



UNIVERSITAT DE
BARCELONA

Departament de Biologia Evolutiva, Ecologia i Ciències Ambientals
Doctorat en Ecologia, Ciències Ambientals i Fisiologia Vegetal

**Resilience of Long-lived Mediterranean
Gorgonians in a Changing World: Insights from
Life History Theory and Quantitative Ecology**

*Memòria presentada per **Ignasi Montero Serra** per optar
al Grau de Doctor per la Universitat de Barcelona*

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A todas las que sueñan con un mundo mejor.

A Latinoamérica.

A Asun y Carlos.

AGRADECIMIENTOS

Echando la vista a atrás reconozco que, pese al estrés del día a día, este ha sido un largo camino de aprendizaje plagado de momentos buenos y alegrías. También ha habido momentos más difíciles, en los cuáles te enfrentas de cara a tus propias limitaciones, pero que te empujan a desarrollar nuevas capacidades y crecer.

Cierro esta etapa agradeciendo a toda la gente que la ha hecho posible, a las oportunidades recibidas, a las enseñanzas de l@s grandes científic@s que me han hecho vibrar en este mundo, al apoyo en los momentos más complicados, a las que me alegraron el día a día, a las que hacen que crea más en mí mismo y, sobre todo, a la gente buena que lucha para hacer de este mundo un lugar mejor y más justo. A tod@s os digo gracias!



GRACIAS!

GRÀCIES!

THANKS!

Advisors' report

Dra. Cristina Linares, professor at Departament de Biologia Evolutiva, Ecologia i Ciències Ambientals (Universitat de Barcelona), and Dr. Joaquim Garrabou, research professor at the Institut de Ciències del Mar (ICM-CSIC), advisors of the PhD thesis entitled “**Resilience of long-lived Mediterranean gorgonians in a changing world: insights from life history theory and quantitative ecology**”,

INFORM, that the research studies developed by **Ignasi Montero Serra** for his Doctoral Thesis have been organized in five chapters, which correspond to five scientific papers listed below: three are already published, one is under review, and the last one is to be submitted in the next months;

and CERTIFY, that the work has been carried out by Ignasi Montero Serra, participating actively in all the tasks: setting the objectives, conceiving and performing the analyses and writing the manuscripts.

Finally, we certify that the co-authors of the publications listed below and that conform this doctoral thesis, will not use these manuscripts in another PhD thesis.

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List and publication status of the chapters of this thesis

CHAPTER I

Montero-Serra I¹, Linares C¹, Doak DF², Ledoux JB^{3,4} & Garrabou J³ (2018) Strong linkages between depth, longevity, and demographic stability. *Proceedings of the Royal Society B*. 285(1873), 20172688. 5-year Impact Factor (2016): 5.417

CHAPTER II

Montero-Serra I¹, Linares C¹, Garcia M³, Pancaldi F³, Frleta-Valic M³, Ledoux JB^{3,4}, Zuberer F⁵, Merad D⁶, Drap P⁶ & Garrabou J³ (2015) Harvesting effects, recovery patterns and management strategies for a long-lived and structural precious coral. *PLoS ONE*. 10(2): e0117250. 5-year Impact Factor (2015): 3.535

CHAPTER III

Montero-Serra I¹, Garrabou J³, Doak DF², Ledoux JB^{3,4} & Linares C¹. Marine protected areas enhance structural complexity but do not buffer the detrimental consequences of ocean warming for an overexploited precious coral. Under review at *Journal of Applied Ecology*. 5-year Impact Factor (2016): 5.989

CHAPTER IV

Montero-Serra I¹, Garrabou J³, Doak DF², Figuerola L¹, Hereu B¹, Ledoux JB^{3,4} & Linares C¹ (2018) Accounting for life-history strategies and timescales in marine restoration. *Conservation Letters*. 11(1), e12341. 5-year Impact Factor (2016): 7.316

CHAPTER V

Montero-Serra I¹, Linares C¹, Mumby PJ⁷, D Macías D⁸, Benoussan N³ & Garrabou J³. Predicting depth-mediated impacts of ocean warming at the end 21st century in Mediterranean coastal ecosystems. In preparation.

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SUMMARY

Temperate benthic communities face cumulative impacts from multiple stressors acting both at local and global scales. Understanding how local management and ocean warming affect the dynamics and resilience of dominant habitat-forming species is central to marine conservation. In this thesis, we combined long-term demographic surveys and large-scale distribution datasets with innovative population and spatial modeling approaches, and meta-analyses to unravel the causes and consequences of extreme life-histories. The final goal was to understand resilience patterns and mechanisms, and to assess the effectiveness of widely used conservation tools such as fishing regulations, marine protected areas (MPAs), and active restoration.

A comparative analysis across marine sessile species revealed strong linkages between depth occurrence, longevity and demographic stability. These results demonstrated a fundamental role of environmental gradients in shaping the evolution of life-history strategies in the ocean and suggest that deep-sea benthic communities tend to be dominated by long-lived species that are very vulnerable to external sources of mortality. In addition, we demonstrate that life-history strategies play a role in important conservation issues such as recovery processes. For instance, recovery times after fishing or restoration actions for long-lived species can take several decades. Thus, accounting for the temporal dimension and life-history tradeoffs in conservation efforts is essential to avoid potential long-lasting impacts and enhance the recovery of damaged ecosystems.

Focusing on the Mediterranean red coral *Corallium rubrum*, an overharvested precious coral emblematic of coralligenous assemblages, we revealed an extremely slow life-history strategy characterized by low reproduction success, high natural survival, and extended longevity. These traits drive a general pattern of slow population dynamics and suggest low resilience to human-driven stressors.

We then show that local and global stressors such as overfishing and warming have strong impacts on different demographic processes of red coral populations.

Harvesting causes dramatic decreases in total biomass and strong shifts in size-class distribution towards populations dominated by small colonies. More broadly, MPAs strongly enhance structural complexity of red coral populations but, contrary to prior expectations, have little effect on their long-term viability and associated extinction risk. Population recovery after harvesting is a very slow process that can take decades. Interestingly, recovery is not driven by sexual reproduction but by the capacity of harvested colonies to survive and regrow new branches. This recovery mechanism may explain the persistence of this historically overexploited long-lived coral.

Unfortunately, novel impacts associated to climate change threaten shallow red coral populations. Recurrent warming-driven mass mortality events had detrimental effects on affected populations, causing long-term declines and potential local extinction. While MPAs have been proposed to enhance the resilience of marine ecosystems to climate change, our simulations suggested that MPAs only have a weak buffering effect to climatic impacts. On the other hand, spatial analyses revealed that future climate change may cause extensive impacts on shallow populations of the red coral and the red gorgonian *Paramuricea clavata* (another emblematic species of coralligenous assemblages with an important structural role) across the Mediterranean Sea. Importantly, we found that the extent and severity of warming impacts is strongly dependent on depth and the global emissions scenarios. Altogether, our results demonstrate a high vulnerability and low demographic resilience of shallow red coral populations to climate change.

The results presented in this thesis suggest that, only by simultaneously acting at both local and global scales, we can ensure the persistence and enhance the structural role of the long-lived Mediterranean red coral. Finally, this thesis also shows the potential of combining long-term large-scale field data, quantitative tools, and principles of life-history theory to provide new perspectives to advance marine conservation in a changing ocean.

RESUM

Els ecosistemes bentònics temperats estan exposats a diversos impactes derivats de l'activitat humana a escales locals i globals. Entendre com eines de gestió en l'àmbit local i els impactes derivats de l'escalfament dels oceans afecten a les dinàmiques i resiliència d'espècies estructurals marines és de vital importància per millorar la seva conservació. Al llarg d'aquesta tesi Doctoral, hem combinat seguiments demogràfics a llarg termini i dades de distribució d'espècies a gran escala amb innovadores tècniques de modelització i meta-anàlisis, per entendre les causes i les conseqüències derivades d'estratègies de vida molt llongeves. L'objectiu final era explorar els patrons i mecanismes claus en la resiliència d'espècies estructurals llongeves i avaluar l'eficàcia de diferents eines de conservació marina com són les regulacions de pesca, les àrees marines protegides i les accions de restauració ecològica.

Mitjançant una anàlisi comparativa d'espècies sèssils marines, vam observar una forta correlació positiva entre la fondària, la longevitat i l'estabilitat demogràfica. Aquests resultats demostren un paper fonamental dels gradients ambientals en els processos d'evolució de les històries de vida en els oceans i suggereixen que les comunitats bentòniques d'ecosistemes profunds tendeixen a ser dominades per espècies llongeves i, per tant, molt vulnerables a pertorbacions externes que incrementen la mortalitat. A més, demostrem que la història de vida de les espècies juga un paper clau en temes importants relacionats amb la conservació com són els temps de recuperació de les poblacions. Per exemple, el temps de recuperació d'espècies llongeves després de pesca o d'accions de restauració ecològica poden ser de l'ordre de dècades. Per tant, és important incorporar l'escala temporal i les històries de vida en les accions de conservació per tal de reduir possibles impactes amb conseqüències negatives a llarg termini i potenciar els processos de recuperació dels ecosistemes malmesos.

La tesi es va enfocar principalment en el corall vermell del Mediterrani, *Corallium rubrum*, com a espècies model, ja que és un corall emblemàtic i amb una funció estructural dintre de les comunitats del coral·ligen. Vam caracteritzar

la seva història de vida, amb un èxit reproductor molt limitat, una baixa mortalitat natural i una longevitat potencial de l'ordre de centenars d'anys. Aquests trets resulten en un patró general de dinàmiques poblacionals molt lentes que, a més, suggereixen una baixa resiliència del corall vermell enfront a perturbacions externes. De fet, perturbacions a escala local com és la pesca i a escala global com és l'escalfament dels oceans tenen un gran impacte en diferents aspectes de la dinàmica del corall vermell.

La pesca del corall vermell ha causat una gran davallada de la biomassa total i canvis en l'estructura de talles cap a poblacions dominades per colònies petites. A gran escala, les àrees marines protegides (AMPs) afavoreixen i incrementen la complexitat estructural de les poblacions de corall vermell però, contràriament a les hipòtesis inicials, les AMPs no afecten de manera significativa en la viabilitat poblacional a llarg termini ni en el risc d'extinció de les poblacions de corall vermell. La recuperació del corall després d'accions de pesca és un procés molt lent que pot trigar dècades. A més, vam demostrar que la recuperació no ve donada per mecanismes de reproducció sexual, sinó per una elevada taxa de supervivència de les colònies pescades i el rebroll de branques noves a partir de les bases de les colònies supervivents. Aquest mecanisme de recuperació podria explicar la persistència del corall vermell al Mediterrani després de mil·lennis de sobreexplotació.

