

Myco- and photobiont associations in crustose lichens in the McMurdo Dry Valleys (Antarctica) reveal high differentiation along an elevational gradient

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Abstract

The climate conditions of the McMurdo Dry Valleys (78° S) are characterized by low temperatures and low precipitation. The annual temperatures at the valley bottoms have a mean range from -30 °C to -15 °C and decrease with elevation. Precipitation occurs mostly in form of snow (3-50 mm a⁻¹ water equivalent) and, liquid water is rare across much of the landscape for most of the year and represents the primary limitation to biological activity. Snow delivered off the polar plateau by drainage winds, dew and humidity provided by clouds and fog are important water sources for rock inhabiting crustose lichens. In addition, the combination of the extremely low humidity and drying caused by foehn winds, confined to lower areas of the valleys, with colder and moister air at higher altitudes creates a strongly improving water availability gradient with elevation.

We investigated the diversity and interaction specificity of myco-/photobiont associations of a total of 232 crustose lichen specimens, collected along an elevational gradient (171-959 m a.s.l.) within the McMurdo Dry Valleys with regard to the spatial distribution caused by climatological and geographical factors. For the identification of the mycobiont and photobiont species three markers each were amplified (nrITS, mtSSU, RPB1 and nrITS, psbJ-L, COX2, respectively). Elevation, associated with a water availability gradient, turned out to be the key factor explaining most of the distribution patterns of the mycobionts. Pairwise comparisons showed *Lecidea cancriformis* and *Rhizoplaca macleanii* to be significantly more common at higher, and *Carbonea vorticosa* and *Lecidea polypycnidophora* at lower, elevations. Lichen photobionts were dominated by the globally distributed *Trebouxia* OTU, *Tr_A02* which occurred at all habitats. Network specialization resulting from mycobiont-photobiont bipartite network structure varied with elevation and associated abiotic factors.

Along an elevational gradient, the spatial distribution, diversity and genetic variability of the lichen symbionts appear to be mainly influenced by improved water relations at higher altitudes.

Keywords

crustose lichens, specificity, spatial patterns, polar desert, foehn winds, humidity

Introduction

The McMurdo Dry Valleys (MDV) in Southern Victoria Land of Continental Antarctica are characterized by an environment that is exceptional also for Antarctica: it is extremely arid and cold, which makes it hostile for most organisms. Thus, life is rare within the valleys of this polar desert, and only few life forms can cope with these extreme conditions (e.g. Adams et al. 2006; Pointing et al. 2009). The main limiting factor for life within the MDV is water availability with fog, clouds, dew and ephemeral melting water of snow patches having important effects on the climatic conditions (Adams et al. 2006; Green et al. 2007; Pannewitz et al. 2005; Stichbury et al. 2011). Among the most diverse macro-organisms present in the MDV are lichens. Lichens represent a classic example of symbiosis, consisting of a fungus (mycobiont) and one or more photosynthetic partners (photobiont). When completely desiccated, lichens are dormant and can survive unfavorable conditions for long periods (Green 2009; Kappen and Valladares 2007). As a consequence they are able to colonize rocks and boulders above melting streams or in the vicinity of snow patches, even in such extreme environments as the MDV (e.g. Green et al. 2011b; Hertel 2007; Ruprecht et al. 2012a; Ruprecht et al. 2010; Sancho et al. 2017; Schroeter et al. 2010). The most successful species are green-algal lichens, as they do not depend on the presence of liquid water for reactivation and can be active below zero degrees, in contrast to cyanobacterial lichens that appear to be completely absent in continental Antarctica (Green et al. 2011a; Kappen 2000; Lange et al. 1986; Schlenz et al. 1997; Seppelt et al. 2010).

Several studies on mycobiont-photobiont interactions in lichens have shown that green-algal as well as cyanobacterial lichenized fungi can show considerable photobiont variability and can have more than one photobiont and even combine green alga and cyanobacteria (Fernandez-Mendoza et al. 2011; Henskens et al. 2012; Nelsen and Gargas 2009; Otalora et al. 2010; Ruprecht et al. 2014; Wornik and Grube 2010). Low photobiont specificity and a high ability to accept different photobionts might be a survival strategy and extend the ecological range of lichens (Blaha et al. 2006; Dal Grande et al. 2017; Leavitt et al. 2015; Ruprecht et al. 2012a; Wirtz et al. 2003). Furthermore, photobiont selection appears to be influenced by abiotic factors like climate (Beck et al. 2002; Fernandez-Mendoza et al. 2011; Peksa and Skaloud 2011; Yahr et al. 2006). At the local scale (for instance along elevational gradients), this may translate into habitat-specific photobiont switches (Vargas Castillo and Beck 2012). Above all, temperature has often been identified as a key factor of photobiont selection of lichens in Antarctica (Green 2009; Kappen and Valladares 2007; Ruprecht et al. 2012a). In warmer regions, myco-/photobiont interactions show increased specificity leading to one-to-one interactions in contrast to more generalist interactions in colder environments (Singh et al. 2017). Thus, it appears that symbiotic interactions in lichens can react very sensitively to environmental change although this conclusion is based on a small database, and these responses have been investigated only in a few species (Allen and Lendemer 2016; Colesie et al. 2014b; Sancho et al. 2017). In general, there is agreement that climatic changes will influence the diversity, abundance and growth of lichens (Sancho et al. 2017) and that lichens therefore represent excellent bioindicators for processes associated with global warming (Alatalo et al. 2015; Allen and Lendemer 2016; Bassler et al. 2016; Sancho et al. 2019).

Over the last decades studies on elevational gradients have re-emerged because the species composition changes remarkably with elevation suggesting a species-specific adaptation to different environmental conditions (Grytnes et al. 2006). They provide steep ecological transitions (e.g. in temperature, humidity and UV radiation) over short distances (Keller et al. 2013; Körner 2007) and several studies suggest that the structure and diversity of communities, the abundance and distribution of species and ecosystem properties and processes can change along elevational gradients (Bassler et al. 2016; Dal Grande et al. 2017;

Grytnes et al. 2006; Junker and Larue-Kontic 2018; Körner 2003; Wolf 1993). For lichens, elevational gradients are reported to show large changes in species composition (Dal Grande et al. 2017; Leavitt et al. 2015), habitat-specific photobiont switches (Vargas Castillo and Beck 2012), and/or microclimatic partitioning of ecologically differentiated fungal and algal gene pools (Nadyeina et al. 2014).

This study focuses on saxicolous crustose lichens in continental Antarctica which are associated with green micro alga as photobionts. In general, these lichens are slow growing and restricted to microhabitats on rock surfaces (Hertel 1998), but nevertheless, due to their poikilohydric nature, they are well adapted to habitats with high insolation and with rapid fluctuations in temperature and water availability (Green et al. 2002; Lange 1997; Lange 2000; Schroeter et al. 2011). Most of the lichens analyzed here belong to the 'lecideoid' lichen group (Hertel 1984) and these species are assigned to the generic name *Lecidea sensu* Zahlbruckner (1925) but they do not necessarily belong to the genus in its strict sense. Due to their inconspicuous growth form, distinguishable only by a few small morphological traits such as spore size and ascus-type, the identification of these lichens is difficult even under best growing conditions (Ruprecht et al. 2019). Extreme climate conditions in cold deserts result in reduced development of the thallus such as chasmolithic growth, a lack of ascomata or sparsely developed ascospores, all features that hamper identification and even the detection of the specimens in the vast landscape (Hertel 2009; Ruprecht et al. 2010). Nevertheless, these pioneers on rocks and pebbles (Hertel 1984; Hertel 1987) belong to one of the most abundant species groups in continental Antarctica (Hertel 2007; Ruprecht et al. 2012b; Ruprecht et al. 2010) and are, therefore, an excellent study system to investigate changes in symbiotic associations along gradients. The present study covers lecideoid lichen species of the genera *Carbonea* Hertel, *Lecanora* Ach., *Lecidella* Körb., *Rhizoplaca* Zopf and the genus *Lecidea* Ach s.str. (Hertel 1984) plus, in addition, lichen samples of the genera *Austrolecia* Hertel, *Buellia* De Not. and *Huea* C.W. Dodge & G.E. Baker which were included, because they often have a similar appearance.

The aim of this study was to analyze the spatial pattern and factors that might affect the distribution of both symbiotic partners within the MDV. We addressed the following objectives: 1) to confirm and extend our knowledge of the abundances of the myco- and photobiont species that have been found by previous, less extensive studies, 2) to investigate the variability of mycobiont/photobiont interactions, in particular, analyze the level of selectivity by using network statistics and 3) to study for mycobiont, photobiont and lichens the relationships with abiotic and climatic factors such as elevation and water status.

Materials & Methods

Study area and sampling sites

This study was conducted as part of the New Zealand Terrestrial Antarctic Biocomplexity Survey (nzTABS, <http://nztabs.ictar.aq>), which was initiated during the International Polar Year 2007–2008, and drew a diverse range of international expertise to profile the biology, geochemistry, geology and climate of the MDV. The study is among the most comprehensive landscape-scale biodiversity surveys undertaken and includes nearly all trophic components found in the MDV ecosystem (Lee et al. 2019). Sampling of soils and biological communities was carried out over two successive Austral summers (2009/10, 2010/11). The geographic area within which lichen samples were collected was the southern part of the MDV (total area: 22700 km², ice-free area: 4500 km²; Levy 2013; Fig. 1a, b). The landscape is a mosaic of glacially formed valleys with intervening high ground, ice-covered lakes, ephemeral streams, arid rocky soils, ice-cemented soils, and surrounding glaciers along the steep scree and

boulder slopes (Fig. 1b-d; Doran et al. 2002; Stichbury et al. 2011; Yung et al. 2014). There are four main valleys (Miers Valley, Garwood Valley, Hidden Valley and Marshall Valley) and some other extensive ice-free areas (Shangri-La). The topography ranges from sea level to more than 2000 m a.s.l. with granite being the dominant rock type on the ridges and hills, whilst the valley floors are covered with glacial drift. The valleys have the typical glaciated form with a U-cross-section with steep sides, often with scree slopes, which reach up to around 600 m in height. To the west the valleys are separated from the polar plateau by the Royal Society Range that has peaks over 4000 m a.s.l. in height. To the east the valleys open out onto the Ross Ice Shelf, which represents a climatically maritime influenced location within the MDV, despite the absence of ice-free sea at any time of the year (Yung et al. 2014).

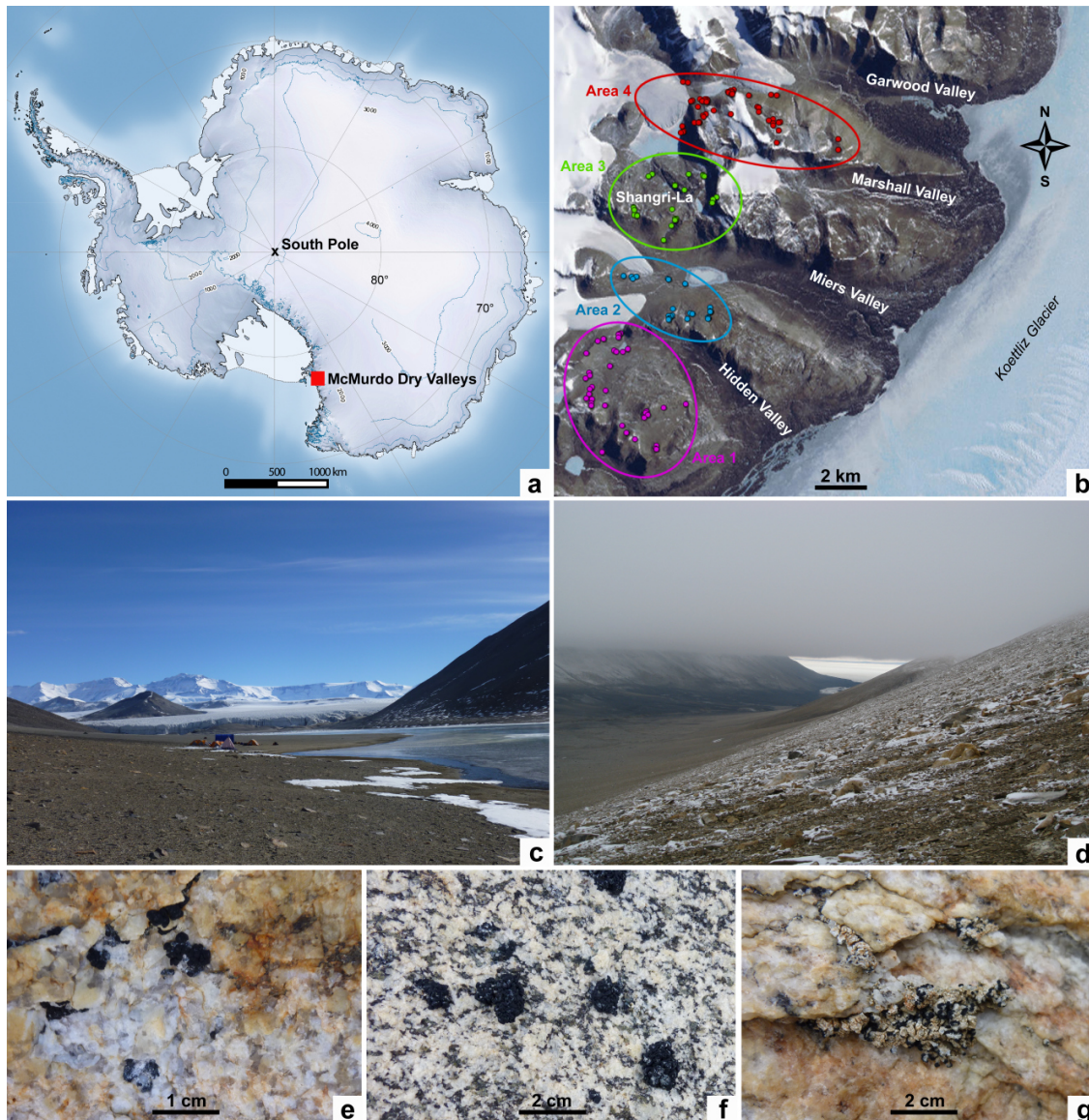


Figure 1: (a) Antarctic continent; investigated area marked with red rectangle (Natural Earth, Qgis), (b) MDV sampling sites defined in four areas (<http://www.gpsvisualizer.com>), (c) Garwood Valley and field camp, (d) Garwood Valley with incoming cloud bank, (e) *Lecidea cancriformis*, (f) *Rhizoplaca macleanii*, (g) *Austrolecia* sp. 1.

