshwater Sawfish may potentially occur in all large rivers of northern Australia from the Fitzroy River, Western Australia, to the western side of Cape York Peninsula, Queensland. It is mainly confined to the main channels of large rivers. It is a marine/estuarine species that spends its first three–four years in freshwater growing to about half its adult size (4 m+). Juveniles and sub-adult Freshwater Sawfish predominantly occur in rivers and estuaries, while large mature animals tend to occur more often in coastal and offshore waters up to 25 m depth. The preferred habitat of this species is mud bottoms of river embayments and estuaries, but they are also found well upstream. They are not found near riparian vegetation. They are usually found in turbid channels of large rivers over soft mud bottoms more than 1 m deep, but they will move into shallow waters when travelling upstream or while hunting prey {Department of the Environment and Energy, 2017 #7522}		Low Unlikely to occur in riverine habitats.
Invertebrates						
Hypochrysops apollo apollo	Apollo Jewel (Wet Tropics Subspecies)	-	V	Shares a mutualistic relationship with Golden Ant (Iridiomyrmex cordatus) and Ant Plant (Myrmecodia beccarii), which occur in coastal woodlands between Cooktown and Ingham in Queensland.	Wildlife Online - Hinchinbrook LGA	Low Outside of species known range

### C-22

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Mammals						
Bettongia tropica	Northern Bettong	E	Е	The Northern Bettong currently occurs in three geographically isolated locations: the Lamb Range, Paluma and Mt Zero, Queensland. Mt Windsor Tableland was known to have an existing population as recently as January 1989. However, despite considerable effort no Northern Bettongs have been seen since this time. Its preferred habitat is tall and medium open eucalypt forest with grassy understorey. These habitat types occur as a narrow fragmented strip along the western edge of wet tropical rainforests {Department of the Environment and Energy, 2017 #7533}.	PMST - Charters Towers LGA, Hinchinbrook LGA, Wildlife Online - Charters Towers LGA	Low Outside of species known range
Dasyurus hallucatus	Northern Quoll	E	-	The Northern Quoll occupies a diversity of habitats across its range which includes rocky areas, eucalypt forest and woodlands, rainforests, sandy lowlands and beaches, shrubland, grasslands and desert. Northern Quoll are also known to occupy non rocky lowland habitats such as beachscrub communities in central Queensland. Northern Quoll habitat generally encompasses some form of rocky areas for denning purposes with surrounding vegetated habitats used for foraging and dispersal. Rocky habitats are usually of high relief, often rugged and dissected but can also include tor fields or caves in low lying areas such as in Western Australia. Eucalypt forest or woodland habitats usually have a high structural diversity containing large diameter trees, termite mounds or hollow logs for denning purposes. Dens are made in rock crevices, tree holes or occasionally termite mounds. Northern Quolls sometimes occur around human dwellings and campgrounds. Northern Quolls appear to be most abundant in habitats within 150 km of the coast {Department of Sustainability Environment Water Population and Communities, 2011 #3587}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	High Likely to occur in forest and woodland habitats supporting rocky areas for denning / breeding.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Dasyurus maculatus gracilis	Spotted-Tailed Quoll (Northern Subspecies)	E	E	The northern subspecies is now thought to be confined to two extant populations: one centered on the Windsor and Carbine Tablelands, Thornton Peak, Mount Finnegan and associated smaller ranges; and the other centered on the Atherton Tablelands and associated mountain ranges. Its habitat is mostly confined to the relatively cool, wet and climatically equable upland closed-forests (mostly above 900 m altitude) that occur in the upper catchments of rivers draining east and west of the Eastern Escarpment in the Wet Tropics bioregion of north-eastern Queensland. It utilises dens for resting and for raising young. Dens have been found in tree hollows, logs, rock crevasses and even among building materials. Maternal dens often have long entrances {Department of the Environment and Energy, 2017 #7534}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in forest to open woodland habitats in elevated areas.