Noves amenaces associades al canvi climàtic estan posant en risc algunes poblacions de poca fondària de corall vermell. Les darreres mortalitats en massa de corall vermell relacionades amb l'escalfament del mar tenen conseqüències molt negatives a llarg-termini, amb importants declivis poblacionals i increment del risc d'extinció local. Mentre les AMPs han estat proposades com a eina per augmentar la resiliència dels ecosistemes marins enfront a perturbacions derivades del canvi global, malauradament, els nostres resultats mostren un paper molt limitat en l'atenuació impactes climàtics. Per altra banda, anàlisis predictives espacials suggereixen que, sota futurs escenaris climàtics, les poblacions de corall vermell i gorgònia vermella *Paramuricea clavata* (una altra espècie estructural i emblemàtica de les comunitats de coral·ligen) podrien patir impactes a gran

escala a diverses regions del Mediterrani. Però el que és més interessant és que la intensitat i extensió dels impactes climàtics estan determinats en gran mesura amb els patrons globals d'emissions de gasos d'efecte hivernacle i amb la fondària a la qual es trobin les poblacions. Aquests resultats confirmen que les poblacions poc profundes de corall vermell són molt vulnerables a l'escalfament dels oceans i tenen una baixa resiliència demogràfica.

Al llarg de la present tesi hem mostrat com, només amb accions de gestió simultànies a escala local i global podrem assegurar la persistència i la funció estructural de les poblacions de corall vermell. Finalment, amb aquesta tesi també demostrarem el gran interès de combinar dades de camp a llarg termini i a gran escala amb tècniques d'ecologia quantitativa i principis de la teoria d'històries de vida per tal de proporcionar noves perspectives i avançar en el camp de la conservació marina en un context de canvi global.

GENERAL INTRODUCTION

Marine biodiversity, functioning and habitat-forming species

Life in the oceans is incredibly diverse in shapes, forms and functions. The complexity of marine ecosystems has long fascinated ecologists and naturalists since ancient times. Besides fulfilling human curiosity, a deep understanding of ecological dynamics and underlying drivers is essential to develop sound evidence-based conservation strategies. A growing body of literature recognizes that biodiversity at all hierarchical levels, from genes to ecosystems, shapes key structural and dynamical properties such as primary productivity, nutrient cycling, functional diversity, trophic complexity, temporal stability, and resilience (Tilman et al. 1996, 2001; Cardinale et al. 2012, Duffy et al. 2016). Importantly, these processes underpin goods and services provide by marine ecosystems to a large proportion of world's human population (Worm et al. 2006, Palumbi et al. 2008, More et al. 2011, Hoegh-Guldberg et al. 2015).

In coastal marine ecosystems, dominant sessile organisms such as seagrass meadows and corals are considered habitat-forming species because of its structural role in providing three-dimensional properties to benthic habitats (Miller et al. 2018). They serve as shelter to several other species and enhance ecosystem services, including fishery productivity, sediment retention, and protection from physical disturbances (Mumby et al. 2004, Worms et al. 2006, Fourqurean et al. 2012, Serrano et al. 2017, Hoegh-Guldberg et al. 2015, Harris et al. 2018). Most habitat-forming species are slow-growing plants and sessile invertebrates that display low reproduction success and high longevity (Garrabou et al. 2002; Linares et al. 2007; Teixidó et al. 2011; Arnaud-Haond et al. 2012, Kersting et al. 2014). Organisms with such slow life-history strategies may be especially vulnerable to the cumulative effects of multiple stressors, including direct and indirect effects of overfishing, coastal pollution, and global change (Hughes & Tanner 2000, Carpenter et al. 2008, Linares & Doak 2010, Short et al. 2011).

Local and global threats to benthic ecosystems

Overexploitation

Humans influence all oceans (Halpern et al. 2008). Overexploitation of marine resources has been an historical major stressor to marine food webs and still impacts a large proportion of the global oceans (Kroodsma et al. 2018). According to FAO estimates, more than half of fish stocks are fully exploited and another third are over-exploited, or depleted (FAO 2014). Besides reported captures, widespread illegal, unreported and unregulated fishing further deplete fish and invertebrates' populations (Pauly et al. 2005, Pauly & Zeller 2016). Benthic habitats are especially vulnerable to destructive fishing practices such as bottom trawling, which can cause major and long-lasting damage to sessile communities (Hiddink et al. 2017).

Coastal ecosystems are further impacted by several local and regional stressors including pollution, coastal development, and global change (Halpern et al. 2008, 2015). Today, the cumulative effects of multiple threats have driven compositional changes, local extinctions, and wholesale destruction of many benthic ecosystems (Jackson et al. 2001, Airoldi & Beck 2007).

Global Change

Since the industrial revolution, the addition of large amounts of CO₂ and other greenhouse gases to the atmosphere is changing the Earth climatic system. The negative consequences of rising ocean temperatures are increasingly notorious at all levels of biological organization – from genes to ecosystems (Scheffers et al. 2016).

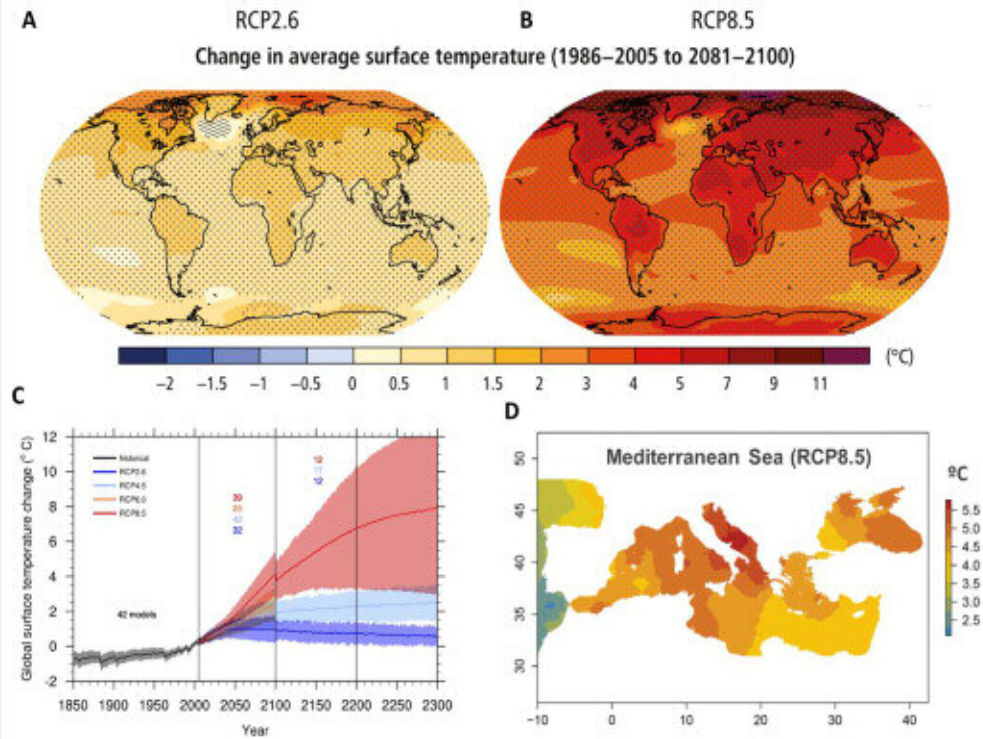


Figure 1. Projections on changes in average temperatures for the end of the 21st century according to IPCC5 scenarios: globally (a, b,c) and for the Mediterranean Sea (d).

Among many drivers linked to ongoing global change, steady increases in sea water temperature and recurrent heat-waves currently imperil structure, function and resilience of world's marine ecosystems (Hoegh-Guldberg & Bruno 2010, Ainsworth et al. 2016, Hughes et al. 2017b).

Widespread changes in the distribution and dynamics of marine organisms have been documented during past decades (Perry et al. 2005, Pinsky et al.

2013), and there is strong evidence that mobile organisms such as fish and invertebrates are tracking temperature trends over broad spatial scales (Pinsky et al. 2013, Montero-Serra et al. 2015). However, ecological responses to warming can be complex, and forecasting potential long-term consequences for habitat-forming species will require a deep understanding on how species-specific life-history, habitat requirements and community interactions shape their

capacity to tolerate and adapt to new thermal environments (Sunday et al. 2015, Rutteford et al. 2015, Kortsch et al. 2015). For instance, risks associated to ocean warming can be particularly high for sessile species with low larval dispersal and long-generation times because they cannot track suitable thermal conditions and thus may experience local population declines and local extinctions. Indeed, during the past three decades, large-scale mass bleaching and mortality events driven by heat waves have impacted iconic ecosystems such as tropical coral reefs (Graham et al. 2008, Hughes et al. 2016, 2017a, 2018); and temperate rocky reefs (Cerrano et al. 2001, Garrabou et al. 2009, Wernberg et al. 2016). These large-scale mortality events had notorious effects at the population and community level (Graham et al. 2016, Wernberg et al. 2016, Hughes et al. 2018a). Yet, for most species, the long-term consequences of recurrent heat-waves remain poorly understood. Worryingly,

field data over large spatial scales suggest that these mortality events have already increased in their frequency and intensity during the past three decades (Hughes et al. 2018b), and there is strong evidence that they will maintain this trend (Oliver et al. 2018, Frölicher & Laufkötter 2018).