Climate of the MDV

The climate of the MDV is, for several reasons, classified as that of a polar desert. First, the mountains at the west are sufficiently high to block seaward flowing ice from the East Antarctic ice sheet from reaching the Ross Sea. In addition, the Transantarctic Mountains provide a precipitation shadow, causing an extremely low humidity and lack of snow or ice cover in the MDV (Monaghan et al. 2005). Annual precipitation is $< 50 \text{ mm a}^{-1}$ water equivalent, with precipitation decreasing away from the coast (Fountain et al. 2010). The major source of liquid water is the seasonal melting of perennial snowbanks and glaciers (Head and Marchant 2014; Stichbury et al. 2011) but, in most cases, this water is not available for lichens that inhabit rock surfaces above the surrounding ground level. MDV climate is best known from the northern valleys, particularly Taylor Valley, because of the McMurdo Long Term Ecological Research programme (McMurdo LTER) that has been active since 1993 (<http://www.mcmllter.org>). The valley floors of the MDV show mean annual temperatures that range from $-30 \text{ }^{\circ}\text{C}$ to $-15 \text{ }^{\circ}\text{C}$ and typically have fewer than 50 days/a where average temperatures exceed $0 \text{ }^{\circ}\text{C}$ (Colesie et al. 2014b; Doran et al. 2002; Ochyra et al. 2008). There is agreement that the air temperature lapse rate is close to $1 \text{ }^{\circ}\text{C}$ decline per 100 m elevation rise, as well as an increase with distance from the coast to the inland of $0.09 \text{ }^{\circ}\text{C}$ per 1 km (McKay 2015). The aspect of the valley slopes has an important impact and north facing slopes are warmer and dryer, south facing slopes are cooler and wetter (Yung et al. 2014). The wind regime is strongly topographically channeled and directed mainly up- or down-valley. During summer, easterly valley winds dominate, due to differential surface heating between the low albedo valley floors and the high albedo ice to the east (Mckendry and Lewthwaite 1990). In winter, wind direction is typically more variable. Cold air pools associated with light winds and very low minimum temperatures ($-50 \text{ }^{\circ}\text{C}$) often occupy topographic low points of the valleys during winter (Doran et al. 2002).

Almost all climate information comes from studies on the valley floors. There are, however, conditions that tend to produce a major difference in water regime between valley floors and intervening mountain ranges. First, there is a tendency at higher elevations for greater snowfall and higher humidity, as shown by the presence of clouds at higher elevations (Fig. 1d). Second, there is the regular occurrence within the valleys of what have traditionally been regarded as katabatic winds (Ayling and McGowan 2006; Mckendry and Lewthwaite 1990) but which are now suggested to be foehn winds albeit generated in a slightly different manner to the classic northern hemisphere foehns (Speirs et al. 2010). In the Taylor Valley, for example, these winds are easily recognizable by their sudden arrival, high intensity (around 15 m s^{-1}), rapidly rising temperature (by around $25 \text{ }^{\circ}\text{C}$ to reach about $0 \text{ }^{\circ}\text{C}$), and rapidly falling relative air humidity to around 20 % (Speirs et al. 2010). These foehn winds also occur in the southern valleys with an example from Miers Valley (Online Resource 2a) showing almost identical characteristics to those in the Taylor Valley. Foehn winds are extremely drying with air vapor pressure deficit rising about 50 times from 0.01 kPa ($-30 \text{ }^{\circ}\text{C}$, 80 % RH) to 0.49 kPa ($0 \text{ }^{\circ}\text{C}$, 20 % RH). They are also topographically constrained within the valleys and can apparently reach altitudes up to almost 500 m (Speirs et al. 2010). The net result of the higher elevation cold, moister air, and the extremely drying foehn winds within the valleys is that the wetness availability gradient is strongly non-linear and, for the purposes of our analyses, we defined an elevational threshold of about 600 m a.s.l. which marks the upper limit of the steeper valley sides.

Sample sources

The present study includes 232 lichen samples (lecioid lichen species of the genera *Carbonea*, *Lecanora*, *Lecidella*, *Rhizoplaca*, *Lecidea* and additionally *Austrolecia*, *Buellia* and *Huea*, which have a similar appearance under the extreme climate conditions) collected in four different areas (Hidden Valley, Miers Valley, Shangri La and Garwood Valley) at 153 different localities with a range of aspects (N-facing slopes, flat areas and plateaus) and elevations (171-959 m a.s.l.; Fig. 1a - g, Table 1).

154 specimens were collected between the years 2009 to 2011 by Ulrike Ruprecht and Roman Türk and are deposited at the herbarium of the University of Salzburg (SZU, Online Resource 1a). An additional 78 lichen samples from the same area were obtained from the study of Perez-Ortega et al. (2012) excluding specimens of the genera *Acarospora*, *Caloplaca*, *Polysporina*, *Sarcogyne* and *Umbilicaria* (see Online Resource 1b).

Please note that for most of the data evaluations, mycobionts and photobionts were treated separately. In some analyses (noted in text), only mycobiont species with $n \geq 10$ (*min10MycoSp*) were used, whilst others included only photobiont haplotypes with $n \geq 10$ (*min10PhoHap*).

Table 1: Site descriptions and specifics of the four regions defined within the MDV, including the range of the coordinates of the sampling sites and areas, minimum, maximum and mean values of the elevation of the sampling sites, and the number of lichen samples per area.

	Sampling area	Range of coordinates of sampling sites	Min., max. & mean of elevation of sampling sites (m a.s.l.)	Number of sampling sites	Number of lichen samples
Area 1	Hidden Valley	S 78.12° - 78.17° E 163.62° - 163.81°	361, 645, 911	39	53
Area 2	Miers Lake	S 78.10° - 78.11° E 163.69° - 163.86	171, 477, 597	28	42
Area 3	Shangri-La	S 78.05° - 78.08° E 163.71° - 163.87°	411, 656, 890	30	51
Area 4	Garwood Valley	S 78.02° - 78.05° E 163.80° - 164.10°	319, 593, 959	56	86

DNA-amplification, purification and sequencing

Total DNA was extracted from the thallus and/or apothecia by using the DNeasy Plant Mini Kit (Qiagen) following the manufacturer's instructions. For all samples, we sequenced and amplified the internal transcribed spacer (ITS) region of the mycobionts' and photobionts' nuclear ribosomal DNA (nrITS). We also amplified additional markers: for the mycobionts the mitochondrial small subunit (mtSSU) and the low-copy protein coding marker RPB1 and, for the photobionts, the chloroplast-encoded intergenic spacer (psbJ-L) and part of the cytochrome oxidase subunit 2 gene (COX2). This was done using newly developed specific primers and PCR-protocols in our project-framework (Ruprecht et al. 2019).

For amplifying nrITS of the mycobiont we used the primers ITS1 (White et al. 1990), ITS1F (Gardes and Bruns 1993), ITS1L (Ruprecht et al. 2019), ITS4 (White et al. 1990), ITS4L (Ruprecht et al. 2019) and for the photobiont 18S-ITS uni-for (Ruprecht et al. 2012a), ITS1T (Kroken and Taylor 2000), ITS1aT (Ruprecht et al. 2014), ITS4T (Kroken and Taylor 2000) and ITS4bT (Ruprecht et al. 2014). For the marker mtSSU the primers CU6 (<https://nature.berkeley.edu/brunslab/tour/primers.html>), mrSSU1 (Zoller et al. 1999), mtSSU for2 (Ruprecht et al. 2010) and mtSSU rev2 (Ruprecht et al. 2010) and for RPB1, fRPB1-C rev (Matheny et al. 2002), gRPB1-A for (Matheny et al. 2002) and RPB1_for_Lec (Ruprecht et al. 2019) were chosen. For the marker COX2, COXIIIF2 and COXIIr (Lindgren et al. 2014) and COX sense (Ruprecht et al. 2019) and for psbJ-L, psbF (Werth and Sork 2010), psbL-sense and psbJ-antisense (Ruprecht et al. 2014) were used. All reactions were performed as described in (Ruprecht et al. 2019). Unpurified PCR-products were sent to Eurofins Genomics/Germany for sequencing.

Phylogenetic analysis

The sequences of the different marker regions listed above were assembled and edited using Geneious version 6.1.8 (Kearse et al. 2012) and aligned with MAFFT v7.017 (Kato et al. 2002) for both symbionts. For the photobiont, the classification and labeling of the different operational taxonomical units (OTUs) followed the concept of Leavitt et al. (2015), reevaluated by Ruprecht et al. (2019).

Phylogenetic relationships of the samples of the present study were calculated from the sequences of the marker nrITS. The other markers could not provide further intraspecific variation and were not available for every specimen; therefore they were excluded in all following analyses using sequence data.

A maximum likelihood analysis was calculated with the IQ-TREE web server (Trifinopoulos et al. 2016), using the model selection algorithm ModelFinder (Kalyaanamoorthy et al. 2017). The BIC (Bayesian information criterion) selected for the best-fit model for the mycobiont alignment TN+I+G4 and for the photobiont K2P+I. Branch supports were obtained with the implemented ultrafast bootstrap (UFBoot) (Minh et al. 2013) (number of bootstrap alignments: 1000, maximum iteration: 1000, minimum correlation coefficient: 0.99). Additionally, a SH-aLRT branch test (Guindon et al. 2010) was performed. Each branch of the resulting tree was assigned with SH-aLRT as well as UFBoot supports; the branches with SH-aLRT < 80 % and/ or UFboot < 95 % were collapsed by adding the command *-minsupnew 80/95* to the script.

Haplotype analysis

We determined the haplotypes (h) of the different mycobiont species and photobiont OTUs by using the function `haplotype()` of the R package `pegas` (Paradis 2010) (note: the function only takes into account transversions and transitions but ignores insertions and deletions). For *min10MycoSp* species and photobiont OTUs with $h \geq 2$ and at least two haplotypes with $n \geq 3$ (*Lecidea cancriformis*, *Lecidella greenii*, *Rhizoplaca macleanii* and photobiont OTU *Tr_A02*), haplotype networks were computed, using the function `haploNet()` of the R package `pegas` (Paradis 2010). The frequencies were clustered in 10% ranges, for example the circles of all haplotypes making up between 20-30 % have the same size.

Analysis of spatial distribution

To analyze how the distribution of the lichen specimens correlated with abiotic factors, the sampling sites of the different lichen species or haplotypes in the investigated areas were compared with respect to their environmental specifics. For this we tested the only relevant variable which was elevation. All other variables such as latitude, longitude, and the BIOCLIM variables generated by Wagner et al. (2017) providing a spatial resolution of 1 km, were not suitable for the relatively small area (data not shown). To assure a minimum group size of 10 sample points, the tests only included the *min10MycoSp* species and *min10PhoHap* haplotypes.

In addition, the elevation of the sample sites of the two most dominant photobiont OTUs (*Tr_A02* and *Tr_S15*) were compared by conducting a nonparametric t-test, using the R function `npar.t.test()` of the package `nparcomp` (Konietschke et al. 2015). We used nonparametric multiple comparisons for relative effects (`mctp-test`; function `mctp()`) of the R package `nparcomp` (Konietschke et al. 2015), which conducts pairwise comparisons of all possible combinations.

Analysis of mycobiont – photobiont associations

The associations between mycobiont and photobiont haplotypes were analyzed by computing bipartite networks, using the R function `plotweb()` of the package `bipartite` (Dormann et al. 2008). For the bipartite network including mycobiont species and photobiont haplotypes the indices H_2' and d' (Blüthgen et al. 2006) were calculated. Both indices are derived from Shannon entropy. H_2' characterizes the degree of complementary specialization or partitioning among the two parties of the entire bipartite network, while d' describes the degree of complementary specialization at species or haplotype level. They both range from 0 for the most generalized to 1 for the most specialized case and were computed using the R functions `H2fun()` and `dfun()` of the package `bipartite` (Dormann et al. 2008).