Hipposideros diadema reginae	Diadem Leaf- Nosed Bat	-	NT	This subspecies is endemic to Queensland, occurring from Cape York Peninsula south to Townsville and inland to Chillagoe, with records from Iron Range, Hinchinbrook Island, Cape Melville, Chillagoe, Cairns and Coen. It roosts throughout the year in caves and disused mines, preferring those with large chambers, high domed ceilings and multiple entrances. They have also been recorded roosting in buildings and culverts. It occurs in a variety of habitat types including lowland rainforest, Melaleuca forests, eucalypt woodland, deciduous vine thickets, and open woodland; where suitable roosts are available throughout its range. Typical foraging sites are at vegetation edges or gaps adjacent to open space {C.Hourigan, 2011 #7535}.	Wildlife Online - Charters Towers LGA, Etheridge LGA	<i>Moderate</i> Likely to occur in forest and open woodland habitats on landforms that supporting cave roosting sites.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Hipposideros semoni	Semon's Leafnosed-bat	E	Е	Distribution for Semon's Leaf-nosed Bat includes coastal Queensland from Cape York to just south of Cooktown. Semon's Leaf-nosed Bat is found in tropical rainforest, monsoon forest, wet sclerophyll forest and open savannah woodland. This species does not have an obligatory requirement for cave roosts. Daytime roost sites include tree hollows, deserted buildings in rainforest, road culverts and shallow caves amongst granite boulders or in fissures {Department of the Environment and Energy, 2017 #7536}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in open forest habitats associated with hollow bearing trees and cave roosting sites.
Macroderma gigas	Ghost Bat	V	V	Ghost bats occur in a wide range of habitats from rainforest, monsoon and vine scrub, to open woodlands in arid areas. These habitats are used for foraging, while roost habitat is more specific. Favoured roosting sites of the ghost bat are undisturbed caves or mineshafts which have several openings {Department of Environment and Heritage Protection, 2013 #7230}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	<i>Moderate</i> Likely to occur in forest and open woodland habitats on landforms that supporting cave roosting sites.
Macrotis lagotis	Greater Bilby	V	E	This species is restricted predominantly to the Tanami Desert, Northern Territory, the Great Sandy and Gibson Deserts, Western Australia and an outlying population between Boulia and Birdsville in south-west Queensland. The species occurs in a variety of habitats, usually on landforms with level to low slope topography and light to medium soils. It occupies three major vegetation types; open tussock grassland on uplands and hills, mulga woodland/shrubland growing on ridges and rises, and hummock grassland in plains and alluvial areas{Department of the Environment, 2015 #3988}.	PMST - Charters Towers LGA	<b>High</b> Likely to occur in tussock grassland habitats.
Mesembriomys gouldii rattoides	Black-footed Tree-rat	V	-	It has been recorded mostly from eucalypt forests and woodlands (but not rainforests) around Mareeba, but there are records sparsely across Cape York Peninsula. In north Queensland, this species mostly occurs in eucalypt forests and woodlands, especially where hollows are relatively plentiful.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	Low Outside of species known range

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Murina florium	Tube-nosed Insectivorous Bat	-	V	Occurs within the wet tropics, and further north in the Iron Range on Cape York Peninsula. The limits to its distribution in Queensland are not well known. It has been recorded from a range of rainforest types, and from rainforest with emergent <i>Eucalyptus grandis</i> , at altitudes from near sea level to 1200m {Department of Environment and Heritage Protection, 2016 #7547}.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Rainforest habitats not present.
Onychogalea fraenata	Bridled Nailtail Wallaby	E	E	The species is recorded in habitats west of the Great Dividing Range in a mixture of tall shrubland and grassy woodland, and on the fertile soils which support open eucalypt forest and woodland, and Brigalow scrub. The species has a preference for scrub edges and adjacent vegetation, grazing and sheltering in the shrubland and grazing the grassy woodland {Strahan, 1995 #185}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in open grassy woodland, tussock grassland and woody shrublands.
Ornithorhynchus anatinus	Platypus	-	SLC	Inhabits freshwater creeks and occasionally lakes in the coastal ranges of eastern Australia and throughout Tasmania. Largely crepuscular in habits, but may be seen throughout the day. Shelters in burrows dug into stream banks and feeds on aquatic invertebrates.	Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<b>High</b> Likely to occur in waterways / streams.
Petauroides volans	Greater Glider	V	V	The Greater Glider has a restricted distribution in eastern Australia, from the Windsor Tableland in north Queensland to central Victoria, with an elevated range from sea level to 1200m above sea level. The species is largely restricted to eucalypt forests and woodlands, with a diet comprising of eucalypt leaves and occasional flowers. It is found in abundance in montane eucalypt forest with relatively old trees and an abundance of hollows. It also favours forests with a diversity of eucalypts to cater for seasonal variation in food abundance {Department of the Environment, 2015 #7227}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	High Likely to occur in riparian forest and open forest to woodland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Petaurus gracilis	Mahogany Glider	E	E	Restricted to the coastal southern Wet Tropics region of northern Queensland. They live in a narrow and highly fragmented band of lowland sclerophyll forest extending around 140 km from Toomulla, north of Townsville, to Tully and up to 40 km inland. Most recorded sightings have been at altitudes below 120 m. The main canopy and sub-canopy trees are eucalypts, bloodwoods and paperbarks and less commonly swamp mahogany and turpentine with an open mid- stratum of smaller trees and shrubs (e.g. wattles, forest siris, golden parrot tree, black she-oak, pandanus) and a grassy ground stratum in which grass trees may be present. The mahogany glider requires a relatively open forest structure for efficient gliding and tends to avoid dense vegetation such as rainforest {Department of Environment and Heritage Protection, 2017 #7548}.		<b>High</b> Likely to occur in riparian forest and open forest to woodland habitats.