Multiple threats to Mediterranean benthic ecosystems

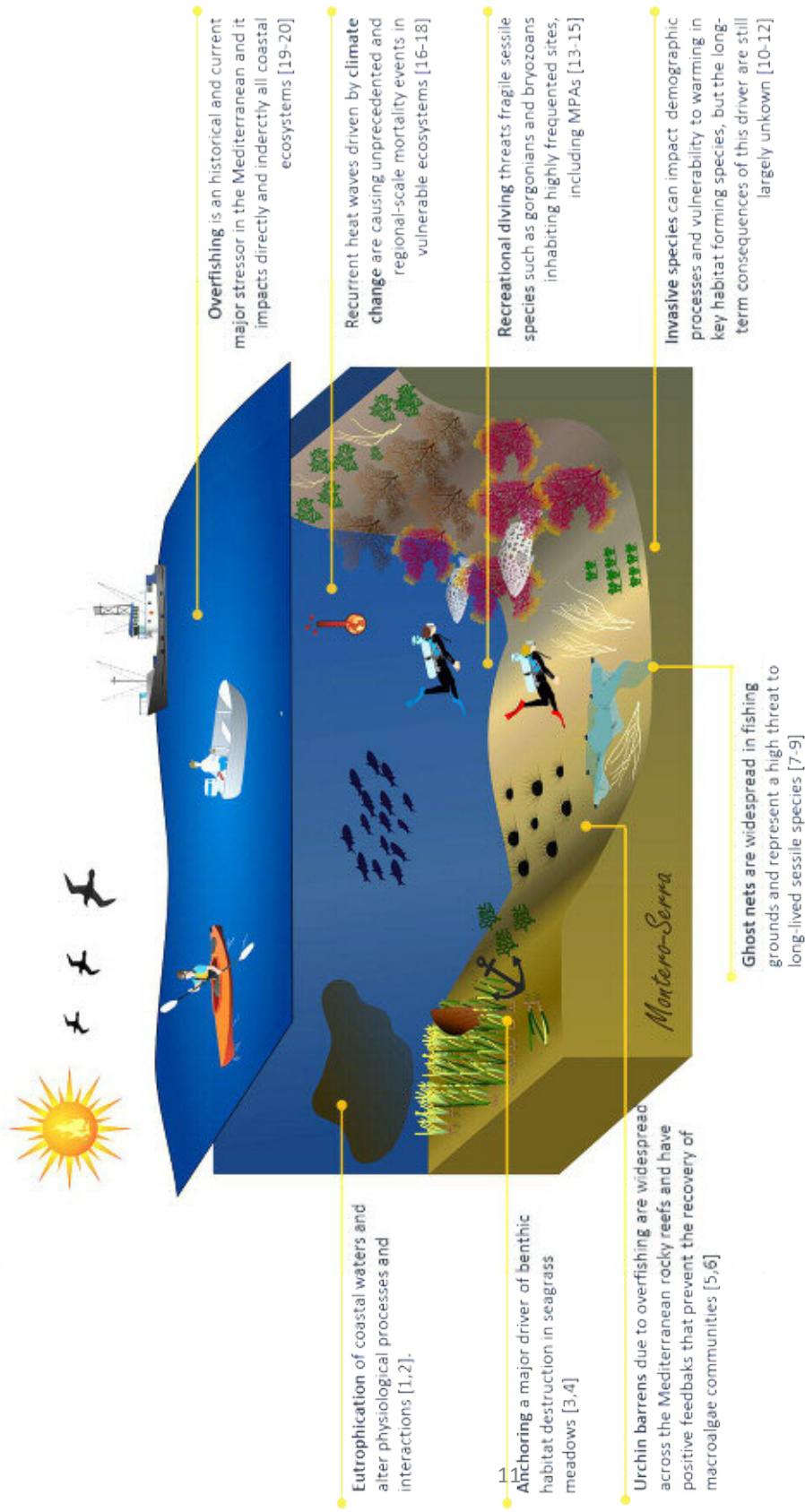


Figure 2. Conceptual example of a Mediterranean benthic ecosystem subjected to multiple local and global scale stressors. Original diagram that includes symbols from IAN Library of Symbols, University of Maryland. References are: [1] Delgado et al. 1999, [2] De'ath & Fabricius 2010, [3] Francour et al. 1999, [4] Montefalcone et al. 2008, [5] Sala et al. 2012, [6] Ling et al. 2015, [7] Ferrigno et al. 2017, [8] Bo et al. 2014, [9] Angiolillo et al. 2015, [10] Cebrian et al. 2012, [11] Kersting et al. 2014, [12] Kersting et al. 2014, [13] Garrabou et al. 1998, [14] Sala et al. 1996, [15] Barker & Roberts 2004, [16] Garrabou et al. 2009, [17] Cerrano et al. 2001, [18] Coa et al. 2009, [19] Coll et al. 2010, [20] Lejeusne et al. 2010.

Local management and resilience of benthic ecosystems: the role of marine protected areas

To cope with escalating effects of multiple stressors to marine ecosystems, conservation efforts at global and local scales are urgently needed (Kennedy et al. 2013). Locally, fishery regulations and marine protected areas (MPAs) can help to reduce or remove threats and are at the forefront of global biodiversity conservation strategies (Halpern 2003, Edgar et al. 2014).

MPAs are spatially explicit units where human uses of the ocean are regulated. This popular term is, however, very vague. Through this thesis, we use the term MPA to define areas where fishing is totally banned (i. e. no-take areas). However, artisanal fisheries could also be allowed in other types of MPA that either regulate specific gears or temporarily closures. In addition, several non-extractive human uses can also be managed within an MPA, including recreational activities such as scuba diving. Figure 3 represents a conceptual model of an MPA that has a gradient of protection levels, from

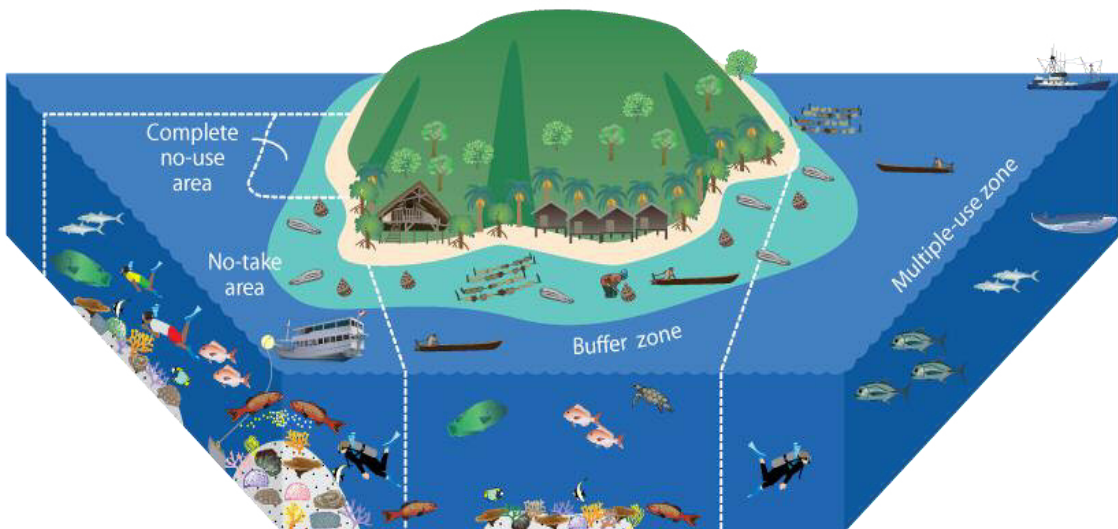


Figure 3. Schematic representation of a coastal marine protected area with different levels of protection. Source: IAN Symbols Library, University of Maryland.

buffering zones that restrict industrial fishing, to no-take areas that ban any type of extractive activity, to complete non-use zone or integral protection areas that prohibit any extractive or recreational activity.

MPAs were first used as a management tool to enhance fisheries productivity and ensure the long-term viability of fished stocks. Ecologists started recognizing community-level responses of fish assemblages to MPA implementations, either via direct trophic interactions or indirect cascading effects. In the 70's, James Estes and colleagues studied the dynamics of Eastern Pacific coastal ecosystems and demonstrated how protection lead to the recovery of sea otter populations and this cascaded into lower trophic levels, with a broad-scale recovery of kelp forests (Estes & Duggings 1995). Since then, ecosystem-level consequences of local protection have been observed in several tropical and temperate regions across the world. For instance, in tropical coral reefs, well enforced MPAs can rebuild parrotfish populations, enhance grazing intensity,

and modulate coral-macroalgae interactions with potential implications for coral reef recruitment and resilience (Mumby et al. 2005, Hughes et al. 2007).

Based on these appealing experiences, MPAs have become a key global conservation tool, with most nations agreeing to commitments made at the World Summit on Sustainable Development in 2002, the fifth World Parks Congress in 2003, and the eighth Conference of the Parties (COP 8) in 2006 to set aside 10–30% of their waters in MPAs by 2012 (Sala et al. 2018).

Today, the perception of MPAs as a panacea that can save vulnerable ecosystems from any stressor is widely embraced. However, empirical evidences that also underscored some limitations of relying on MPAs to protect habitat-forming marine species (Sala et al. 2012, Christiansen et al. 2014, Toth et al. 2014).

In this context, ecologists should provide sound evidence-based frameworks to understand and predict the positive effects of MPAs, but also

to identify their potential limitations to protecting marine biodiversity from stressors that cannot be managed locally. There are several important aspects to consider and unknowns that should be scientifically addressed to advance towards the design of sound scientific and effective management strategies.

A well-developed theory based on empirical studies states that MPAs are most effective at rebuilding depleted populations of targeted fish when all fishing is prohibited, enforcement is strong, and marine reserves are large, old, and isolated (Halpern 2003, Edgar et al. 2014, Gil et al. 2017). For habitat-forming species that are targeted for exploitation, such as the case of precious corals, oysters, clams or sponges, the same principles of design and enforcement of MPAs may apply (Michelli et al. 2010), although in some cases the time periods needed can be of the order of decades (Linares et al. 2010, 2012, Keiser et al. 2018).

The catastrophic consequences of ongoing climate change push for a critical debate in marine conservation to assess when MPAs can enhance

resilience on structural benthic species. MPAs can remove several local threats and, in some cases, we may expect some buffering capacity via indirect mechanisms, including changes in population size-structures (Linares et al. 2010), portfolio effects (Bernhard & Leslie 2013), and competition release. For instance, in tropical coral reefs, enhanced herbivory within reserves plays a crucial role in facilitating coral recruitment and recovery (Mumby et al. 2007). However, in many cases, these mechanisms are not clear and current evidences of large scale bleaching events suggest that MPAs may not provide increased resistance in affected benthic communities (Hughes et al. 2017a). Under which circumstances protected reefs can show increased resistance and recovery is a major research question that needs to be addressed.

How does this thesis contribute to current challenges? Insights from quantitative tools and life-history theory

Principles of life-history theory

Life-history strategies define lifetime patterns of energy allocation to growth, reproduction and maintenance of individuals. These traits can differ in their relative magnitude and timing, and their combination determines the overall fitness at the population level, which is subjected to natural selection and drives evolution (Stearns 1977, 1989). Life history theory studies mechanisms underlying the great diversity of strategies coexisting on Earth. Despite a longstanding interest from ecologists and evolutionary

biologists to disentangle what drives life-history variation, some fundamental questions remain partially unresolved. In addition, the potential for life-history principles to address marine conservation issues is generally overlooked.

Besides its crucial role in linking individual traits to population-level fitness and providing insights into evolutionary process, life-history theory can play a crucial role as a framework to address conservation challenges (Griffith et al. 2016). In particular, the demographic differences that arise from life history tradeoffs can shape responses to conservation efforts, including the effectiveness of management strategies (Morris & Doak 2002), likelihood of biological

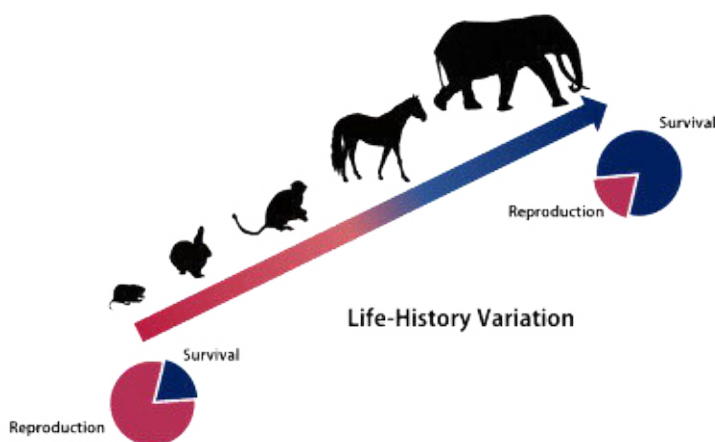


Figure 4. Example of life-history variation along the *fast-slow continuum* in mammal species.

invasions (Kerr et al. 2016), and the responses of species to global change (Jiguet et al. 2007, Linares and Doak 2010, Salguero-Gomez et al. 2012).