Phylogenetic species diversity of the interaction partners was quantified by calculating a number of further metrics listed in Table 2, including the indices NRI (Net relatedness index), PSV (Phylogenetic species variability) and PSR (Phylogenetic species richness).

Table 2: Diversity metrics compared in this study, citations and descriptions of each, and the used R functions.

Metric	Definition	References	Description and interpretation of values	R function, R package
d'	Specialization index	(Blüthgen et al. 2006)	Degree of interaction specialization at species level (values range from 0 = most generalized case to 1 = most specialized case)	<code>dfun()</code> , <code>bipartite</code> (Dormann et al. 2008)
NRI	Net relatedness index	(Webb 2000; Webb et al. 2002)	Comparison of phylogenetic distances among all members of a community (pos. values = phylogenetic clustering; neg. values = phylogenetic evenness)	<code>ses.mpd()</code> , <code>picante</code> (Kembel et al. 2010)
PSV	Phylogenetic species variability	(Helmus et al. 2007)	Degree to which species in a community are phylogenetically related (values range from 0 = increased relatedness to 1 = decreased relatedness)	<code>psd()</code> , <code>picante</code> (Kembel et al. 2010)
PSR	Phylogenetic species richness	(Helmus et al. 2007)	PSV multiplied by species richness SR (number of species in a sample); SR after discounting species relatedness (values range from 0 = increased relatedness to SR = decreased relatedness)	<code>psd()</code> , <code>picante</code> (Kembel et al. 2010)

Analysis of DNA polymorphism

For each identified mycobiont and photobiont species with more than one sample, we calculated the haplotype as well as the nucleotide diversity using p-distances with `DnaSP v5` (Librado and Rozas 2009). Gaps and missing data were excluded. We focused on h/N (number of haplotypes, h , divided by number of samples, N , per species), Hd (haplotype diversity, the probability that two randomly chosen haplotypes are different; Nei 1987) and π (nucleotide diversity, average number of nucleotide differences per site between two randomly chosen DNA sequences; Nei and Li 1979).

In order to analyze the dependence of haplotype and nucleotide diversity values on elevation, we used the defined threshold of 600 m a.s.l. The h/N , Hd , d' , and PSV of those *min10MycoSp* species with mean values above this threshold were grouped together and then compared to those species with mean values below 600 m a.s.l., using the R function `nonpartest()` of the package `npmv` (Ellis et al. 2017), which performs nonparametric comparisons of multivariate samples. (Note: π , NRI, and PSR were excluded because of high correlations ($r \geq 0.85$) with h/N (π), Hd (π), d' (π) and PSV (NRI and PSR)).

Results

Phylogenetic analysis

The molecular phylogenies for the mycobiont (Online Resource 2b) and the photobiont (Online Resource 2c) are based on the marker nrITS, because the additional markers (mycobiont: mtSSU, RPB1; photobiont: psbJ-L, COX2) showed little sequence variation in this area. Both analyses include only accessions from the study sites (Online Resource 1a, b) to present the various species- and diversity levels.

Mycobiont: The final data matrix for the phylogeny comprised 232 single sequences of the marker nrITS with a length of 610 bp and included sequences of the families *Lecanoraceae*, *Teloschistaceae*, *Catillariaceae*, *Caliciaceae* and *Lecideaceae*. The phylogenetic tree was midpoint rooted and shows a total of 25 strongly supported clades on species level, assigned to eight genera. The backbone is not supported and therefore the topology will not be discussed. The genera *Huea*, *Austrolecia*, *Buellia* and *Lecidea* are clearly assigned to their family level and are strongly supported. The genera *Carbonea*, *Lecidella* and *Rhizoplaca* assigned to the family *Lecanoraceae* each form highly supported clades, but do not form an independent clade. The clade of the genus *Lecanora* is divided in five species (*L. cf. mons-nivis*, *L. fuscobrunnea*, *L. sp. 1*, *L. sp. 2* and *L. sp. 3*), *Carbonea* in three species (*C. vorticosa*, *C. sp. URm1* and *C. sp. 2*), *Austrolecia* in three species (*A. sp. 1*, *A. sp. 2* and *A. sp. 4*), *Buellia* in four species (*B. frigida*, *B. sp. 1*, *B. sp. 2* and *B. sp. 3*) and *Lecidea* in seven species (*L. andersonii*, *L. cancriformis*, *L. lapicida*, *L. polypycnidophora*, *L. sp. 5*, *L. sp. 6* and *L. UCR1*). The samples allocated to the genera *Lecidella*, *Rhizoplaca* and *Huea* were, in each genus, monospecific (*Lecidella greenii*, *Rhizoplaca macleanii* and *Huea sp. 1*). The taxonomical assignment of the obtained sequences was made using the following literature: Perez-Ortega et al. (2012); Ruprecht et al. (2012b); Ruprecht et al. (2010).

Photobiont: The final data matrix for the phylogeny comprised 222 single sequences of the marker nrITS with a length of 578 bp. The phylogenetic tree was midpoint rooted and shows four strongly-supported clades, all of them belonging to the genus *Trebouxia*. They were assigned to OTU level (Puillandre et al. 2012) using the system of Leavitt et al. (2015), reevaluated by (Ruprecht et al. 2019) and assigned to *Tr_A02*, *Tr_S02*, *Tr_S15*, *Tr_S18*. The OTU *Tr_A02* was by far the most common, being present in 202 (91 %) of our 222 samples. Photobiont sequences taken from Perez-Ortega et al. (2012) which were labelled only with numbers were included in our system of assigning the haplotypes to the appropriate OTUs (Ruprecht et al. 2019) and therefore renamed.

Haplotype analysis

The following analyses were based on the nrITS sequences of myco- and photobionts. The number of haplotypes differed significantly between myco- and photobionts. We identified 48 different mycobiont but only 17 different photobiont haplotypes. The most frequent mycobiont haplotype was *Lecidella greenii_h01* with 28 samples, the most frequent photobiont haplotype was *Tr_A02_h01* with 87 samples. Additionally, some of the mycobiont species appear to be more diverse (number of different haplotypes) than others. Fehler! Verweisquelle konnte nicht gefunden werden. & 3 show bar charts that give the number of samples per mycobiont species/photobiont OTU per area and per haplotype.

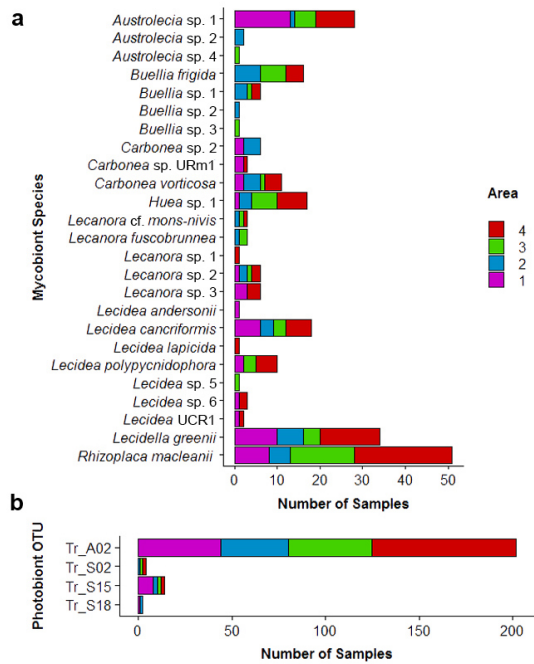


Figure 2: Number of samples per species/OTU and area (cf. Fig. 1b). (a) mycobiont species (total sample size: $n = 232$), (b) photobiont OTUs (total sample size: $n = 222$).

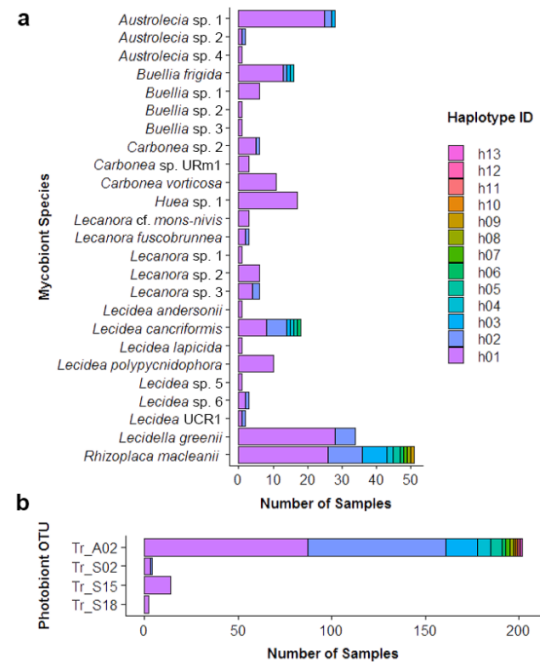


Figure 3: Number of samples per species/OTU and haplotype. (a) mycobiont species (total sample size: $n = 232$), (b) photobiont OTUs (total sample size: $n = 222$).

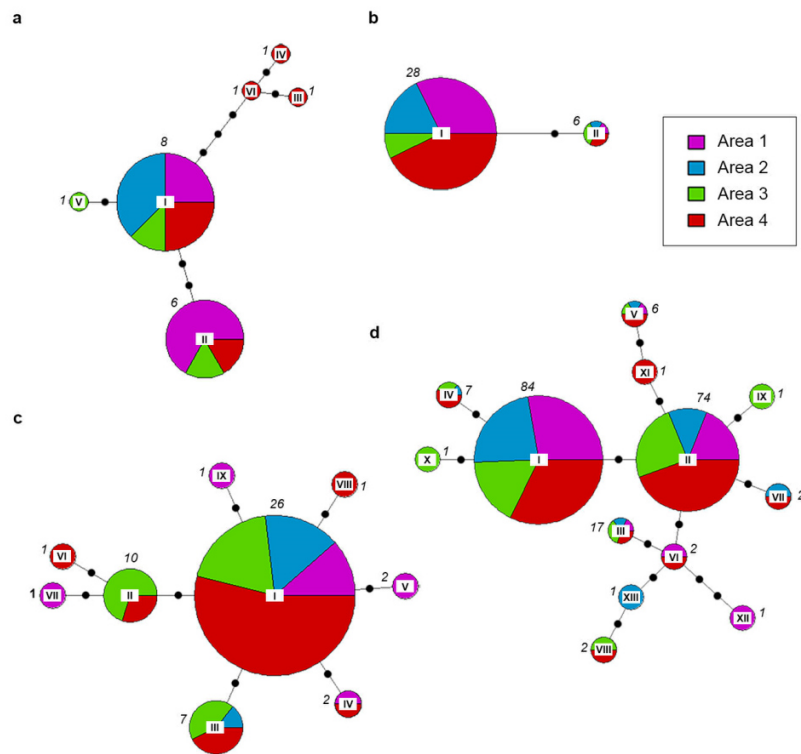


Figure 4: Haplotype networks of those mycobiont species/ photobiont OTUs with $h \geq 2$ and at least two haplotypes with $n \geq 3$: (a) *Lecideia cancriformis*, (b) *Lecidella greenii*, (c) *Rhizoplaca macleanii* and the photobiont OTU *Tr_A02*, (d) showing the spatial distribution within Area 1 to Area 4 (cf. Fig. 1b & 2). Roman numerals at the center of the pie charts refer to the haplotype IDs, italic numbers next to the pie charts to the total number of samples per haplotype. The circle sizes reflect relative frequency within the species/OTU; in doing so, frequencies were clustered in ten, so that for example the circles of all haplotypes making up between 20-30 % have the same size.

Three different mycobiont species (*Lecidea cancriformis*, *Lecidella greenii* and *Rhizoplaca macleanii*) and the most common photobiont OTU (*Tr_A02*) met the required criteria defined above for the construction of haplotype networks ($h \geq 2$ and at least two haplotypes with $n \geq 3$). In Fig. 4 the respective haplotype networks show the spatial location within the four areas. As shown in Fig. 2 for mycobiont species/photobiont OTU, the distribution again turned out to be rather uniform, with most of the haplotypes found in all of the four areas.

Analysis of spatial distribution

For 12 of the 28 pairwise comparisons for the mycobionts species (*min10MycoSp*) and photobiont haplotypes (*min10PhoHap*) the mctp-tests (pairwise comparisons of all possible combinations) for elevation showed significant differences, which are also visually recognizable when comparing the maps of the sample locations where the sample locations for each species are shown separately with varying colors indicating their elevations (*min10MycoSp* and *min10PhoHap* summarized in the respective OTUs; Online Resource 2d). The pairs with significant differences as well as the associated p-values are given in Online Resource 1c.

For the photobionts the mctp-test showed significant differences for only for two of the six pairwise comparisons (*Tr_A02_h01* and *Tr_A02_h03*, *Tr_A02_h01* and *Tr_S15_h01* and, in each case, *Tr_A02_h01* has lower elevation values; Online Resource 1c). The nonparametric t-test for comparing the elevation of the sample locations on OTU level resulted in a significant difference between the two groups for *Tr_A02* and *Tr_S15* ($p = 0.005$) with sampling sites of the OTU *Tr_S15* being higher.