Petrogale sharmani	Sharman's Rock- Wallaby	V	V	Sharman's rock-wallaby has a restricted distribution, centred on the Seaview and Coane Ranges, west of Ingham in north- east Queensland. It inhabits rocky slopes, rocky outcrops, boulder piles, cliff lines and gorges, usually associated with tropical woodland with a grassy understorey {Department of Environment and Heritage Protection, 2015 #7232}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	Low Outside of species known range.

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SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Phascolarctos cinereus	Koala (NSW, ACT & QLD - excluding SE QLD)	V	V	The Koala has a fragmented distribution throughout eastern Australia from north-east Queensland to the Eyre Peninsula in South Australia. In NSW it mainly occurs on the central and north coasts with some populations in the west of the Great Dividing Range. Inhabits eucalypt woodlands and forests. Koalas Feed on the foliage of more than 70 eucalypt species and 30 non-eucalypt species, but in any one area will select preferred browse species. The preferred tree species vary widely on a regional and local basis. Some preferred species include Forest Red Gum Eucalyptus tereticornis, Grey Gum E. punctata. In coastal areas, Tallowwood E. microcorys and Swamp Mahogany E. robusta are important food species, while in inland areas White Box E. albens, Bimble Box E. populnea and River Red Gum E. camaldulensis are favoured {NSW National Parks and Wildlife Service, 1999 #43; NSW National Parks and Wildlife Service, 2003 #31}{Office of Environment & Heritage, 2015 #7299}. Hawks Nest and Tea Gardens Population and population in the Pittwater LGA listed as Endangered under the NSW TSC Act{Office of Environment & Heritage, 2016 #7300}{Office of Environment & Heritage, 2013 #7301}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA	High Likely to occur in riparian forest and open forest to woodland habitats. Previous records in corridor options.
Pteropus conspicillatus	Spectacled Flying-fox	V	V	The Spectacled Flying-fox occurs in north-eastern Queensland, north of Cardwell with past records from Brisbane and Chillagoe. The Spectacled Flying-fox feeds on fruits and blossom, primarily in the canopy vegetation of a wide range of vegetation communities, including closed forest, gallery forest, eucalypt open forest and woodland, Melaleuca thickets, coastal swamps, mangroves, vegetation in urban settings, and commercial fruit crops. It roosts in large camps {Department of the Environment, 2015 #3675}.	PMST - Charters Towers LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA, Hinchinbrook LGA	<i>Moderate</i> May occur in riparian, open forest and woodland habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Pteropus poliocephalus	Grey-headed Flying-fox	V	-	Occurs in the coastal belt from Rockhampton in central Queensland to Melbourne in Victoria. The species is widespread throughout their range in summer, whilst in autumn it occupies coastal lowlands and is uncommon inland. Occurs in subtropical and temperate rainforests, tall sclerophyll forests and woodlands, heaths and swamps as well as urban gardens and cultivated fruit crops. Roosting camps are generally located within 20 km of a regular food source and are commonly found in gullies, close to water, in vegetation with a dense canopy. Feed on the nectar and pollen of native trees, in particular Eucalyptus, Melaleuca and Banksia, and fruits of rainforest trees and vines {Office of Environment & Heritage, 2015 #7341} {Department of the Environment, 2016 #7342}.	PMST - Hinchinbrook LGA	Low Outside of species known range.