In the present thesis, we provide new insights into the role of the environment in life-history evolution and explore potential applications of life-history theory to inform conservation issues, including assessments of the effectiveness of marine protected areas, ecological restoration actions, and anticipating recovery periods.

Quantitative Ecology

Quantitative ecology is a growing field that focuses on statistical and computational methods for addressing questions in fundamental and applied ecology. It differs from theoretical ecology which focuses on simpler mechanistic models and aims to find general rules across model biological systems. Rather, quantitative ecology tools are fitted to empirical data and often used to address specific environmental issues. Common examples of their application include modeling dynamics of natural

populations, forecasting the potential impacts climatic change, and predicting the spread of invasive species. Will an exotic species thrive in a new territory? What are the best management options to eradicate invasive species or to facilitate population recovery? Those are examples of important questions that population modeling can help to answer.

Population models

Population models are flexible quantitative tools with multiple relevant applications (Griffith et al. 2016). The basic rationale behind them is that populations experience a net growth resulting from a balance between the entrance of new individuals to the population by reproduction and immigration, and the reduction of individuals by death or emigration. However, often individuals show divergent traits and populations are best modeled when individuals are classified into age, size or physiological state. A structured population model tracks the dynamics not only of total population size, but also the distribution of the variables that differentiate individuals. The choice of

the suitable stages depends on the type of organism and should be informed by biological knowledge. In corals, for instance, stages are generally best determined by size rather than age, because they often exhibit size-dependent fecundity, growth and mortality rates (Hall & Hughes 1996, Tsounis et al. 2006), but also due to the challenges associated to age-determination in clonal organisms (Hughes & Jackson 1990, Hughes 1984).

Matrix population models

Matrix population models are a specific type of population model that uses transition matrices, which are a form of algebraic representation for synthesizing a larger number of often repetitious algebraic computations. The matrix describes all the ways that an individual in one stage class at present time can contribute to another stage class in the next year (or any time step defined in the model). Matrix models are very popular in ecology because they are relatively simple, easy to construct and provide useful information. For instance, elasticity analysis of matrix models is an

important tool in conservation biology, allowing quantifying the relative contribution of each vital rate to the overall population growth, and this can be highly informative to assess conservation priorities (Morris & Doak 2002).

The main intrinsic limitation of the matrix models comes from discretizing individual states into a finite list of options. All individuals within a given stage therefore are assumed to be identical, in terms of their possible fates and the odds for each possible fate. Sometimes this is reasonable but for many sessile species can include arbitrary decisions. For instance, for organisms often described according to a continuous variable such as size, like plants or corals, defining discrete classes can be arbitrary and potentially misleading (Hughes et al. 1984, Merow et al. 2014).

Integral Projection Models

Dealing with this issue requires models that can accommodate a continuous individual-level state variable (Fig.5). Easterling (2000) show how to construct such an Integral Projection

Model (IPM), using continuous stage classes and discrete time, and provides sensitivity and elasticity formulas analogous to those for matrix models. While the change to a continuous structuring variable has very little impact on the behavior of the model, IPMs retain much of the analytical machinery that makes the matrix model appealing (Ellner & Rees 2005). In this thesis, we fitted integral projection

models to long-term demographic data on the red coral *Corallium rubrum* to describe his life-history and vulnerability to local and global stressors, and provide novel potential applications of these models to anticipate expected recovery times after fishing and active restoration actions.

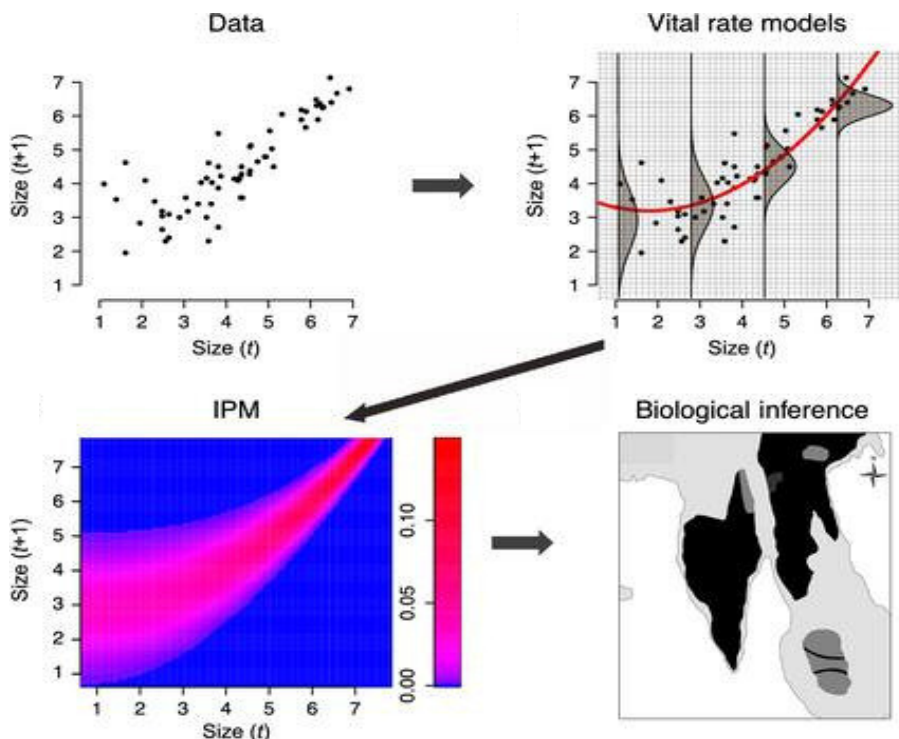


Figure 5. Integral projection models allow integrating continuous size-based individual data to perform a way arrange of biological inference (Merow et al. 2014).

Study system and model species

Mediterranean Sea

The Mediterranean Sea hosts high levels of biodiversity and sustain coastal economies providing ecosystem services to 143 million of people from 21 countries (Coll et al. 2010, NOAA 2015). Mediterranean marine habitats have been historically damaged by the cumulative action of multiple stressors such as overfishing, habitat destruction, pollution and invasive species (Lejeusne et al. 2010, Claudet & Fraschetti 2010). In addition, temperate regions are global hotspots of present and future climate change (Oliver et al.

2018). Due to its geographic position and semi-enclosed configuration, the Mediterranean is warming at much faster rates than the observed mean trends for the global oceans (Vargas-Yanez et al. 2008) and the occurrence of hot extremes has increased by more than two-fold (Oliver et al. 2017). These heat waves had impacted Mediterranean benthic communities at broad-scales, triggering widespread mass mortality events across vulnerable ecosystems such as seagrass meadows or coralligenous habitats (Garrabou et al. 2009).

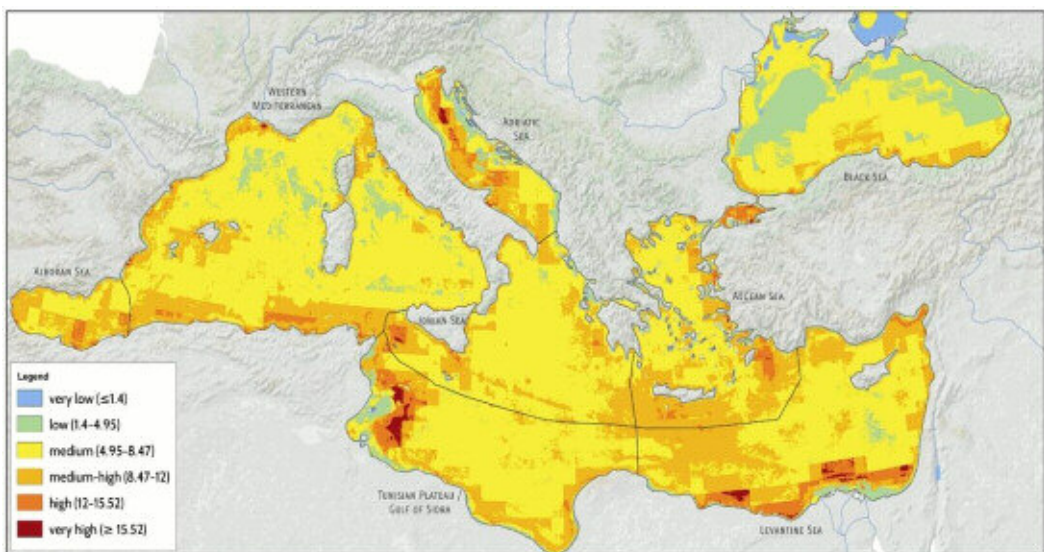


Figure 6. Cumulative stressors on Mediterranean marine ecosystems (Micheli et al. 2013)

Mediterranean coralligenous habitats

Coralligenous habitats are bioconstructions formed by the accumulation of calcareous encrusting algae growing at low irradiance levels. They are highly diverse assemblages that harbor more than 1600 species (Ballesteros 2006). These communities are dominated by long-lived algae and sessile invertebrates, which exhibit low dynamics and belong to a variety of taxonomic groups including anthozoans, porifera, bryozoans and tunicates (Teixidó et al. 2011). Coralligenous habitats can be found in hard bottoms across all Mediterranean coasts and have a bathymetrical distribution ranging from 20 to 120 m depth depending on the local environmental variables (Ballesteros 2006, Martin et al. 2014).

Mediterranean anthozoans are among the most charismatic and ecological important species found in coralligenous habitats. They form relatively dense populations that provide structural complexity to benthic habitats and enhance

biodiversity (Ponti et al. 2014, 2016). In this dissertation, we will focus on the Mediterranean red coral *Corallium rubrum*, which is a long-lived octocoral with an important cultural and ecological role in the Mediterranean Sea. The red coral as well as other coralligenous species are affected by several disturbances affect the Mediterranean red coral populations. Destructive fishing practices, ghost fishing, anchoring, over-frequent recreational diving, mucilaginous algal aggregates and algal invasions. More recently, several mass mortalities related to positive seawater temperature anomalies have been reported across NW Mediterranean populations (Cerrano et al. 2001, Garrabou et al. 2001, 2009).

Mediterranean red coral

The red coral, *Corallium rubrum* (Linnaeus 1758) is an emblematic precious coral endemic to the Mediterranean and adjacent Atlantic waters. Precious corals are cnidarian of the anthozoan group (animals with plant-like shape) that build an internal hard carbonate-skeleton which is very appreciated for jewelry. They are completely azooxanthellate and mostly feed on detrital organic matter and zooplankton (Tsounis et al. 2006, Picciano & Ferrier-Pagès 2007).