Fig. 5 shows the elevational distribution of the different mycobiont species and photobiont OTUs at their sample sites for *min10MycoSp* species and *min10PhoHap* haplotypes. The mycobiont species *Lecidea cancriformis* and *Rhizoplaca macleanii* had a significant tendency to higher elevations, whilst *Carbonea vorticosa*, *Lecidea polypycnidophora* and *Lecidella greenii* had a significant tendency to lower elevations. One haplotype of the photobiont, OTU *Tr_S15*, was restricted to higher elevations. The remaining mycobiont species and photobiont haplotypes have an intermediate distribution.

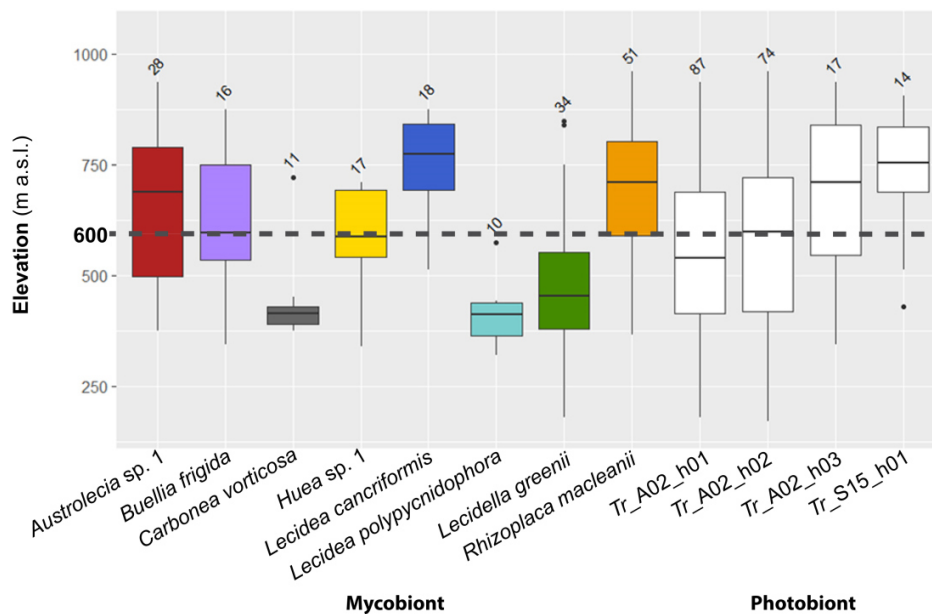


Figure 5: Boxplots showing the elevation of the sample sites of the *min10MycoSp* species and *min10PhoHap* haplotypes. Numbers in italics refer to sample sizes. The elevational threshold of 600 m a.s.l. is highlighted with a dashed line.

Analysis of mycobiont-photobiont associations

The bipartite network was calculated for all associations between the mycobiont species (*min10MycoSp*; lower level) and the respective photobiont haplotypes (higher level; Fig. 6). The H_2' (overall level of complementary specialization of all interacting species) of this network shows a low value of 0.226 which indicates a network with mostly generalized interactions (as opposed to specialized). This was mainly caused by the dominant occurrence of the three most abundant haplotypes (h01-h03) of the most common photobiont OTU *Tr_A02* (91 % of 222 accessions). All mycobiont species were additionally associated to a variety of other and less abundant haplotypes (h04-h13) of this OTU *Tr_A02*. Furthermore, some of the mycobionts were associated with accessions of the distantly related OTUs *Tr_S02*, *Tr_S15* and *Tr_S18*. The network matrix shows the number of associations between the mycobiont species and photobiont haplotypes (Online Resource 1d). The individual d' values (complementary specialization at species or haplotype level) are ranging from 0 to 1 and are presented in Table 3.

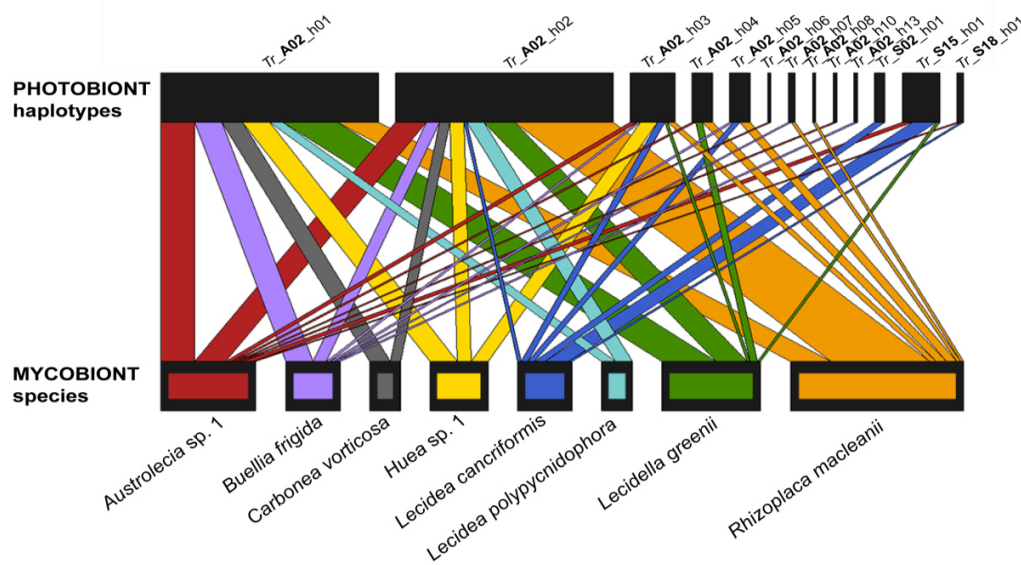


Figure 6: Bipartite network including the *min10MycoSp* species as well as the haplotypes of their associated photobionts with an $H_2' = 0.226$. Rectangles represent species, and the width is proportional to the number of samples. Associated species are linked by lines, whose width is proportional to the number of associations.

Species richness of mycobiont vs photobiont

The number of different symbiotic partners at haplotype level (SR) as a function of the number of mycobiont haplotypes (h in Table 3) is illustrated in Fig. 7 for the *min10MycoSp* species. The two variables show a correlation of $r = 0.701$; thus, highly variable mycobionts tend to be associated with a higher number of photobiont haplotypes.

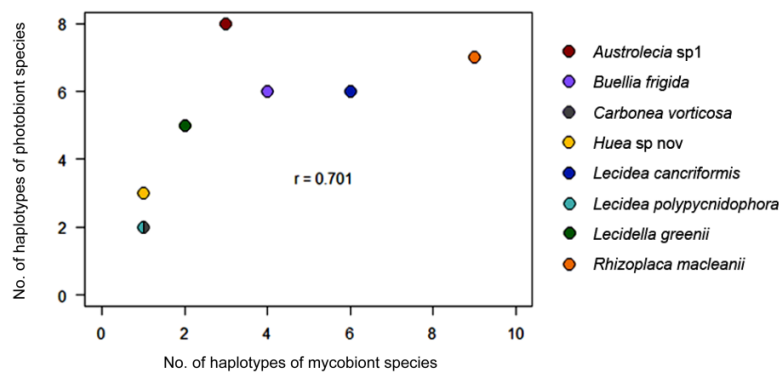


Figure 7: Scatterplot of the species richness (SR) of photobiont haplotypes dependent on the number of haplotypes (h) of each *min10MycoSp* species (*Carbonea vorticosa* and *Lecidea polypycnidophora* share the same coordinates) including the Pearson correlation coefficient (r).

Analysis of DNA polymorphism and nonparametric comparisons of multivariate samples

Analyses of DNA polymorphism and nonparametric comparisons of multivariate samples were achieved for *min10MycoSp* and *min10PhoHap* including the parameters h/N (number of haplotypes, h , divided by number of samples, N , per species), Hd (haplotype diversity), and π (nucleotide diversity), d' (specialization index), NRI (net relatedness index), PSV (phylogenetic species variability), SR (species richness) and PSR (phylogenetic species richness; Table 3).

Table 3: Diversity (left) and specificity indices (right) for the different mycobiont species and photobiont OTUs: N, number of sequences; h, number of haplotypes; h/N , quotient of h and N ; Hd, haplotype diversity; π , nucleotide diversity; d' , specialization index; NRI, net relatedness index; PSV, phylogenetic species variability; SR, species richness; PSR, phylogenetic species richness.

Mycobiont Species	N	h	h/N	Hd	π	d'	NRI	PSV	SR	PSR
<i>Austrolecia</i> sp. 1	28	3	0.107	0.204	0.00069	0.053	-0.127	0.454	8	3.636
<i>Austrolecia</i> sp. 2	2	2	1.000	1.000	0.00791	0.140	-	-	1	-
<i>Austrolecia</i> sp. 4	1	1	1.000	-	-	0.000	-	-	1	-
<i>Buellia frigida</i>	16	4	0.250	0.350	0.00098	0.105	2.210	0.029	6	0.173
<i>Buellia</i> sp. 1	6	1	0.167	0.000	0.00000	0.014	-0.784	0.670	3	2.011
<i>Buellia</i> sp. 2	1	1	1.000	-	-	0.000	-	-	1	-
<i>Buellia</i> sp. 3	1	1	1.000	-	-	0.036	-	-	1	-
<i>Carbonea</i> sp. 2	6	2	0.333	0.333	0.00073	0.163	-0.354	0.511	4	2.046
<i>Carbonea</i> URm1	3	1	0.333	0.000	0.00000	0.000	0.895	0.011	2	0.023
<i>Carbonea vorticosa</i>	11	1	0.091	0.000	0.00000	0.064	0.895	0.011	2	0.023
<i>Huea</i> sp. 1	17	1	0.059	0.000	0.00000	0.120	1.266	0.023	3	0.069
<i>Lecanora</i> cf. <i>mons-nivis</i>	3	1	0.333	0.000	0.00000	0.305	0.819	0.046	2	0.092
<i>Lecanora fuscobrunnea</i>	3	2	0.667	0.667	0.00129	0.391	-1.239	1.000	2	2.000
<i>Lecanora</i> sp. 1	1	1	1.000	-	-	0.366	-	-	1	-
<i>Lecanora</i> sp. 2	6	1	0.167	0.000	0.00000	0.048	1.285	0.015	3	0.045
<i>Lecanora</i> sp. 3	6	2	0.333	0.533	0.00104	0.254	1.590	0.025	4	0.099
<i>Lecidea andersonii</i>	1	1	1.000	-	-	0.366	-	-	1	-
<i>Lecidea cancriformis</i>	18	6	0.333	0.719	0.00388	0.595	-1.273	0.665	6	3.990
<i>Lecidea lapicida</i>	1	1	1.000	-	-	0.000	-	-	1	-
<i>Lecidea polypycnidophora</i>	10	1	0.100	0.000	0.00000	0.059	0.895	0.011	2	0.023
<i>Lecidea</i> sp. 5	1	1	1.000	-	-	1.000	-	-	1	-
<i>Lecidea</i> sp. 6	3	2	0.667	0.667	0.01028	0.147	-	-	1	-
<i>Lecidea</i> UCR1	2	2	1.000	1.000	0.00193	0.491	0.866	0.023	2	0.045
<i>Lecidella greenii</i>	34	2	0.059	0.299	0.00062	0.049	0.027	0.415	5	2.074
<i>Rhizoplaca macleanii</i>	51	9	0.176	0.692	0.00185	0.155	2.528	0.033	7	0.234
Photobiont OTU	N	h	h/N	Hd	π	d'	NRI	PSV	SR	PSR
Tr_A02	202	11	0.054	0.659	0.00174	0.367	1.235	0.485	42	20.375
Tr_S02	4	2	0.500	0.500	0.00189	0.721	-0.336	0.441	2	0.882
Tr_S15	14	1	0.071	0.000	0.00000	0.493	-0.965	0.529	7	3.705
Tr_S18	2	1	0.500	0.000	0.00000	0.344	-1.188	1.000	2	2.000

The lowest values of Hd and π were developed by four mycobiont species of *min10MycoSp* (*C. vorticosa*, *L. polypycnidophora*, *L. greenii* and *Huea* sp. 1) which occur at the lowest and intermediate elevations. In contrast, species that were found only at the higher elevations (*R. macleanii*, *L. cancriformis*) showed the opposite behavior and had the highest values for Hd and π . These results were supported by nonparametric tests of the different haplotype and nucleotide diversity values which showed a significant difference between *min10MycoSp* samples below the threshold of 600 m a.s.l. (*Carbonea vorticosa*, *Huea* sp. 1, *Lecidea polypycnidophora* and *Lecidella greenii*) and those above (*Austrolecia* sp. 1, *Buellia frigida*, *Lecidea cancriformis* and *Rhizoplaca macleanii*) in terms of h/N , Hd , d' and PSV (ANOVA type test p-value: 0.009). The scatterplots of the indices as a function of elevation as well as the Pearson correlation coefficients are presented in Online Resource 2e.