Rhinolophus robertsi	Large-eared Horseshoe-bat	E	E	The Greater Large-eared Horseshoe Bat occurs only in northern Queensland, from the Iron Range southwards to Townsville and west to the karst regions of Chillagoe and Mitchell-Palmer. The Greater Large-eared Horseshoe Bat is found in lowland rainforest, along gallery forest-lined creeks within open eucalypt forest, Melaleuca forest with rainforest understorey, open savannah woodland and tall riparian woodland of Melaleuca, Forest Red Gum (E. tereticornis) and Moreton Bay Ash (E. tesselaris). Daytime roosting habitat for the Greater Large-eared Horseshoe Bat includes caves and underground mines located in rainforest, and open eucalypt forest and woodland. Roosts have also been observed in road culverts, and it is suspected that the species uses basal hollows of large trees, dense vegetation, rockpiles and areas beneath creekbanks {Department of the Environment and Energy, 2017 #7524} {Churchill, 2008 #6883}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in open forest habitats associated with hollow bearing trees and cave roosting sites.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Saccolaimus saccolaimus nudicluniatus	Bare-rumped Sheathtail Bat	CE	E	Known to occur in north-eastern Queensland and the monsoonal tropics of the Northern Territory. In Queensland, it occurs from Ayr to the Iron Range, including Magnetic and possibly Prince of Wales Islands. Most records are near- coastal mostly in eucalypt forests and woodlands. Known to be associated with coastal lowland rainforests, and more open forests dominated by Eucalyptus or Corymbia species interspersed with coastal lowland rainforest {Department of the Environment and Energy, 2017 #7539}.	HINCHINDIOOK LGA	Low Rainforest habitats not present.
Sminthopsis archeri	Chestnut Dunnart	-	NT	This predator was thought to be restricted to southern Papua New Guinea and Cape York, until it was recorded woodlands 200 km west of Townsville. It is found in open woodland to tall open forest and heathlands {Van Dyck and Strahan, 2008}.	Wildlife Online - Etheridge LGA	Moderate May occur in open woodland and tall open forest habitats
Sminthopsis douglasi	Julia Creek Dunnart	E	E	Endemic to north-western Queensland, where it occurs in the Mitchell Grass Downs and Desert Uplands bioregions. It is closely associated with tussock grasslands on cracking clay soils, with habitat quality associated particularly with increasing densities of cracks and holes, and with the extent and density of grass cover {Department of the Environment and Energy, 2017 #7539}.	PMST - Charters Towers LGA	Low Outside of species known range.
Tachyglossus aculeatus	Short-beaked Echidna	-	SLC	Occurs throughout most terrestrial habitats across mainland Australia and Tasmania, where its favoured ant and termite prey are available.	Wildlife Online - Etheridge LGA, Hinchinbrook LGA	High Likely to occur in all terrestrial habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Taphozous australis	Coastal Sheathtail-bat	-	V	Occurs along a very narrow coastal zone in Queensland from Shoalwater Bay, through Cape York Peninsula, to Moa Island in the Torres Strait. Taphozous australis also occurs on numerous coastal islands off Queensland throughout its range. Its presence depends on coastal roosts, preferring sea caves and rocky clefts. Also known to roost in disused mines, boulder piles, rock fissures, concrete bunkers, and occasionally in buildings. It forages above the canopy in areas of coastal dune scrubland, melaleuca swamps, open eucalypt forest, grasslands, coastal heathland, monsoon forests, and mangroves on lowlands and foothills {Department of the Environment and Energy, 2017 #7539}.	Wildlife Online - Hinchinbrook LGA	Low Coastal habitats not present
Xeromys myoides	False Water-rat	V	V	The only known False Water Rat populations in Australia are in coastal areas of the Northern Territory and Queensland. It is found in coastal wetlands such as lagoons, swamps and sedged lakes close to fore dunes. It forages amongst the mangroves at night when the tide is low, and when the tide rises it returns to the adjacent sedgelands for shelter. The False Water Rat builds large mud nests like termite mounds, up to 60 centimetres high and usually in areas where they can escape the highest of tides. They often use exposed tree roots to form the foundation for the mounds {Australian Government, 2003 #3581}	PMST - Hinchinbrook LGA	Low Coastal habitats not present

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SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	ΗΑΒΙΤΑΤ	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Reptiles						
Acanthophis antarcticus	Common Death Adder	-	V	The Common Death Adder occurs from the Gulf region of the Northern Territory across to central and eastern Queensland and New South Wales, and through to the southern parts of South Australia and Western Australia. Once abundant in many areas, this species has experienced a dramatic reduction in numbers. Within this range the species is found in a wide variety of habitats in association with deep leaf litter, including rainforests, wet sclerophyll forests, woodland, grasslands, chenopod dominated shrublands, and coastal heathlands {Department of Environment and Resource Management, 2011 #3598}.	Wildlife Online - Charters Towers LGA, Wildlife Online - Etheridge LGA	<i>Moderate</i> May occur in forest and SEVT habitats.