Red coral populations are patchily distributed in crevices, overhangs or cave entrances (Riedl 1984, Fig. 7). The bathymetrical distribution varies considerably in different areas, from shallow water populations formed in some nutrient-rich locations in the NW Mediterranean, up to a maximum of 1000m depth, as it has been recently discovered in Maltese waters (Carpine & Grasshoff 1975, Riedl 1984, Rossi et al. 2008, Costantini et al. 2010, Knittweis et al. 2016).

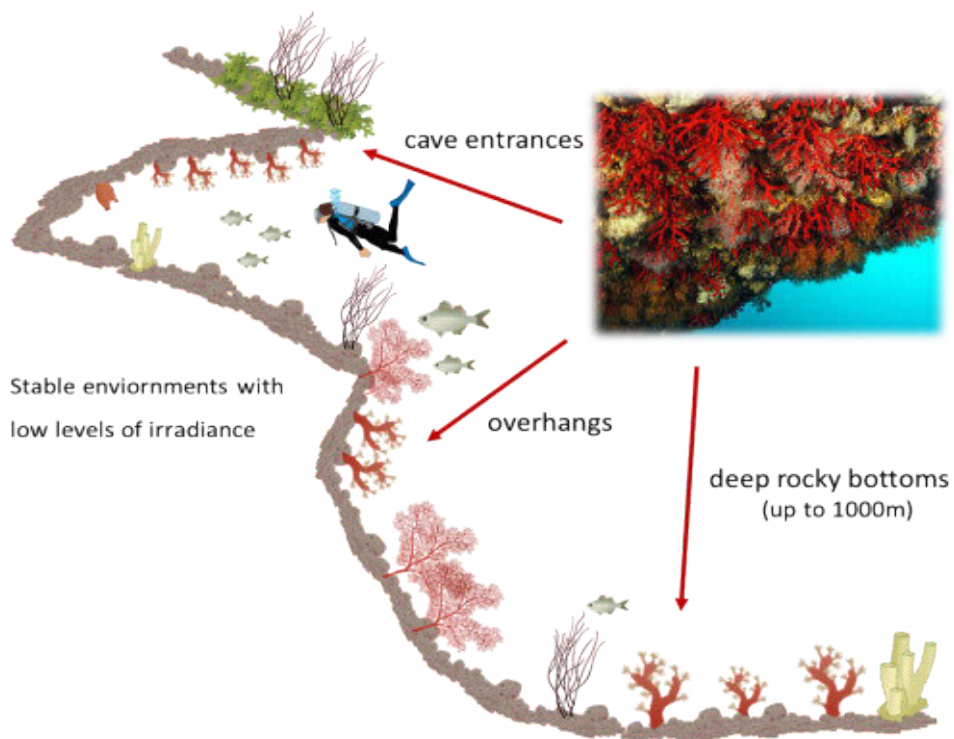


Figure 7. Habitats and bathymetric distribution of the Mediterranean red coral.

The red coral is a gonochoric internal brooder. Females brood larvae internally and then release them from late June and throughout July (Tsounis et al. 2006, Torrents & Garrabou 2011). *C. rubrum* reaches sexual maturity at relatively small sizes (about 2 mm in diameter and 3 cm in height) which corresponds to ages between 7 and 10 years old (Torrents et al. 2005, Gallmetzer et al. 2010). As commonly observed for marine sessile invertebrates, red coral colony sizes are correlated to reproductive outputs: the number of polyps per colony increases exponentially with size and large colonies also show higher proportion of gravid polyps (Torrents et al. 2005, Tsounis et al. 2006).

Population genetic studies revealed a very restricted gene flow (effective dispersal was estimated in 22 and 32 cm) (Ledoux et al. 2010a). Thus, red coral populations are mainly closed at ecological timescales and recovery from larvae coming from external sources is limited (Costantini et al. 2007, Ledoux et al., 2010a,b).

Due to historical overexploitation and present widespread unsustainable fishing practices, most red coral shallow populations are depleted and can be considered functionally impaired or even ecologically extinct (Bruckner 2009, Garrabou et al. 2017a).

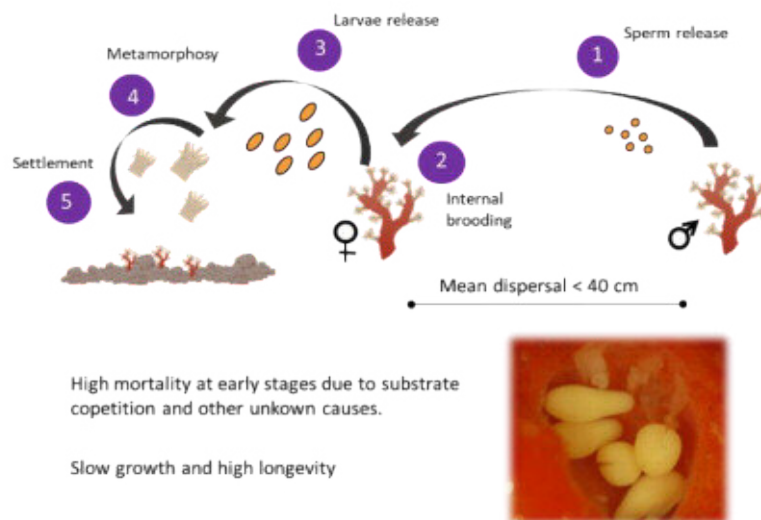


Figure 8. Life-cycle and reproduction in the Mediterranean red coral.

Aims and outline of the thesis

The present Doctoral Dissertation aims to implement cutting-edge modeling tools to address several issues in fundamental and applied ecology. In particular, we combine population and spatial models, long-term datasets, and global meta-analyses to unravel drivers in life-history evolution and their demographic consequences. Focusing on the long-lived red coral, we explore demographic consequences of harvesting and population recovery mechanisms, the long-term consequences of recurrent climatic disturbances and assess the effectiveness of local management tools such as marine protect areas in the face of local and global threats. We then investigate how life-history theory can shape the outcomes of active restoration efforts, and show how population models can be used to anticipate a crucial aspect of conservation actions such as the expected recovery times. Finally, we develop novel spatial modeling approaches to explore how depth will mediate the expected impacts of

present and future climate change in two main gorgonian species across the Mediterranean Sea. This thesis provides a broad overview on the demographic drivers of resilience for the Mediterranean red coral and other long-lived invertebrates to local and global stressors and shows how quantitative tools can be used assess the effectiveness of current marine conservation efforts in a changing environment.

The present Doctoral Dissertation has been conceived as a whole; however, the topics have been structured in five chapters intended for publication in international peer-reviewed scientific journals. Chapters 1, 2, and 4 are already published, but the format has been unified here to preserve the consistency of the present document. All figures, tables and text have been preserved as appear in the published papers. Cited literature has been unified and appears at the end of the document.

Chapter I: Strong linkages between depth, longevity, and demographic stability across marine sessile species

A comparative analysis across more than 200 marine sessile species revealed strong linkages between depth occurrence, longevity and demographic stability. These results demonstrated a fundamental role of environmental gradients in shaping the evolution of life-history strategies in the ocean. On the other hand, we showed that high longevity lead to dampened population dynamics. Focusing on the Mediterranean red coral, *Corallium rubrum*, which is an emblematic and highly overfished precious coral, we quantified the extremely slow life-history strategy of this species and the high vulnerability to external mortality sources of these strategies.

Chapter II: Harvesting effects, recovery patterns and management strategies for a long-lived and structural precious coral

Like other precious corals worldwide, the red coral *Corallium rubrum* has been historically overfished and its conservation is currently a worldwide

concern. However, the processes underlying their recovery after fishing remain poorly understood. We examined harvesting effects and recovery mechanisms on red coral populations and revealed high impacts with large decreases in biomass and strong size-class distribution shifts towards populations dominated by small colonies. The observed general pattern of low recruitment and high mortality of new recruits demonstrated limited effects of sexual reproduction on population recovery. Adversely, low mortality of partially harvested adults and a large proportion of colonies showing new branches highlighted the importance of re-growth in the recovery process. The demographic projections obtained through stochastic models confirmed that the recovery rates of *C. rubrum* can be strongly modulated depending on harvesting procedures.

Chapter III: Marine protected areas enhance structural complexity but do not buffer the detrimental consequences of climate change for an overexploited precious coral

We used long-term data on multiple red coral populations and Integral Projection Models to show that local management and warming-induced mass mortality events have different demographic consequences. Warming-driven mass mortality events had detrimental effects on the viability of red coral populations, causing long-term declines and leading to a potential local extinction. On the other hand, removal of local stressors through MPAs strongly enhanced the structural complexity of protected coral populations. Contrary to our expectations, however, local management had little effect on the long-term viability and associated extinction risk of coral populations. Simulations combining the effects of both global warming and harvesting under multiple climatic scenarios, revealed that MPAs have a weak buffering effect to extreme climatic impacts.

Chapter IV: Accounting for life-history strategies and timescales in marine restoration

Understanding the drivers of restoration success is a central issue for marine conservation. Here, we explored the role of life-history strategies of sessile marine species in shaping restoration outcomes and their associated timescales. A transplantation experiment for the red coral *C. rubrum* was highly successful over a relatively short term due to high survival and reproductive potential of the transplanted colonies. However, demographic projections predict that from 30 to 40 years may be required for fully functional *C. rubrum* populations to develop. More broadly, a comprehensive meta-analysis revealed a negative correlation between survival after transplanting and growth rates among sessile species. As a result, simulated dynamics for a range of marine sessile invertebrates predict that minimum recovery times are positively associated with survival rates. These results demonstrate a tradeoff between initial transplantation efforts and the speed of recovery.

Chapter V: Predicting depth-mediated impacts of ocean warming at the end 21st century in Mediterranean coastal ecosystems

Understanding the effects of global warming on the biogeography and vertical distribution of marine organisms remains challenging. Here, using a 3D ocean model, we show that present and future climate shapes the bathymetrical distribution of two dominant gorgonians of the Mediterranean Sea: *Paramuricea clavata* and *Corallium rubrum*. Under present conditions, local temperatures are negatively correlated to the upper distribution ranges of both species. In the future, using Bioclimatic Envelope Models, we predict a large decrease in habitat suitability based on 21st century climate change scenarios. The frequency and intensity of climatic impacts, however, will strongly depend on depth, basin (Western vs. Eastern Mediterranean), and emissions scenario. Overall, gorgonian populations will be highly impacted at shallow depths from 5 to 25m, but there will be regional differences: in the warmer Eastern Mediterranean Sea, the

impacts will be extensive and reach deeper habitats; and in the generally colder Western basin, there will be heterogeneous patterns of warming across space and depth.