Discussion

The McMurdo Dry Valleys at 78° S in Victoria Land, Antarctica are known as the largest continuous ice-free landscape on the continent and are mostly colonized by lithic and soil related microbial communities (Bottos et al. 2014; Colesie et al. 2014a; De Los Rios et al. 2014; Lee et al. 2019; Pérez-Ortega et al. 2012; Yung et al. 2014). The only comprehensive evaluation to date for saxicolous lichens was that of Pérez-Ortega et al. (2012) which showed a high number of mycobiont species (26) and, in contrast, a low number of photobiont species (four). Here we use a new and much larger dataset focused specifically on the lecideoid lichen group that includes some other species with similar morphologies in the same and adjacent areas and the respective data of Pérez-Ortega et al. (2012; Fig. 1b). A total of 25 mycobiont species were identified with a composition that was mostly similar to the previous study but including a few additional specimens (*Lecanora fuscobrunnea*, unknown *Lecidea* and *Buellia* species). They were all associated with the same set of common photobionts (four species), dominated (91%) by the more recently reclassified OTU *Trebouxia* A02 (Leavitt et al. 2015), equivalent to the species *T.* sp. URa2 (Ruprecht et al. 2012a). The evaluation of Pérez-Ortega et al. (2012) included not only the same set of lecideoid and morphologically similar lichen genera (*Lecidea*; *Carbonea*, *Lecanora*, *Lecidella*, *Rhizoplaca*; *Austrolecia*; *Buellia*; *Huea*) as investigated in this study, but also five additional lichen genera (*Acarospora*; *Caloplaca*; *Polysporina*; *Sarcogyne*; *Umbilicaria*).

In our study, we found that the different mycobiont species and photobiont OTUs within the MDV appear to be relatively evenly distributed across all four primary sample sites (Fig. 2), which is in basic agreement to the previous study. However, our results show distinct patterns for distribution, genetic diversity and specificity. These results contrast with Pérez-Ortega et al. (2012) where the distribution of mycobionts and photobionts was independent of elevation and other abiotic factors. A clear trend has now emerged showing that the distribution of species/OTUs is significantly related to elevation, using 600 m a.s.l. as a defined threshold dividing higher and lower sites. The mycobiont species *Carbonea vorticosa*, *Lecidea polypycnidophora* and *Lecidella greenii* were found almost exclusively below and *Lecidea cancriformis* and *Rhizoplaca macleanii* above the threshold (Fig. 5), which was supported by mctp-tests for pairwise comparisons (Online Resource 1c).

In contrast, the dominant photobiont OTU *Tr_A02* is distributed everywhere whilst the remaining and distantly related OTUs (*Tr_S02*, *Tr_S15* and *Tr_S18*) are mostly restricted to the higher elevation habitats (cold and humid; Fig. 5, Online Resource 2c). This result is in agreement with Dal Grande et al. (2018), who reported clear elevational preferences for some *Trebouxia* taxa at the OTU level at a mountain range in central Spain covering an elevational gradient of 1400 m. He suggested that altitude plays a prominent role in shaping the community structure of these algae. The distribution of *Tr_S15* (equivalent to the species *T.* URa1), only occurring in the cold and humid areas was surprising because the first specimen of this species described was found

at the climatically most extreme and dry habitats in the Darwin area 80° S (Ruprecht et al. 2012a). However, lichen photobionts seem to have clear ecological preferences and niche differentiations. This was also shown for various *Asterochloris* lineages (Trebouxiophyceae) which have diverging preferences in terms of rain and sun exposure (Peksa and Skaloud 2011) whilst Nadyeina et al. (2014) reported elevational partitioning in the distribution of different gene pools of the photobiont of the lungwort lichen *Symbiochloris reticulata* (Trebouxiophyceae; Skaloud et al., 2016). In general, several studies suggest a strong genetic association of lichen-associated algae with climatic factors and substrate (e.g. Fernandez-Mendoza et al. 2011; Vargas Castillo and Beck 2012; Yahr et al. 2006), and this has been interpreted as evidence for ecological specialization (Muggia et al. 2014; Ruprecht et al. 2012a).

The mycobiont species *min10MycoSp* not only show clear spatial differentiation with respect to elevation for species and OTUs but also for variables expressing the genetic diversity and specialization towards both symbiotic partners. A higher elevation correlates with a higher number of haplotypes (Hd) and an increased nucleotide diversity (π) which leads to a greater intraspecific diversity within the mycobionts (Table 3). These differences are also partially reflected by a higher d' , PSV, PSR and a low NRI which show a low relatedness to the co-occurring photobiont, associated with the rarely occurring and highly differentiated other OTUs *Tr_S02*, *Tr_S15* and *Tr_S18*. Consequently, mycobionts with a high genetic diversity have a higher number of interacting partners. These findings are partially supported by the study of Singh et al. (2017), who reported climate as a selective pressure in terms of increased specificity of myco-/photobiont interactions.

Our study has also shown that highly variable mycobionts are associated with a larger number of photobiont haplotypes (Fig. 7). If we focus on the species which are significantly distributed either below or above the threshold of 600 m a.s.l. three main scenarios emerged: (1) mycobionts with low genetic diversity (*Carbonea vorticosa*, *Lecidea polypycnidophora* and *Lecidella greenii*) are associated with one photobiont OTU *Tr_A02*, and were found in only the lower area; (2) a mycobiont with a high genetic diversity (*Rhizoplaca macleanii*) is still solely associated to one photobiont OTU (*Tr_A02*) and is only located at the high elevated areas and (3) the mycobiont with the highest genetic diversity (*Lecidea cancriformis*) is associated with highest number of photobiont OTUs, in the high elevation sites. These findings are in agreement with the known distribution of *L. greenii* and *Tr_A02* (*Trebouxia URa2*), which, so far, have only been reported for sites in the more northern parts of the Ross Sea region and have never been found at the most extreme southern environments like the Darwin area (Ruprecht et al. 2012a; Ruprecht et al. 2012b). In contrast, *L. cancriformis* is one of the most widespread lichens, being distributed all over Continental Antarctica and is associated with all known photobiont species (Castello 2003; Ruprecht et al. 2012a; Ruprecht et al. 2010).

The above results suggest that the mycobionts are dependent on the availability of climatically adapted photobionts. However, the mycobionts seem to have also their unique climate specific preferences because they do not make use of the whole climate niche of the associated photobionts. These findings are only partially in line with previous studies (e.g. Romeike et al. 2002; Wirtz et al. 2003) that suggest that in extreme environments like the Antarctic continent there might be a selection pressure against photobiont specificity so that more versatile mycobionts are favored. Flexibility concerning the partner choice has been considered as an adaptive strategy to survive harsher environmental conditions (Leavitt et al. 2013; Singh et al. 2017; Werth and Sork 2010).

To explain the myco- and photobiont distribution we need to understand what abiotic factors control terrestrial life in these polar ecosystems. For the MDV Lee et al. (2019) state clearly that one of the most important factors governing the distribution of taxa in

the MDV is temperature, which is accepted to be inversely correlated to elevation (McKay 2015). However, much less is known about the conditions for wetness and humidity. The available wetness index for the MDV quantifies the expected wetness of a unit within the watershed by calculating the amount of possible water flowing into that unit from estimated snow fall (Stichbury et al. 2011). For rock associated lichens this is not relevant, because they are not connected to this source of water. They are, therefore, dependent on moisture provided by the very low precipitation, infrequent melting snow (Head and Marchant 2014) and humidity provided by incoming fog and clouds from the sea. Additionally, it is now clear if the occasional foehn wind events cause severe drying within the valleys at altitudes up to 500 m. At higher elevations there is cold and moister air and this establishes a strong moisture availability gradient with elevation (Speirs et al. 2010; Fig. 1d). Our results suggest that our defined elevation threshold of about 600 m a.s.l. is a reasonable level which marks the shift from lower, dryer to higher, more humid conditions. Habitat aspect is also known to be important. Yung et al. (2014) described large differences with respect to just aspect for their chasmoendolithic microbial communities at Miers Valley. Similar results were also reported for the more maritime site, Botany Bay (Seppelt et al. 2010). We did not find any impact of other topographical features such as distance to coast, slope, aspect and substrate. The collection sites were mainly N-facing or on plateaus, our transects were narrow and consistently only five to ten km inland plus the underlying rock in the whole area is granite and the investigated lichens are restricted to siliceous rock (Ruprecht et al. 2012b; Ruprecht et al. 2010). However, our sampling was equally distributed below and above the threshold of 600 m a.s.l., so the differences found for species distribution, genetic diversity and specificity appears to be due to the changing climate conditions, particularly moisture, along the elevational gradient.

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Online Resource 1

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Online Resource 1a: Samples used in this study, with information on haplotypes, collecting localities and Genbank accession numbers of different markers. To avoid redundant data accumulation, for every haplotype only one reference sequence was uploaded to Genbank.

voucher_ID	area	elevation	latitude	longitude	Mycobiont					Associated green micro algae (<i>Trebouxia</i>)					
					species_name	HT	Accession numbers			OTU_id	HT	Accession numbers			
							nrITS	mtSSU	RPB1			nrITS	psbJ-L	COX2	
T46643	4	602	-78.031	163.865	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	-	Tr_A02	02	MK970698	-	-
T46647a	4	794	-78.033	163.898	<i>Buellia</i> sp. 1	01	MK970689	-	-	-	Tr_A02	01	MK970699	-	-
T46647b	4	794	-78.033	163.898	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	MK970665	MN023039	MN023053	-	Tr_A02	02	MK970698	-	-
T46651	4	374	-78.028	163.851	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	-	-	-	-	-	-	-	-
T46659	4	363	-78.023	163.903	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	-	Tr_A02	02	MK970698	-	-
T46672	4	397	-78.024	163.898	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	-	Tr_A02	02	MK970698	-	-
T46673	4	365	-78.027	163.851	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	-	Tr_A02	01	MK970699	-	-
T46676	4	359	-78.028	163.851	<i>Lecidella greenii</i> Ruprecht & Türk	02	MK970671	-	-	-	Tr_A02	01	MK970699	-	-
T46677	4	843	-78.027	163.848	<i>Lecidea cancriformis</i> Dodge & Baker	04	MK970681	-	-	-	Tr_A02	05	MK970696	-	-
T46678	4	843	-78.028	163.850	<i>Rhizoplaca macleanii</i> (Dodge) Castello	08	MK970669	-	-	-	Tr_A02	01	MK970699	-	-
T46679	4	790	-78.036	163.971	<i>Rhizoplaca macleanii</i> (Dodge) Castello	04	MK970664	MN023037	-	-	Tr_A02	05	MK970696	-	-
T46680	4	360	-78.036	163.971	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	MN023043	MN023053	-	Tr_A02	01	MK970699	-	-
T46681	4	376	-78.032	163.951	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	-	Tr_A02	02	MK970698	-	-
T46683	4	381	-78.028	163.824	<i>Huea</i> sp. 1	01	MK970690	-	MN023048	-	Tr_A02	02	MK970698	-	-
T46684	4	870	-78.044	163.986	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	-	-	Tr_S02	01	MK970693	-	-
T46685	4	870	-78.044	163.986	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	MN023056	-	Tr_S02	01	MK970693	-	MN023030
T46687	4	340	-78.028	163.853	<i>Huea</i> sp. 1	01	MK970690	-	-	-	Tr_A02	02	MK970698	-	-
T46699	4	405	-78.020	163.815	<i>Huea</i> sp. 1	01	MK970690	MN023041	-	-	Tr_A02	01	MK970699	-	-
T46701	4	390	-78.020	163.805	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	MN023034	-	-	Tr_A02	02	MK970698	-	-
T46706	4	379	-78.028	163.821	<i>Lecidella greenii</i> Ruprecht & Türk	02	MK970671	-	MN023054	-	Tr_A02	02	MK970698	-	-