Delma mitella		V	NT	The Atherton Delma is known from the eastern side of the Atherton Tablelands in northeastern Queensland. The Atherton Delma is known only from tall open forests and rainforest interfaces in the Herberton, Ravenshoe and Paluma districts {Australian Government, 2008 #3979}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in open forest habitats
Denisonia maculata	Ornamental Snake	V	V	The species is known only from the Brigalow Belt North and parts of the Brigalow Belt South biogeographical regions. The core of the species' distribution occurs within the drainage system of the Fitzroy and Dawson Rivers. The Ornamental Snake's preferred habitat is within, or close to, habitat that is favoured by its prey - frogs. The species is known to occur in RE 11.4.3, 11.4.6, 11.4.8, 11.4.9, 11.3.3, and 11.5.16 {Department of the Environment, 2015 #3980}.	PMST - Charters Towers LGA, Etheridge LGA Wildlife Online - Charters Towers LGA	Low Outside of species known range (Brigalow Belt).

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Egernia rugosa	Yakka Skink	V	V	The Yakka Skink is commonly found in cavities under and between partly buried rocks, logs or tree stumps, root cavities and abandoned animal burrows. The species often takes refuge in large hollow logs and has been known to excavate deep burrow systems, sometimes under dense ground vegetation. In cleared habitat, this species can persist where there are shelter sites such as raked log piles, deep gullies, tunnel erosion/sinkholes and rabbit warrens. The species has also been found sheltering under sheds and loading ramps. This species is not generally found in trees or rocky habitats {Department of Sustainability Environment Water Population and Communities, 2011 #2273}.	PMST - Charters Towers LGA, Etheridge LGA, Hinchinbrook LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in forest to open woodland habitats.
Lampropholis mirabilis	-	-	NT	Distributed from Magnetic Island and parts of adjacent mainland, QLD. Found among granitic rocks at the edges of, or in clearings, rain- and monsoon forest, vine thickets and denser woodland habitats {Cogger, 2000 #20}.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in SEVT.
Lerista ameles	-	-	V	Known from one locality 30km east of Mt Surprise, QLD. Inhabits woodland in loose soil beneath rocks on low, weathered granite outcrops {Cogger, 2000 #20}.	Wildlife Online - Etheridge LGA	Low Outside of species known range
Lerista cinerea	Vine-thicket Fine- lined Slider	-	V	Occurs in leaf litter and loose soils in vine-thickets and adjacent forests (Greer 1983).	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in SEVT habitats.
Lerista hobsoni	Hobson's fine- liner slider	-	V	Found in scattered open forest localities in the Upper Burdekin Drainage Basin. In associated IBRA regions of Brigalow Belt North, Desert Uplands and Einasleigh Uplands.	Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in open forest habitats.
Lerista vanderduysi	Leaden-bellied fine-line slider	V	V	Typically found in semi-evergreen vine thickets.	Wildlife Online - Etheridge LGA	<i>Moderate</i> May occur in SEVT habitats.

SCIENTIFIC NAME	COMMON NAME	EPBC ACT	NC ACT	HABITAT	DATA SOURCE	LIKELIHOOD OF OCCURRENCE IN CORRIDOR OPTIONS
Lerista vittata	-	V	V	Lerista vittata was first described at Mount Cooper Station, approximately 80 km south-east of Charters Towers, Queensland. It has since been discovered 100–200 km NNW of Hughenden on the Chudleigh Plateau, though populations are fragmented and appear to be distinctively different. The species is found in Semi-evergreen Vine Thickets of the Brigalow Belt (North and South) and Nandewar Bioregions, an ecological community listed under the Environment Protection and Biodiversity Conservation Act 1999 as Endangered. It is also found more broadly, in spinifex communities as well as eucalyptus dominated woodlands with well-developed grassy understories {Department of the Environment, 2015 #3978}.	PMST - Charters Towers LGA, Etheridge LGA Wildlife Online - Charters Towers LGA	<i>Moderate</i> May occur in SEVT habitats.
Lygisaurus rococo (syn. Carlia rococo)	Chillagoe litter- skink	-	NT	Currently known only from limestone rock outcrops near Chillagoe, north-east QLD. Found amongst ground litter which has accumulated between rock outcrops {Cogger, 2000 #20}.	Wildlife Online - Etheridge LGA	Low Outside of species known range
Phyllurus gulbaru	Gulbaru Gecko	-	E	Has a very restricted distribution in Queensland, found in rocky slopes and gullies in rainforest habitat. The species is located near a UNESCO World Heritage site. The Gulbaru gecko was discovered in 2001. The species has only been found at three sites despite extensive survey {Department of the Environment and Energy, 2017 #7539}.	LGA	Low Outside of species known range

Key: CE = Critically Endangered, E = Endangered, V= Vulnerable, NT = Near Threatened, M = Migratory, SLC = Special Least Concern