CHAPTER I

Strong linkages between depth, longevity and demographic stability across marine sessile species

Ignasi Montero-Serra, Cristina Linares, Daniel F. Doak,
Jean-Baptiste Ledoux and Joaquim Garrabou

INTRODUCTION

From sequoias to small desert shrubs to naked mole rats, the study of species with unusually long lifespans has long fascinated evolutionary ecologists (Buffenstein 2005, Baudisch et al. 2013, Jones et al. 2014, Colchero et al. 2017). Understanding the conditions where extreme life histories dominate and the potential consequences of these singular strategies for the dynamics of natural populations is central to ecology and evolution, and may also have important implications for conservation biology (Griffith et al. 2016). However, to date ecologists have only found weak correlations between extrinsic factors and longevity, and we still have relatively few full demographic descriptions for extremely long-lived species.

In terrestrial landscapes, high longevity is common in relatively inhospitable environments characterized by low availability of resources, such as desert and alpine ecosystems (Vasek 1980, Forbis &

Doak 2004, Garcia et al. 2011). However, some species of extreme longevity inhabit highly productive habitats (e.g, the coastal redwood), and climate or habitat factors generally provide only weak predictions of life history traits (Stearns 1977, McIntyre et al. 1995, Morrison & Hero 2003, Moles et al. 2007). The marine realm is perhaps a more promising setting to examine the external factors driving life history variation than are terrestrial habitats, due to the set of physical-chemical properties that lead to strong and predictable depth-related gradients at smaller spatial scales than are observed in other systems (Garrabou et al. 2002). The main abiotic factors controlling bottom-up processes in marine communities (light, food availability, temperature, and disturbance intensity) all generally decrease with depth (e.g., Sanders 1968, Barnes & Hughes 1999, Smith & Brown 2002); these factors are also often invoked as likely selective agents favoring short or long lifespans (Munch & Salinas 2009). If environment plays a role in shaping the evolution of longevity,

we therefore expect an increase of lifespan along depth gradients. Yet, quantitative analyses addressing this hypothesis are scarce.

In this study, we first explore longevity patterns of marine habitat-forming species, asking how predictable longevity is across depths and whether there are consistent consequences of individual longevity for population dynamics. We discuss these results in the light of similar analyses for terrestrial plant species. Next, we conduct a focused analysis of the iconic Mediterranean red coral, examining aspects of population biology of an extremely long-lived species that may be general for high-longevity species. Our findings represent an important step towards a better understanding on the role of environmental conditions as drivers of the evolution and maintenance of longevity and have implications for the conservation of dominant structural species found at deep habitats, where demographic data are scarce.

METHODS

Review of demographic studies

Life history strategies are complex combinations of multiple traits that describe the timing and magnitude of the species' reproduction, growth, and survival, which in turn generate life history metrics such as longevity, generation time, and net reproductive rate. For two reasons, we have focused here on longevity as a proxy for this broader diversity in life history strategies. First, several comparative demography studies have shown that longevity is highly correlated to many other demographic traits; in particular, longevity is positively correlated to age at maturity and generation time and negatively correlated to reproductive output and growth (Gaillard et al. 2016, Salguero-Gomez et al. 2016). Second, unlike many complicated life history traits that require detailed demographic study to estimate, longevity can be estimated in quite simple ways and is widely reported, making it

especially suited to broad-scale comparisons.

To construct our dataset, we searched the literature for studies reporting longevity estimates and demographic data by using several combinations of keywords. We restricted our analysis to marine sessile species and used the terms “marine sessile”, “coral”, “hexacoral”, “octocoral”, “macroalgae”, “seaweed”, “bivalve” and “bryozoan”. These were combined with either “demographic model” and “matrix model” or “longevity” and “lifespan” in the Web of Knowledge. We found 144 suitable studies reporting data for 241 species (Table S9). We include studies of polychaetes that form cemented tubes, and thus are effectively sessile, as well as studies of bivalves that are and are not strictly sessile.

Maximum longevity is estimated through different methodologies. Some studies directly report estimates of maximum observed lifespan using growth rings or

geochemical analysis or both. Others report standard demographic data in the form of a matrix model or annual survival rates of adult individuals or colonies. The latter were used to derive potential maximum lifespan based on demographic simulations using the same methodology as for the red coral (see Section on Demographic analysis below). When demographic data were based on non-annual transitions, annual lifespan was calculated as $L_{ann} = L * 12 / T$, where L_{ann} is the annual lifespan, L is the non-corrected lifespan value, and T is the period described by each transition in months.

As a proxy for habitat, we used depth of occurrence, quantified by the maximum depth reported for the species. While mean or median depth would be more informative, accurate information on these central depths is not available for a wide range of species. Maximum depth is more commonly reported and likely to be correlated with a species' characteristic depth range. Maximum depths were obtained by

searching in the literature including the keywords “depth”, “depth range” and “maximum depth” combined to the species scientific name (Table S9).

Demographic variability using matrix population models

To quantify temporal variance in population growth, we used stochastic demographic models based on at least 5-year study periods (4 annual transition matrices) for marine sessile species ($n = 9$) and terrestrial plants ($n = 25$) (Table S10). For each species, we simulated the fate of 1000 initial individuals for 1000 years, assuming equal probabilities for each reported annual transition, and computed changes in population size by dividing the total number of individuals at each time (t) by the previous number ($t-1$). We then log-transformed these realized annual lambda values and computed their variance. Two long-term demographic studies on bivalve species reported only observed annual lambdas, thus we computed

variance in population growth directly from these observed lambdas after a log-transformation.

Statistical analysis

We used Pearson and Spearman’s rank correlations and linear models to assess the relationships between depth and longevity as well as longevity and population growth variability. We also explored multiple combinations of potential predictors and their interactions by fitting a set of multiple linear models and using Akaike Information Criteria corrected for small sample sizes (AICc) to select the best models. First, a set of linear models was fitted to the whole longevity dataset to test for effects of depth while accounting for different methodological differences in longevity calculations and also to explore potential significant interactions between depth and taxonomic class. We also ran separate set of models to test for other taxonomic levels (Phylum or Order) had more explanatory power or shifted the results. A second set

of models was fit using only the species for which estimates were based on demographic models. For these species, we explored which longevity definition (see section on Demographic Analysis) had a better fit with maximum depth. Finally, to assess the effect of longevity on demographic variability, a third set of models was fit to sessile species for which four or more annual transition models were available, setting variance in population growth as a response variable and maximum longevity, habitat, and their interaction as potential predictors. In these and all other statistical tests, longevity was log-transformed for analysis to achieve more linear results.

Study Case: The Mediterranean red coral *Corallium rubrum*

Natural history

Precious corals are long-lived species found in several seas across the world. The red coral *Corallium rubrum* is a long-lived octocoral of the Mediterranean Sea. It is a depth-generalist species commonly

observed in the NW Mediterranean at shallower depths in overhangs and caves but its bathymetrical distribution can reach up to 1000m deep (Knittweis et al. 2016). Due to the high value of its carbonate skeleton for the ornamental jewelry industry, *C. rubrum* has been intensively harvested for millennia. Exploitation - both legal and illegal - as well as ongoing climate change are current threats to this species (Garrabou et al. 2001, 2009, 2017, Linares et al. 2012, Bramanti et al. 2013, Montero-Serra et al. 2015).

Study area and demographic monitoring

A total of 1144 coral colonies from eight *C. rubrum* populations along the North-Western Mediterranean region were individually monitored using photographic techniques (n = 30 photoquadrats of 20x20cm at each population) over periods ranging from 3 to 10 years between 2003 and 2011 (Table S1, see Montero-Serra et al. 2018 for detailed information).

Demographic Analysis

Based on the photographic time-series data, we could estimate size-dependent annual survival, fecundity and growth rates. We used a set of Generalized Mixed Effects Models (GLMMs) to describe annual survival and fecundity rates. Because the goal was to describe the general life-history of the red coral, population and year were included in the models as random factors. Growth rates are extremely low for this species (Garrabou & Harmelin 2002, Marschal et al. 2004), thus a non-linear growth/shrinkage model was fit to data of initial and final colony sizes measured for 247 colonies in two populations over periods of 7-8 years. Due to relatively rare ‘extreme shrinkage’ of some colonies, we divided the growth/shrinkage process into normal growth and extreme retrogression, each with their own mean change of size (see SI). The growth/shrinkage rates model and the fixed effects from the survival and reproduction GLMMs, which included colony size as a fixed

effect and population and year as random factors, were then used to construct a deterministic Integral Projection Model (hereafter IPM). We analyzed this model as a large matrix based on 80 colony size classes (see Montero-Serra et al. 2018, and Supplementary Methods for a full description on model construction).

Maximum potential lifespan was estimated by deterministic simulations using the demographic IPMs. Specifically, we simulated the fate of 100 individuals starting at the same stage and recording the number of years until 50% (mean lifespan) and 5% (maximum potential lifespan) of individuals were still alive. To explore the expected mean and maximum lifespan conditional on starting at different life-stages, we run three set of simulations: starting with 100 recruits, starting with 100 of the smallest size-class adults (first reproductive individuals, Torrents et al. 2005), and starting with 100 largest size-class adults. In all results reported in main text, we use

adult maximum longevity. To explore how robust these longevity metrics are to changes in the dimension of the transition matrix derived from the IPMs, we computed all these metrics using matrices constructed with 10 to 200 size classes.

To assess the elasticity of deterministic population growth to each vital rate, we performed a perturbation analysis of size-dependent survival, fecundity, and mean normal growth and mean extreme shrinkage, using an 80 size-class model. Each size-dependent vital rate was separately increased or decreased (by 1% and 5%). A mean demographic model was derived that included each perturbation and its deterministic population growth rate (dominant eigenvalue) was computed. Finally, the size dependent elasticity of each vital rate to lambda was calculated using:

— —

Where a is the vital rate, j the size-specific interval, λ the deterministic

population growth, and e_{aj} is the elasticity for vital rate a_j . Since underlying vital rates were perturbed separately (e.g, mean growth, mean extreme shrinkage, and survival), no compensation to control for shifting effects on other vital rates was required (see approach 4 in Griffith et al. 2017).

We calculated mean age-specific survival and fecundity rates by simulating a cohort of 1000 new recruits and computing at each time step the survival rate (N_t/N_{t-1}) and the fecundity (R_t/N_{t-1}) where N is the total adult abundance and R is the total number of recruits.