T46709	3	600	-78.073	163.717	<i>Buellia</i> sp. 3	01	MK970688	MN023047	-	Tr_A02	02	MK970698	-	-
T46710	3	600	-78.073	163.717	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	MK970666	-	-	Tr_A02	02	MK970698	-	-
T46713	4	350	-78.028	163.843	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	MN023043	MN023053	Tr_A02	02	MK970698	-	-
T46716	4	412	-78.040	163.802	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	MN023043	MN023053	Tr_A02	02	MK970698	-	-
T46717	4	442	-78.040	163.806	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T46718	4	442	-78.040	163.807	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	-	-	Tr_A02	02	MK970698	-	-
T46719	4	425	-78.038	163.804	<i>Carbonea</i> sp. URm1	01	MK970657	-	-	Tr_A02	02	MK970698	-	-
T46726	3	890	-78.071	163.785	<i>Austrolecia</i> sp. 4	01	MK970653	-	-	Tr_A02	01	MK970699	-	-
T48769	1	370	-78.126	163.700	<i>Lecidella greenii</i> Ruprecht & Türk	02	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48770	1	373	-78.127	163.690	<i>Lecanora</i> sp. 3	01	MK970659	-	-	Tr_A02	06	MK970701	-	-
T48772	1	393	-78.120	163.684	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48773	1	377	-78.127	163.674	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	MN023043	-	Tr_A02	01	MK970699	MN023065	-
T48774	1	526	-78.135	163.626	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T48776	1	783	-78.165	163.753	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	MN023046	MN023057	Tr_A02	03	MK970702	-	-
T48777	1	766	-78.166	163.755	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	MN023046	MN023058	Tr_S15	01	MK970692	MN023065	MN023031
T48778a	1	700	-78.149	163.769	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T48778b	1	700	-78.149	163.769	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	03	MK970702	-	-
T48779	1	757	-78.164	163.755	<i>Rhizoplaca macleanii</i> (Dodge) Castello	04	MK970664	MN023037	-	Tr_A02	05	MK970696	-	-
T48781	1	745	-78.128	163.620	<i>Lecidea</i> sp. 6	02	MK620097	-	-	Tr_A02	01	MK970699	-	-
T48782	1	679	-78.123	163.642	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	MN023046	-	Tr_S18	01	MK970695	MN023070	MN023032
T48783	1	621	-78.122	163.653	<i>Huea</i> sp. 1	01	MK970690	MN023041	MN023048	Tr_A02	01	MK970699	-	-
T48784	1	423	-78.121	163.683	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	MN023043	-	Tr_A02	01	MK970699	MN023065	-
T48785	1	429	-78.133	163.666	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023054	Tr_S15	01	MK970692	-	-
T48786	1	448	-78.167	163.651	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48787	1	361	-78.127	163.678	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023054	Tr_A02	01	MK970699	-	-
T48788	1	397	-78.120	163.684	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	MN023033	MN023050	Tr_A02	01	MK970699	-	-
T48789	1	860	-78.152	163.739	<i>Carbonea</i> sp. 2	01	MK970654	-	MN023051	Tr_S15	01	MK970692	MN023069	MN023031
T48790a	1	415	-78.120	163.682	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	MN023033	-	Tr_A02	01	MK970699	-	-
T48790b	1	415	-78.120	163.682	<i>Lecanora</i> sp. 3	01	MK970659	-	-	Tr_A02	12	MK970703	-	-
T48791a	1	373	-78.120	163.686	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48792	1	415	-78.120	163.682	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	02	MK970698	-	-
T48793a	1	831	-78.153	163.731	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	MN023056	Tr_S15	01	MK970692	-	MN023031
T48793b	1	831	-78.153	163.731	<i>Carbonea</i> sp. 2	01	MK970654	-	-	Tr_S15	01	MK970692	MN023069	MN023031
T48793c	1	831	-78.153	163.731	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48794a	1	905	-78.151	163.735	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_S15	01	MK970692	MN023069	MN023031
T48794b	1	905	-78.151	163.735	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T48795a	1	853	-78.161	163.714	<i>Rhizoplaca macleanii</i> (Dodge) Castello	07	MK970668	-	-	Tr_A02	02	MK970698	-	-
T48795b	1	853	-78.161	163.714	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	02	MK970698	-	-
T48795c	1	853	-78.161	163.714	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_S15	01	MK970692	MN023069	-
T48796	1	831	-78.159	163.698	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	02	MK970698	-	-
T48797	1	814	-78.156	163.689	<i>Rhizoplaca macleanii</i> (Dodge) Castello	05	MK970667	-	-	Tr_A02	02	MK970698	-	-
T48798	1	911	-78.150	163.736	<i>Rhizoplaca macleanii</i> (Dodge) Castello	05	MK970667	-	-	Tr_A02	02	MK970698	-	-
T48799b	1	743	-78.145	163.620	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	-	-	Tr_A02	02	MK970698	-	-
T48799c	1	743	-78.145	163.620	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	-	Tr_S15	01	MK970692	-	-
T48800	1	729	-78.146	163.631	<i>Lecanora</i> sp. 2	01	MK970662	MN023036	MN023052	Tr_A02	02	MK970698	-	-
T48801a	1	732	-78.144	163.626	<i>Lecanora</i> sp. 3	01	MK970659	-	-	Tr_A02	01	MK970699	-	-
T48801c	1	732	-78.144	163.626	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	MN023040	-	Tr_A02	01	MK970699	-	-
T48802	1	732	-78.147	163.629	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-

T48803a	1	775	-78.148	163.630	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48803b	1	775	-78.148	163.630	<i>Carbonea</i> sp. URm1	01	MK970657	MN023034	-	Tr_A02	01	MK970699	-	-
T48804	1	775	-78.148	163.630	<i>Rhizoplaca macleanii</i> (Dodge) Castello	09	MK970670	-	MN023053	Tr_A02	01	MK970699	-	-
T48805	1	732	-78.144	163.626	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	02	MK970698	-	-
T48806	1	568	-78.142	163.628	<i>Carbonea</i> sp. URm1	01	MK970657	MN023034	-	Tr_A02	01	MK970699	-	-
T48807	1	554	-78.141	163.631	<i>Lecidea andersonii</i> Filson	01	MK970673	MN023042	MN023060	Tr_A02	03	MK970702	-	-
T48809	1	540	-78.148	163.657	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48811a	1	557	-78.142	163.657	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	Tr_A02	02	MK970698	-	-
T48812a	1	540	-78.138	163.619	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48812b	1	540	-78.138	163.619	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	Tr_A02	01	MK970699	-	-
T48813	1	557	-78.142	163.657	<i>Lecidea</i> UCR1	01	MK970675	MN023044	-	Tr_A02	01	MK970699	-	-
T48815	1	516	-78.137	163.627	<i>Austrolecia</i> sp. 1	02	MK970652	-	-	Tr_A02	02	MK970698	-	-
T48816	2	171	-78.100	163.810	<i>Buellia</i> sp. 1	01	MK970689	-	-	Tr_A02	02	MK970698	-	-
T48817a	2	546	-78.113	163.785	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T48817b	2	546	-78.113	163.785	<i>Carbonea</i> sp. 2	02	MK970655	-	-	Tr_A02	03	MK970702	-	-
T48818	1	596	-78.148	163.811	<i>Austrolecia</i> sp. 1	03	MK970651	-	-	Tr_A02	01	MK970699	-	-
T48820	2	575	-78.114	163.780	<i>Carbonea</i> sp. 2	01	MK970654	-	-	Tr_A02	02	MK970698	-	-
T48821	2	589	-78.114	163.779	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	MK970665	-	-	Tr_A02	02	MK970698	-	-
T48823a	2	401	-78.098	163.710	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	-	-	-	-	-	-	-
T48824	2	345	-78.097	163.707	<i>Buellia</i> sp. 1	01	MK970689	-	-	Tr_A02	01	MK970699	-	-
T48825	2	498	-78.097	163.691	<i>Carbonea</i> sp. 2	01	MK970654	MN023035	-	Tr_A02	01	MK970699	-	-
T48826	2	424	-78.097	163.717	<i>Lecanora fuscobrunnea</i> Dodge & Baker	01	MK970661	-	-	Tr_S02	02	MK970694	MN023068	-
T48827a	2	345	-78.097	163.707	<i>Buellia frigida</i> Darbishire	04	MK970687	-	-	Tr_A02	01	MK970699	-	-
T48827b	2	345	-78.097	163.707	<i>Huea</i> sp. 1	01	MK970690	MN023041	-	Tr_A02	03	MK970702	-	-
T48828a	2	406	-78.110	163.858	<i>Lecidella greenii</i> Ruprecht & Türk	02	MK970671	-	MN023054	-	-	-	-	-
T48828b	2	406	-78.110	163.858	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	02	MK970698	-	-
T48829	2	453	-78.111	163.858	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48831	2	451	-78.111	163.858	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	MN023033	-	Tr_A02	01	MK970699	-	-
T48832	2	526	-78.112	163.824	<i>Carbonea</i> sp. 2	01	MK970654	-	MN023051	Tr_A02	03	MK970702	-	-
T48834	2	513	-78.114	163.854	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	-	Tr_S15	01	MK970692	-	-
T48835	2	522	-78.112	163.820	<i>Buellia</i> sp. 1	01	MK970689	-	-	Tr_A02	01	MK970699	-	-
T48836	2	179	-78.099	163.778	<i>Lecanora cf. mons-nivis</i> Darbishire	01	MK970658	-	-	Tr_A02	01	MK970699	-	-
T48837	2	179	-78.098	163.777	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	Tr_A02	01	MK970699	-	-
T48839	2	514	-78.114	163.854	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	Tr_A02	01	MK970699	-	-
T48840	3	722	-78.068	163.861	<i>Buellia</i> sp. 1	01	MK970689	-	-	Tr_A02	02	MK970698	-	-
T48841a	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-
T48841b	3	722	-78.068	163.861	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	-	MN023050	Tr_A02	02	MK970698	-	-
T48843a	3	836	-78.066	163.870	<i>Lecidea cancriformis</i> Dodge & Baker	01	MK970677	-	-	Tr_S02	01	MK970693	-	-
T48843b	3	836	-78.066	163.870	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_S15	01	MK970692	MN023069	-
T48843c	3	836	-78.066	163.870	<i>Lecidea cancriformis</i> Dodge & Baker	05	MK970678	-	-	-	-	-	-	-
T48843d	3	836	-78.066	163.870	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	01	MK970699	-	-
T48843e	3	836	-78.066	163.870	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	03	MK970702	-	-
T48844	3	749	-78.067	163.863	<i>Lecidella greenii</i> Ruprecht & Türk	02	MK970671	-	-	Tr_A02	02	MK970698	-	-
T48849	3	509	-78.077	163.783	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48851	3	587	-78.074	163.793	<i>Lecanora fuscobrunnea</i> Dodge & Baker	02	MK970660	MN023037	-	Tr_A02	02	MK970698	-	-
T48852	3	587	-78.075	163.792	<i>Huea</i> sp. 1	01	MK970690	-	-	Tr_A02	01	MK970699	-	-
T48853	3	594	-78.075	163.791	<i>Huea</i> sp. 1	01	MK970690	-	MN023048	Tr_A02	01	MK970699	-	-
T48855a	4	688	-78.036	163.837	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-

T48855b	4	688	-78.036	163.837	<i>Lecidea cancriformis</i> Dodge & Baker	03	MK970680	MN023046	MN023059	Tr_S15	01	MK970692	MN023069	MN023031
T48857a	4	688	-78.036	163.827	<i>Lecidea</i> sp. 6	01	MK970684	MN023045	-	Tr_A02	01	MK970699	-	-
T48857b	4	688	-78.036	163.827	<i>Buellia</i> sp. 1	01	MK970689	-	-	Tr_S15	01	MK970692	-	MN023031
T48857c	4	688	-78.036	163.827	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	02	MK970698	-	-
T48858	4	319	-78.025	163.975	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	-	-	-	-	-
T48859	4	319	-78.025	163.986	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	-	MN023053	Tr_A02	02	MK970698	-	-
T48860a	4	692	-78.036	163.836	<i>Huea</i> sp. 1	01	MK970690	-	MN023048	Tr_A02	01	MK970699	-	-
T48860b	4	694	-78.036	163.836	<i>Lecidea</i> sp. 6	01	MK970684	MN023045	MN023064	Tr_A02	01	MK970699	-	-
T48860c	4	694	-78.036	163.836	<i>Huea</i> sp. 1	01	MK970690	-	-	Tr_A02	02	MK970698	-	-
T48860d	4	694	-78.036	163.836	<i>Huea</i> sp. 1	01	MK970690	-	-	Tr_A02	01	MK970699	-	-
T48861	4	380	-78.024	163.892	<i>Lecanora</i> sp. 3	01	MK970659	-	-	Tr_A02	01	MK970699	-	-
T48862	4	875	-78.037	163.978	<i>Lecidea cancriformis</i> Dodge & Baker	06	MK970682	-	-	-	-	-	-	-
T48864	4	850	-78.039	163.989	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	-	-	Tr_A02	03	MK970702	-	-
T48865a	4	874	-78.036	163.990	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	MK970666	-	-	Tr_A02	02	MK970698	MN023067	-
T48867a	4	936	-78.043	164.104	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	05	MK970696	-	-
T48867b	4	936	-78.043	164.104	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	01	MK970699	-	-
T48867c	4	936	-78.043	164.104	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	MK970666	-	-	Tr_A02	03	MK970702	-	-
T48869	4	849	-78.037	163.978	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	-	-	-	-	-
T48872	4	673	-78.030	163.951	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	-	-	-	-	-	-
T48873	4	455	-78.024	163.893	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023054	Tr_A02	02	MK970698	-	-
T48874	4	375	-78.024	163.899	<i>Lecidea</i> UCR1	02	MK970676	MN023044	MN023062	Tr_A02	11	MK970697	MN023066	-
T48875	4	375	-78.024	163.900	<i>Lecidea lapicida</i> (Ach.) Ach. subsp.	01	MK970683	-	-	Tr_A02	01	MK970699	-	-
T48876a	4	380	-78.024	163.900	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	MN023040	-	Tr_A02	02	MK970698	-	-
T48876b	4	380	-78.024	163.900	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	Tr_A02	01	MK970699	-	-
T48877	4	385	-78.024	163.900	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	MK970656	-	MN023050	Tr_A02	02	MK970698	-	-
T48879	3	419	-78.057	163.747	<i>Lecidella greenii</i> Ruprecht & Türk	01	MK970671	-	MN023055	-	-	-	-	-
T48880	3	443	-78.058	163.740	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970674	-	-	-	-	-	-	-
T48881a	3	541	-78.061	163.791	<i>Huea</i> sp. 1	01	MK970690	-	MN023049	Tr_A02	03	MK970702	-	-
T48881b	3	541	-78.061	163.791	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	MK970666	-	-	Tr_A02	01	MK970699	-	-
T48882	3	575	-78.057	163.817	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	MK970663	-	-	Tr_A02	01	MK970699	MN023065	-
T48883a	3	671	-78.058	163.847	<i>Huea</i> sp. 1	01	MK970690	-	-	Tr_A02	02	MK970698	-	-
T48883b	3	671	-78.058	163.847	<i>Lecidea</i> sp. 5	01	MK620099	-	MN023063	Tr_A02	09	MK970700	-	-
T48885	3	666	-78.057	163.844	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	MK970666	-	-	Tr_A02	02	MK970698	-	-
T48886	4	464	-78.025	163.932	<i>Austrolecia</i> sp. 1	01	MK970650	-	-	Tr_A02	02	MK970698	-	-
T48887a	3	710	-78.070	163.711	<i>Lecidea cancriformis</i> Dodge & Baker	02	MK970679	MN023046	-	Tr_S15	01	MK970692	MN023069	-
T48887b	3	710	-78.070	163.711	<i>Huea</i> sp. 1	01	MK970690	-	-	Tr_A02	03	MK970702	-	-
T48888b	3	629	-78.073	163.717	<i>Lecanora cf. mons-nivis</i> Darbishire	01	MK970658	-	-	Tr_A02	01	MK970699	-	-
T48900	4	610	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	MK970663	-	-	Tr_A02	02	MK970698	-	-

Online Resource 1b: Additional samples taken from Perez-Ortega et al. (2012) and used in this study, with information on haplotypes, collecting localities and Genbank accession numbers.

Laboratory code	area	elevation	latitude	longitude	Mycobiont			Associated green micro algae		
					species_name	HT	Accession numbers nrITS	OTU_id	HT	Accession numbers nrITS
s106	2	566	-78.113	163.782	<i>Lecanora</i> sp. 2	01	JX036037	Tr_A02	02	JX036159
s111	2	520	-78.114	163.854	<i>Buellia frigida</i> Darbshire	01	JX036042	Tr_A02	01	JX036164
s112	2	520	-78.114	163.854	<i>Buellia</i> sp. 2	01	JX036043	Tr_A02	01	JX036165
s113	2	520	-78.114	163.854	<i>Lecidea cancriformis</i> Dodge & Baker	01	JX036044	Tr_S15	01	JX036166
s114	2	520	-78.114	163.854	<i>Lecidea cancriformis</i> Dodge & Baker	01	JX036045	Tr_A02	05	JX036167
s115	3	839	-78.066	163.870	<i>Lecidella greenii</i> Ruprecht & Türk	02	JX036046	Tr_A02	03	JX036168
s117	3	839	-78.066	163.870	<i>Buellia frigida</i> Darbshire	01	JX036048	Tr_A02	03	JX036169
s118	3	839	-78.066	163.870	<i>Buellia frigida</i> Darbshire	01	JX036049	Tr_A02	02	JX036170
s119	3	411	-78.083	163.768	<i>Austrolecia</i> sp. 1	01	JX036050	Tr_A02	01	JX036171
s120	3	411	-78.083	163.768	<i>Lecidea polypycnidophora</i> Ruprecht & Türk	01	JX036051	Tr_A02	02	JX036172
s121	4	384	-78.024	163.900	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	JX036052	Tr_A02	01	JX036173
s122	2	413	-78.110	163.787	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	JX036053	Tr_A02	01	JX036174
s123	2	413	-78.110	163.787	<i>Carbonea vorticosa</i> (Flörke) Hertel	01	JX036054	Tr_A02	01	JX036175
s124	3	666	-78.057	163.844	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	JX036055	Tr_A02	02	JX036176
s125	3	666	-78.057	163.844	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	JX036056	Tr_A02	02	JX036177
s126	3	541	-78.061	163.791	<i>Huea</i> sp. 1	01	JX036057	Tr_A02	03	JX036178
s127	4	874	-78.036	163.990	<i>Buellia frigida</i> Darbshire	01	JX036058	Tr_A02	07	JX036179
s128	3	581	-78.075	163.790	<i>Austrolecia</i> sp. 1	01	JX036059	Tr_A02	10	JX036180
s165	2	465	-78.097	163.691	<i>Buellia frigida</i> Darbshire	01	JX036070	Tr_A02	02	JX036191
s166	2	597	-78.097	163.691	<i>Buellia frigida</i> Darbshire	01	JX036071	Tr_A02	01	JX036192
s168	4	688	-78.036	163.837	<i>Austrolecia</i> sp. 1	02	JX036073	Tr_A02	04	JX036194
s171	4	688	-78.036	163.837	<i>Lecanora</i> sp. 2	01	JX036076	Tr_A02	01	JX036197
s173	2	521	-78.114	163.854	<i>Lecanora</i> sp. 2	01	JX036078	Tr_A02	01	JX036199
s174	4	597	-78.033	163.849	<i>Buellia frigida</i> Darbshire	01	JX036079	Tr_A02	02	JX036200
s175	4	597	-78.033	163.849	<i>Lecanora</i> sp. 2	01	JX036080	Tr_A02	04	JX036201
s176	3	658	-78.073	163.712	<i>Buellia frigida</i> Darbshire	01	JX036081	Tr_A02	01	JX036202
s177	2	521	-78.114	163.854	<i>Austrolecia</i> sp. 2	01	JX036082	Tr_A02	01	JX036203
s178	2	521	-78.114	163.854	<i>Austrolecia</i> sp. 2	02	JX036083	Tr_A02	01	JX036204
s179	3	540	-78.063	163.809	<i>Lecanora</i> sp. 2	01	JX036084	Tr_A02	01	JX036205
s180	3	540	-78.063	163.809	<i>Buellia frigida</i> Darbshire	01	JX036085	Tr_A02	01	JX036206
s181	2	453	-78.111	163.858	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036086	Tr_A02	04	JX036207
s182	4	688	-78.036	163.837	<i>Austrolecia</i> sp. 1	01	JX036087	Tr_A02	06	JX036208
s184	3	615	-78.073	163.717	<i>Austrolecia</i> sp. 1	01	JX036089	Tr_A02	02	JX036210
s187	2	575	-78.114	163.778	<i>Buellia frigida</i> Darbshire	02	JX036092	Tr_A02	13	JX036213
s189	3	719	-78.071	163.712	<i>Buellia frigida</i> Darbshire	01	JX036094	Tr_A02	02	JX036215
s190	4	464	-78.025	163.899	<i>Lecanora</i> sp. 3	02	JX036095	Tr_A02	02	JX036216
s191	4	419	-78.025	163.900	<i>Lecanora</i> sp. 3	02	JX036096	Tr_A02	01	JX036217
s192	4	481	-78.030	163.834	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036097	Tr_A02	01	JX036218

s195	4	874	-78.036	163.992	<i>Buellia frigida</i> Darbshire	01	JX036100	Tr_A02	01	JX036221
s197	3	541	-78.061	163.791	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	JX036101	Tr_A02	01	JX036222
s198	3	541	-78.061	163.791	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	JX036102	Tr_A02	01	JX036223
s199	4	375	-78.024	163.900	<i>Austrolecia</i> sp. 1	01	JX036103	Tr_A02	02	JX036224
s200	4	375	-78.024	163.900	<i>Austrolecia</i> sp. 1	01	JX036104	Tr_A02	02	JX036225
s201	4	375	-78.024	163.900	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	JX036105	Tr_A02	02	JX036226
s202	4	360	-78.027	163.839	<i>Lecanora</i> cf. <i>mons-nivis</i> Darbshire	01	JX036106	Tr_A02	08	JX036227
s203	4	360	-78.027	163.839	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036107	Tr_A02	01	JX036228
s205	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	JX036108	Tr_A02	04	JX036229
s206	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	JX036109	Tr_A02	02	JX036230
s207	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	JX036110	Tr_A02	08	JX036231
s208	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036111	Tr_A02	02	JX036232
s209	3	722	-78.068	163.861	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036112	Tr_A02	01	JX036233
s212	4	710	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036115	Tr_A02	02	JX036236
s213	4	710	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036116	Tr_A02	02	JX036237
s214	4	710	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036117	Tr_A02	02	JX036238
s215	3	710	-78.070	163.711	<i>Lecanora fuscobrunnea</i> Dodge & Baker	01	JX036118	Tr_A02	02	JX036239
s221	2	508	-78.112	163.823	<i>Buellia frigida</i> Darbshire	01	JX036124	Tr_A02	01	JX036245
s229	4	710	-78.034	163.845	<i>Huea</i> sp. 1	01	JX036132	Tr_A02	03	JX036251
s230	4	710	-78.034	163.845	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036133	Tr_A02	02	JX036252
s231	4	710	-78.034	163.845	<i>Austrolecia</i> sp. 1	01	JX036134	Tr_A02	03	JX036253
s232	4	710	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036135	Tr_A02	01	JX036254
s233	4	710	-78.034	163.845	<i>Rhizoplaca macleanii</i> (Dodge) Castello	02	JX036136	Tr_A02	04	JX036255
s235	2	589	-78.113	163.778	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036138	Tr_A02	02	JX036257
s236	2	589	-78.113	163.778	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036139	Tr_A02	02	JX036258
s237	2	589	-78.113	163.778	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036140	Tr_A02	07	JX036259
s266	3	583	-78.075	163.791	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036141	Tr_A02	04	JX036260
s267	4	597	-78.033	163.849	<i>Buellia frigida</i> Darbshire	01	JX036142	Tr_A02	01	JX036261
s269	2	566	-78.113	163.782	<i>Huea</i> sp. 1	01	JX036143	Tr_A02	01	JX036262
s270	2	566	-78.113	163.782	<i>Huea</i> sp. 1	01	JX036144	Tr_A02	01	JX036263
s271	4	959	-78.047	164.104	<i>Rhizoplaca macleanii</i> (Dodge) Castello	03	JX036145	Tr_A02	02	JX036264
s272	4	809	-78.035	163.978	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036146	Tr_A02	01	JX036265
s273	4	688	-78.030	163.949	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036147	Tr_A02	01	JX036266
s274	4	874	-78.036	163.990	<i>Rhizoplaca macleanii</i> (Dodge) Castello	01	JX036148	Tr_A02	01	JX036267
s275	4	874	-78.036	163.990	<i>Lecanora</i> sp. 1	01	JX036149	Tr_A02	03	JX036268
s300	4	481	-78.030	163.834	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036150	Tr_A02	01	JX036269
s301	4	481	-78.030	163.834	<i>Lecidella greenii</i> Ruprecht & Türk	01	JX036151	Tr_A02	01	JX036270
s95	4	375	-78.024	163.900	<i>Rhizoplaca macleanii</i> (Dodge) Castello	06	JX036152	Tr_A02	04	JX036271
s96	3	585	-78.075	163.791	<i>Buellia frigida</i> Darbshire	03	JX036153	Tr_A02	05	JX036272
s97	2	575	-78.114	163.814	<i>Austrolecia</i> sp. 1	01	JX036154	Tr_S18	01	JX036273

Online Resource 1c: The pairs of *min10MycoSp* species and *min10PhoHap* haplotypes that showed significant differences with regards to elevation of sampling sites, as well as the associated p-values of the mctp-tests. The pairs are listed line-by-line; the species with the samples found in higher elevations is given on the left, respectively. PH = photobiont haplotypes

Elevation of Sample Sites in Pairwise Comparison:		p-value	
higher	lower		
Mycobiont Species	<i>Austrolecia</i> sp. 1	<i>Carbonea vorticosa</i>	0.012
	<i>Austrolecia</i> sp. 1	<i>Lecidea polypycnidophora</i>	0.001
	<i>Buellia frigida</i>	<i>Carbonea vorticosa</i>	0.027
	<i>Buellia frigida</i>	<i>Lecidea polypycnidophora</i>	0.003
	<i>Huea</i> sp. 1	<i>Lecidea polypycnidophora</i>	0.049
	<i>Lecidea cancriformis</i>	<i>Carbonea vorticosa</i>	0.000
	<i>Lecidea cancriformis</i>	<i>Huea</i> sp. 1	0.017
	<i>Lecidea cancriformis</i>	<i>Lecidea polypycnidophora</i>	0.000
	<i>Lecidea cancriformis</i>	<i>Lecidella greenii</i>	0.000
	<i>Rhizoplaca macleanii</i>	<i>Carbonea vorticosa</i>	0.001
	<i>Rhizoplaca macleanii</i>	<i>Lecidea polypycnidophora</i>	0.000
	<i>Rhizoplaca macleanii</i>	<i>Lecidella greenii</i>	0.002
PH	Tr_A02_h03	Tr_A02_h01	0.021
	Tr_S15_h01	Tr_A02_h01	0.011