RESULTS

Depth, longevity and population stability

The comparison of maximum lifespan revealed a great diversity of life histories across marine sessile species (Fig. 1). Hydrozoans,

bryozoans, cirripeds and polychaetes displayed the shortest lifespans, ranging from weeks to several decades. These generally short-lived taxa have received significantly less attention in the literature compared to other major taxa such as bivalves and corals. The other groups had a

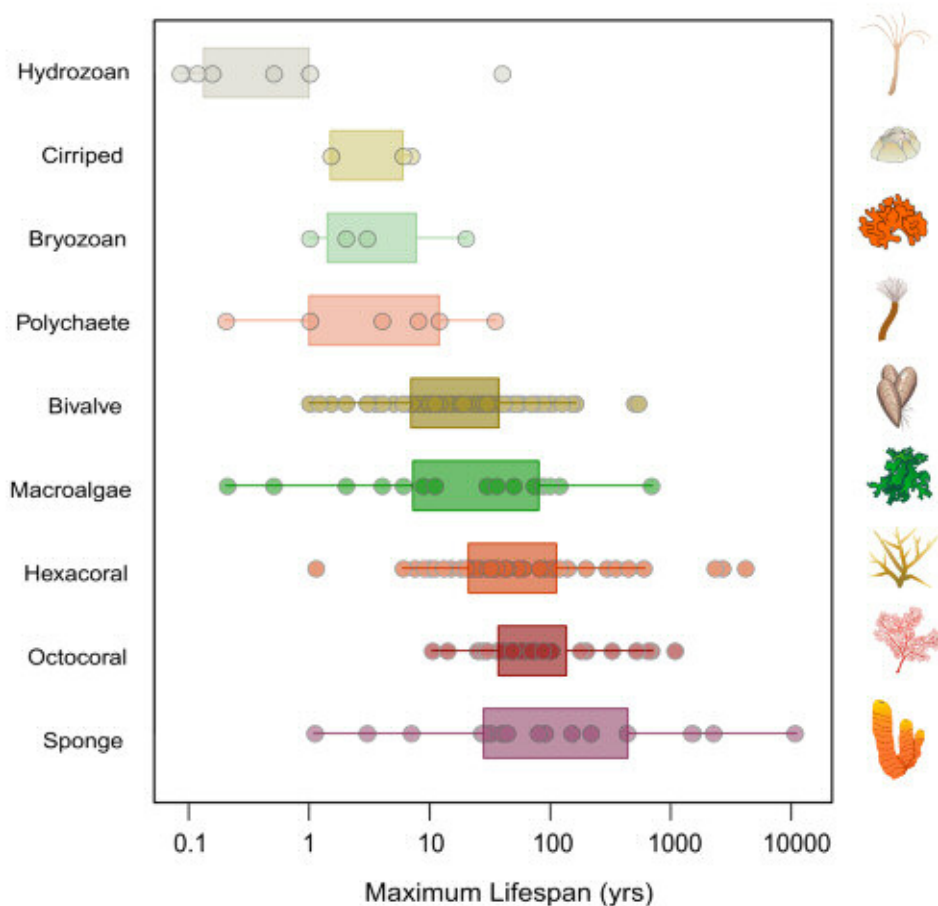


Figure 1. Longevity patterns across marine sessile species (n = 241). Data were obtained from the literature and correspond to the maximum potential lifespan reported for each species for the main marine sessile taxa. See Table S9 for specific species data.

great range of lifespans, up to hundreds of years for some calcareous macroalgae and bivalves and to thousands of years for cnidarians and sponges. Most macroalgae species ranged from less than one year to several decades, but an outlier with exceptional longevity is the Antarctic calcareous algae *Clathromorphum nereostratum*. Marine bivalve's lifespans ranged from less than 1 yr to more than 500 yrs for the antarctic scallop *Arctica islandica* and the giant deep-sea oyster *Neopycnodonte zibrowii*,

which is the longest-lived known non-clonal animal. Modular and structural taxa such octocorals, hexacorals and sponges were dominated by longer lived species. Extreme cases were the deep-sea coral *Gerardia sp.*, found in Hawaii, with an estimated age of 4700 yrs and the longest-lived known marine species, the sponge *Monorhaphis chuni*, observed at 1000m depth with an estimated age of 11000 yrs (Roark et al. 2009; Jochum et al. 2012).

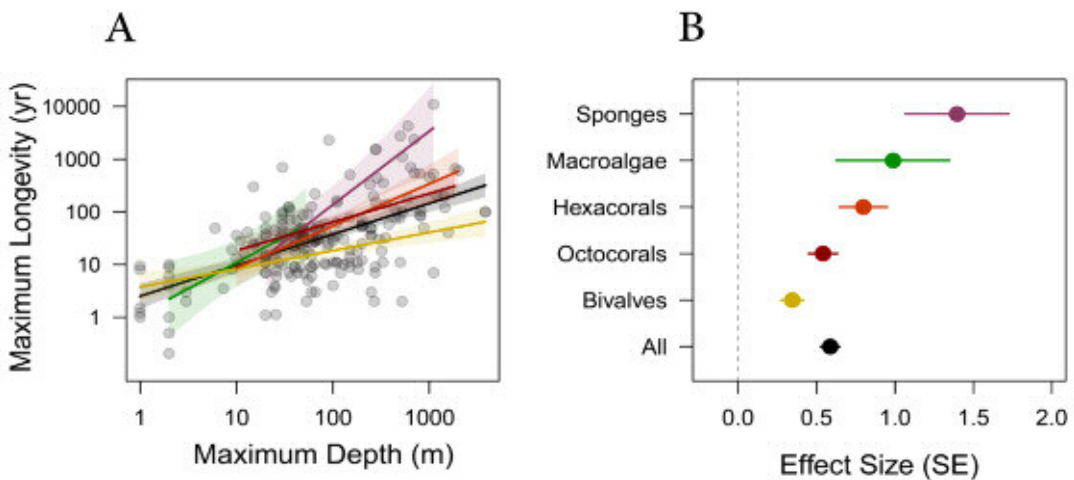


Figure 2. Relationship between maximum depth of occurrence and maximum longevity across marine sessile species. Colors correspond to the labels in panel B, and lines show the best fit linear model for each taxon. Black line shows the linear relationship for all taxa. Shaded areas represent standard errors. B. Slope corresponds to the estimated slope of maximum depth on maximum longevity for each taxon, with error bars showing ± 1 standard error.

We correlated $\log(\text{maximum lifespan})$ with $\log(\text{maximum depth})$ and found strong positive relationships in the five main taxa for which enough data were available to perform meaningful analyses: octocorals (Spearman's $\rho = 0.673$; $P < 0.001$; $n = 35$), hexacorals (Spearman's $\rho = 0.592$; $P < 0.001$; $n = 40$), sponges (Spearman's $\rho = 0.647$; $P < 0.001$; $n = 17$), macroalgae (Spearman's $\rho = 0.600$; $P = 0.023$; $n = 14$), and bivalves (Spearman's $\rho = 0.491$; $P < 0.001$; $n = 93$) (Fig. 2, Table S2). The positive relationship between depth and lifespan was consistent across different measures of longevity, although adult maximum longevity showed the strongest relationships with depth (Table S3). This relationship is also consistent when accounting for taxon or matrix dimension; the best supported model includes an interaction between depth and taxon, with significantly larger slopes for sponges ($P = 0.0131$) and macroalgae ($P = 0.0308$) than for the other taxa (Table S4) (Table S4; overall model statistics: $R^2 = 0.511$; $F = 18.12$; $df = 185$; $P <$

0.001). Use of other taxonomic levels does not change this qualitative result, and model fit is best for Class (Table S5). We also ran quantile regressions between $\log(\text{depth})$ and $\log(\text{longevity})$ to see if depth is a more powerful regulator of extreme life history values, but found no significant differences in the slope of the depth ~ longevity relationship for 10% up to 90% quantiles (Table S6).

To assess population consequences of longevity, we regressed variance in realized $\log(\lambda)$ on longevity across marine sessile invertebrates and terrestrial plants and found a similar, strongly negative relationship for both. Short-lived species displayed higher variability in population growth over time than did longer lived species ($n = 34$; Spearman $\rho = -0.73$; $P < 0.001$). The pattern was consistent in terrestrial plants ($n = 25$, Spearman $\rho = -0.68$; $P = 0.0002$) and marine sessile invertebrates ($n = 9$; Spearman $\rho = -0.67$, $P = 0.0214$) (Fig. 3, Table S7).

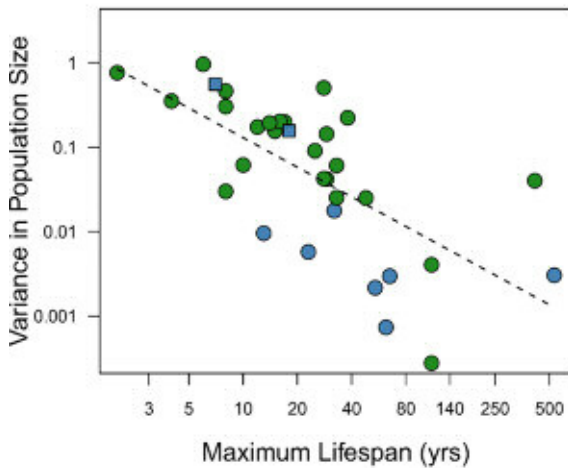


Figure 3. Relationship between temporal variability in population growth and maximum longevity in terrestrial plants (green, $n = 25$, Spearman $\rho = -0.68$; $P = 0.0002$) and marine sessile (blue, $n = 9$; Spearman $\rho = -0.67$, $P = 0.0214$) species, and overall ($n = 34$; Spearman $\rho = -0.73$; $P < 0.0001$). Squared blue dots correspond to two bivalve species whose variance was computed directly from observed annual lambdas.

The extreme life-history of *C. rubrum* and its population dynamics

The red coral *C. rubrum* displayed slow dynamics over the eight populations studied along the NW Mediterranean (Fig. S1), a pattern

driven by extremely high adult survival, ranging between 98.05% and 99.76% from the smallest to the largest reproductive *C. rubrum* colonies (Fig. 4a). Monitoring of post-recruitment survival revealed that relative to later survival rates, annual survival is low during the first years after recruitment with a mean of 69.5% (Fig. 4c). However, by the age of four, young colonies already displayed a high survival probability of 92.2%. Overall, mortality rates decreased rapidly during the early stages of the red coral lifetime, following a Type III survivorship curve, and then decreased steadily, more similar to Type II survivorship (Fig. 5a). Recruitment was very limited; from 246 quadrats (about 30 per population) individually monitored for recruitment at the eight studied populations, 83.7% showed no annual recruitment. The remaining 16.3% of quadrats were divided into 10.7% with only

a single recruit present, 5.1% with 2-44 recruits, and only 0.5 with five or more recruits (Fig. 4b, 4d). Mean annual per capita recruitment, weighted by the relative abundance of different sized individuals, was

0.077. This is considerably lower than the average per capita recruitment for a range of marine sessile invertebrates for which there standard demographic data was available ($n = 28$, Median = 0.547-