Online Resource 1d Network matrix giving the number of associations between the mycobiont species and photobiont haplotypes

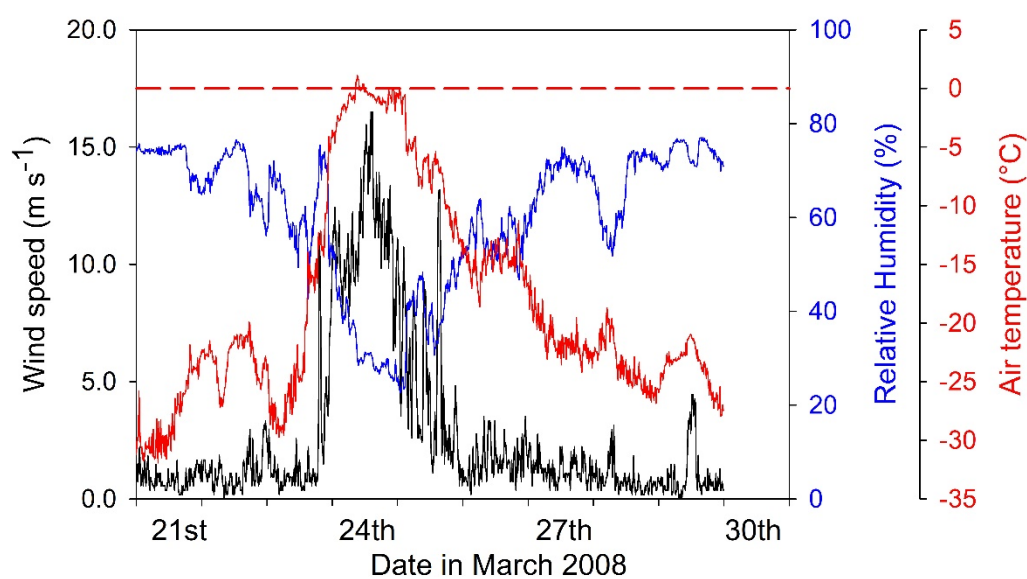
	Photobiont Haplotypes																
	Tr_A02_h01	Tr_A02_h02	Tr_A02_h03	Tr_A02_h04	Tr_A02_h05	Tr_A02_h06	Tr_A02_h07	Tr_A02_h08	Tr_A02_h09	Tr_A02_h10	Tr_A02_h11	Tr_A02_h12	Tr_A02_h13	Tr_S02_h01	Tr_S02_h02	Tr_S15_h01	Tr_S18_h01
<i>Austrolecia</i> sp. 1	10	9	2	1	-	1	-	-	-	1	-	-	-	-	-	3	1
<i>Austrolecia</i> sp. 2	2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Austrolecia</i> sp. 4	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Buellia frigida</i>	8	4	1	-	1	-	1	-	-	-	-	-	1	-	-	-	-
<i>Buellia</i> sp. 1	3	2	-	-	-	-	-	-	-	-	-	-	-	-	-	1	-
<i>Buellia</i> sp. 2	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Buellia</i> sp. 3	-	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Carbonea</i> sp. 2	1	1	2	-	-	-	-	-	-	-	-	-	-	-	-	2	-
<i>Carbonea</i> URm1	2	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Carbonea vorticosa</i>	6	3	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Huea</i> sp. 1	8	4	5	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecanora</i> cf. <i>mons-nivis</i>	2	-	-	-	-	-	-	1	-	-	-	-	-	-	-	-	-
<i>Lecanora fuscobrunnea</i>	-	2	-	-	-	-	-	-	-	-	-	-	-	-	1	-	-
<i>Lecanora</i> sp. 1	-	-	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecanora</i> sp. 2	3	2	-	1	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecanora</i> sp. 3	3	1	-	-	-	1	-	-	-	-	1	-	-	-	-	-	-
<i>Lecidea andersonii</i>	-	-	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecidea cancriformis</i>	-	1	2	-	2	-	-	-	-	-	-	-	-	3	-	7	1
<i>Lecidea lapicida</i>	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecidea polypycnidophora</i>	4	5	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecidea</i> sp. 5	-	-	-	-	-	-	-	-	1	-	-	-	-	-	-	-	-
<i>Lecidea</i> sp. 6	3	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
<i>Lecidea</i> UCR1	1	-	-	-	-	-	-	-	-	-	1	-	-	-	-	-	-
<i>Lecidella greenii</i>	16	9	1	2	-	-	-	-	-	-	-	-	-	-	-	1	-
<i>Rhizoplaca macleanii</i>	12	29	2	3	3	-	1	1	-	-	-	-	-	-	-	-	-

Online Resource 2

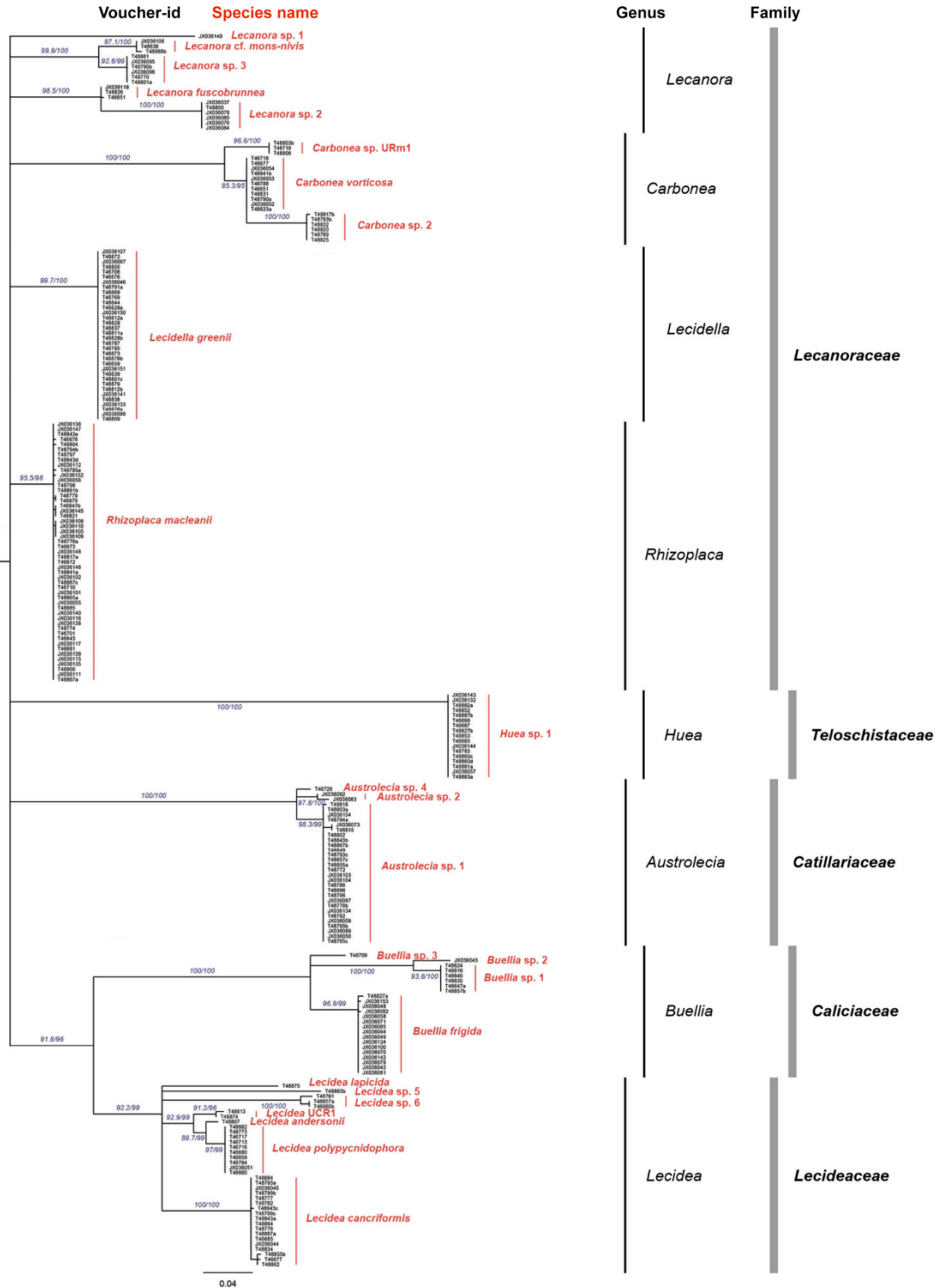
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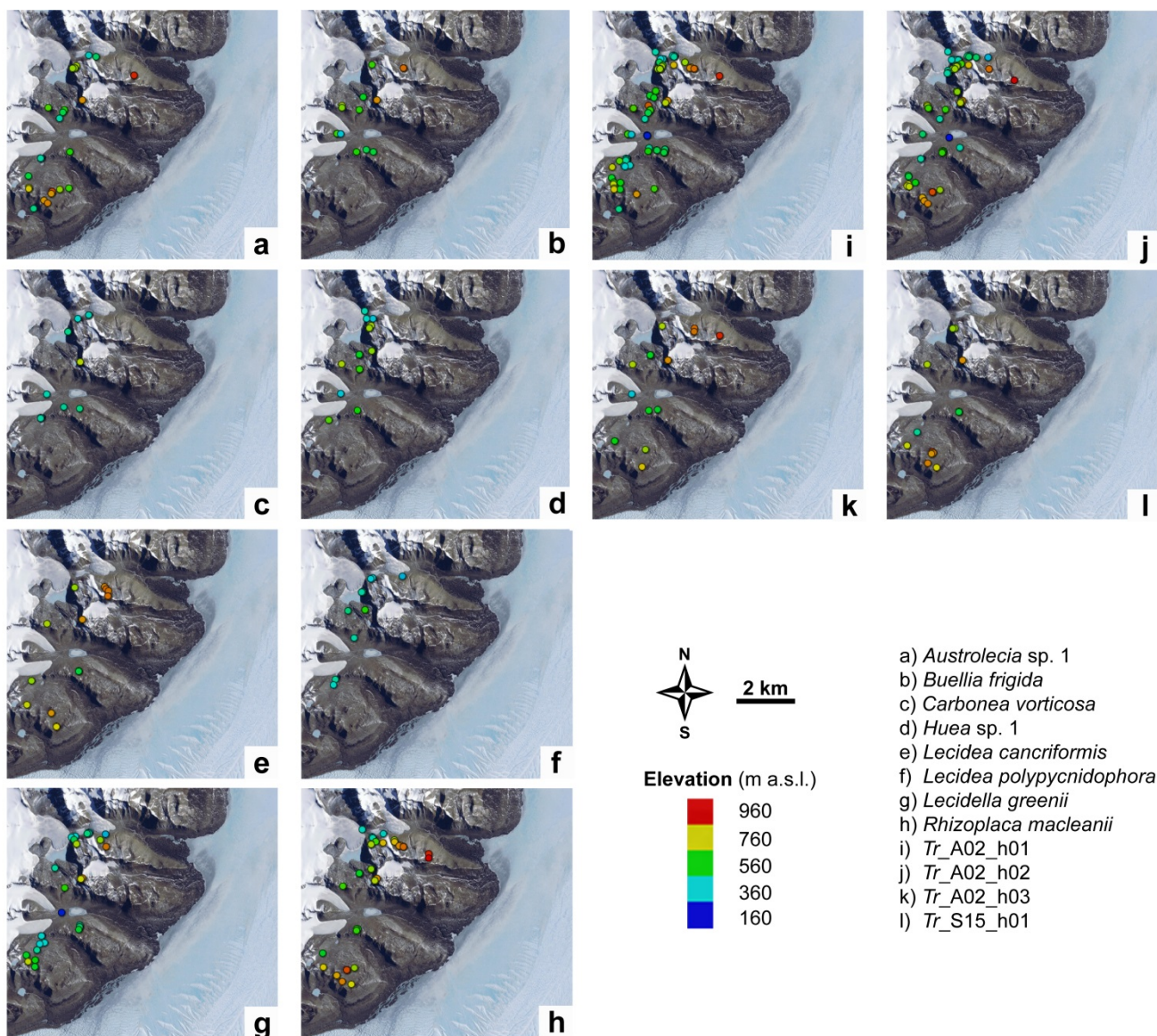
Online Resource 2a: Foehn event in Miers Valley during March 2008



Online Resource 2b: Phylogeny of all mycobionts specimen based on the marker ITS and calculated with IQ-tree. Numbers in italic refer to SH-aLRT and UFboot supports. Branches with SH-aLRT < 80 % and UFboot < 95 % were collapsed.



Online Resource 2d: Sample locations of the mycobiont species (*min10MycoSp*) and photobiont haplotypes (*min10PhoHap*) within the MDV (maps generated with <http://www.gpsvisualizer.com>).



Online Resource 2e: Scatterplots of (a) h/N , (b) Hd , (c) π , (d) d' , (e) NRI , (f) PSV and (g) PSR dependent on elevation means for *min10MycoSp*. r gives the Pearson correlation coefficient.