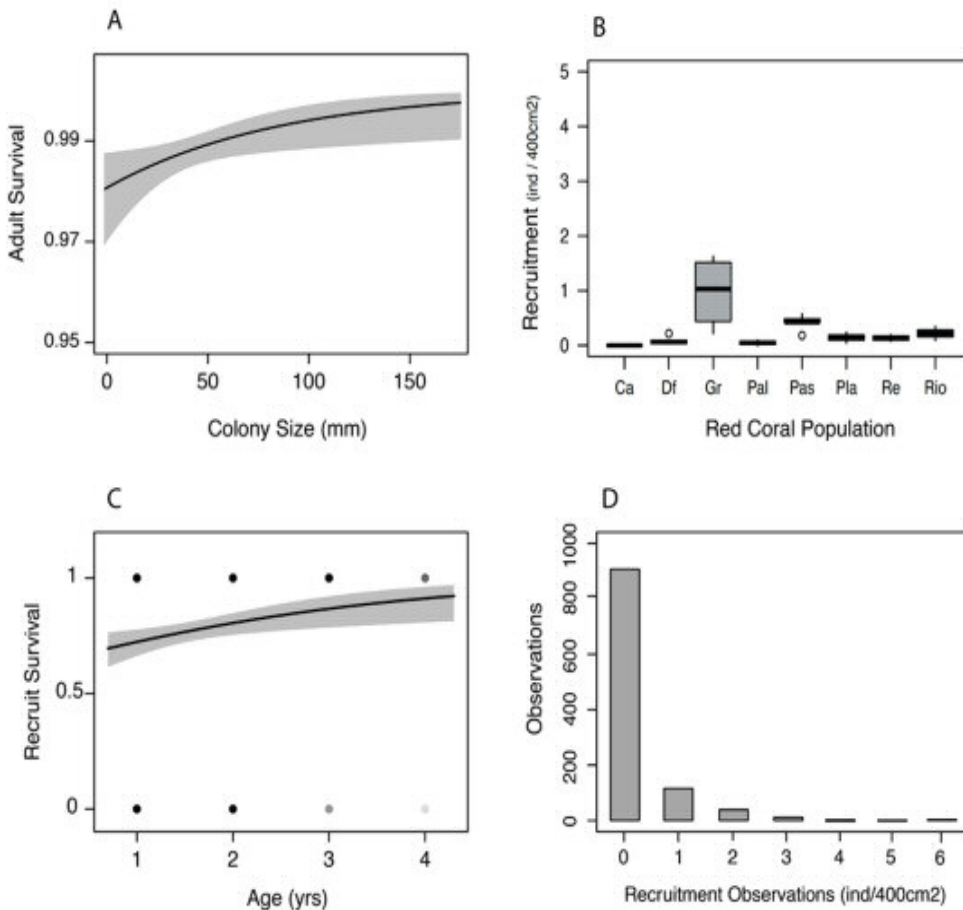


Figure 4. Long-term demographic traits of the red coral *Corallium rubrum*. A. Adult survival probability depending on colony size. The line represents a logistic Generalized Linear Mixed Effects Model including population and year as random effects and colony size as a fixed effect. B. Mean recruitment rates in eight *C. rubrum* populations. C. Age-dependent post-recruitment survival probability. D. Frequency of annual recruitment observations in eight *C. rubrum* populations. In A and C, grey areas show ± 1 standard error.

0.727, Mean = 5.14 ± 4.02 (SD), Table S8), although higher than the same estimate for 6 other species (Table S8).

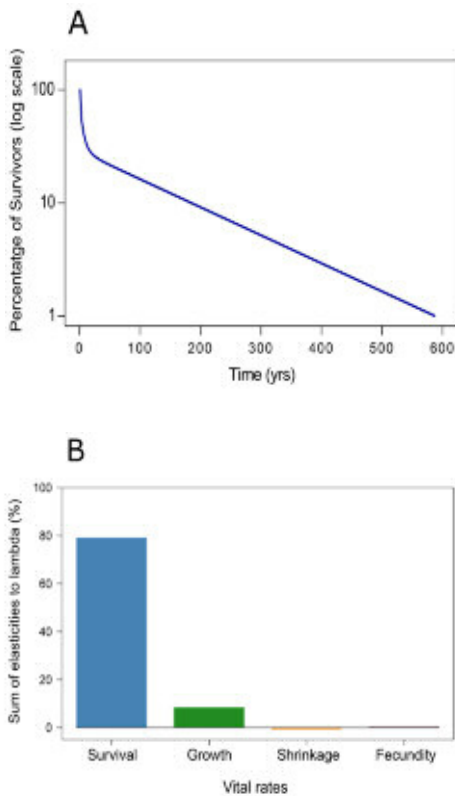


Figure 5. Demographic analyses for the red coral *Corallium rubrum*. A. Survivorship curve based on a deterministic simulation of 100 individuals. B. Patterns of elasticity to the asymptotic population growth rates of different vital rates, with each bar indicating elasticity summed across all size classes. Other vital rates have even lower elasticities (Fig. S4).

Demographic simulations revealed high maximum potential future longevity for *C. rubrum* colonies, ranging from 501 to 532 yrs for small and large adult respectively (Fig. 5a, S2). The mean expected lifespan was also relatively high for adult red coral colonies and ranged from 98 yrs for small adult colonies to 129 yrs for the larger red coral colonies. Our simulations also revealed potential artefacts due to matrix dimensionality. When using relatively small transition matrices (<60 size classes) simulations underestimated values of maximum and mean longevity, the values stabilized around 60 to 80 size-classes and were consistent thereafter (Fig. S2).

Perturbation analysis revealed a disproportionately large influence of survival rates on asymptotic population growth (Fig. 5b). Elasticity of lambda to fecundity, mean growth and mean extreme shrinkage, as well as all other vital rates, was markedly lower (Fig. S4). Model fitting of age or size dependent fecundity and survival

showed no evidence of senescence (Fig. S3).

DISCUSSION

Demography is a powerful tool to understand the evolution of life histories and has provided important insights into plant, mammal, and bird's strategies (Griffith et al. 2016, McDonald et al. 2017). However, a lack of long-term and broad spatial scale data on the dominant species of marine ecosystems, particularly those living at deeper depths, hinders our understanding of the processes shaping the dynamics of these threatened ecosystems (Donovaro et al. 2017). In this study, we analyzed the long-term dynamics of the long-lived red coral *Corallium rubrum* and published data for a broad range of marine sessile invertebrates and macroalgae. The results revealed strong positive relationships between depth, longevity and demographic stability, showing strong habitat-determined predictability of key demographic processes in marine ecosystems.

Marine sessile species have a wide range of maximum lifespans across taxa and habitats, ranging from weeks in some intertidal species to hundreds of years for the red coral *C. rubrum*, and several thousands of years for some deep-sea corals and sponges (Fig. 1). Our most striking finding was that maximum lifespan was strongly and positively correlated with a species' maximum depth occurrence in hexacorals, octacorals, sponges, and although weaker, also in bivalves (Fig. 2). These results demonstrate a strong role of habitat features at shaping the distribution of longevity patterns.

In the shallowest waters, short-lived species dominate (Fig. 2), perhaps due to high disturbance rates and competition in high-energy environments where fast-growing species have a distinct advantage. In contrast, species displaying diverse life-history strategies coexist at intermediate depths, including short-lived invertebrates that also occur intertidally and longer-lived massive corals, macroalgae and seagrasses

(Arnaud-Haond et al. 2012; Darling et al. 2012). The high energy availability due to constant solar irradiation may allow the presence of a large spectrum of life-history strategies that have diversified to occupy multiple niches, following analogous processes than those shaping tropical rainforest plant communities (Connell 1978). Finally, in deeper waters, environmental constraints seem to favor the dominance of slower life-histories and longer lifespans. Depth gradient determines changes in environmental factors such as light and food availability, temperature, and disturbance intensities. In turn, these factors are likely to control the evolution of lifespans through several mechanisms. First, deeper habitats are generally less productive, limiting food supply for consumers and favoring slow growth and longer lifespans (Larson 2001). Empirical studies exploring physiological tradeoffs have shown that the elevated metabolic costs of fast strategies can be detrimental in the absence of high availability of resources (Reznick et al. 2000).

Also, as light is strongly depth-dependent, organisms dwelling at deeper habitats may experience a “competition release”, mainly by the absence of macroalgae and other fast-growing autotroph organisms that are constrained to the photic zone (Zabala & Ballesteros 1989). Similarly, deeper habitats are less diverse and thus organisms could face less predation pressure, pushing selective forces to favour somatic maintenance and therefore life extension (Healy et al. 2014). Finally, the relative predictability of deep environments may enable species to survive with lower reproductive and mortality rates. Shallow habitats are more exposed to external sources of mortality due extreme physical disturbances (i. e. hurricanes and storms) and temperature fluctuations (Garrabou et al. 2002; Bridge et al. 2013), making them less suitable for organisms whose long-term success rely on extremely high survival of adult stocks (Linares et al. 2007).

Previous anecdotal data on longevity in sessile and mobile marine

organisms support our findings. For instance, the highest longevity reported to date corresponds to the sponge *Monorhaphis chuni*, found at 1000m depth with an estimated age of 11000 yrs (Jochum et al. 2012). A similar pattern has been also observed in marine fish (Cailliet et al. 2001). Indeed, a recent study based on radiocarbon dating on eye tissue revealed that, the deep-resident Greenland shark *Somniosus microcephalus*, which can live up to 400 yrs, may be the longest-lived vertebrate on Earth (Nielsen et al. 2016). Overall, these results demonstrate a strong role of depth-related environmental gradients in shaping life-history strategies across marine taxa. The strength of this pattern is striking and contrasts with the weak environmental correlates of longevity or other life history traits that seem typical in terrestrial systems (Stearns 1977, McIntyre et al. 1995, Morrison & Hero 2003, Moles et al. 2007). Nonetheless, our findings open new questions regarding the existence of potentially analogous processes driving the evolution of life-history

strategies along elevation and/or productivity gradients in terrestrial landscapes. Further analyses exploring the relationship between depth and different environmental parameters could provide a more mechanistic understanding of this general pattern of increasing species maximum longevity towards deeper marine waters.

At least three demographic mechanisms may contribute to the strong relationship observed between depth and longevity in marine sessile organisms. First, species may show relatively indiscriminate settlement. Under this scenario, the longevity pattern may arise because faster-growing, shorter-lived species out-compete slow species in productive, shallow sites, while at deeper sites only slow-growing species with high longevity may persist. Second, species may have evolved specific settlement cues so that larvae of long-lived species preferentially settle in either deep or shallow sites. Finally, to some extent these life history patterns could result from

demographic compensation, reflecting phenotypic plasticity in life histories (i. e. fecundity, growth or survival rates) as a function of local constraints on population growth rates (Villellas et al. 2015). In reality, all three mechanisms are likely to contribute to this matching of life history to environment.

The extreme life-history features of the red coral *Corallium rubrum*

While our survey shows that mesophotic and deep-sea ecosystems are predictably inhabited by species of extreme longevity, we have few detailed studies of the population ecology of these taxa. We found that *C. rubrum*, a widely distributed Mediterranean red coral, showed a consistent pattern of slow population dynamics driven by extremely high survival, recruitment limitation, and apparently negligible senescence (Fig. 5 & S3a). The species revealed a maximum potential longevity up to 532 yrs, overlapping with lifespan ranges of other deep-sea organisms. Although previous research had shown that *C.*

rubrum settlement can be highly heterogeneous (Bramanti et al. 2005, Santangelo et al. 2012), in this study we observed a much more homogenous pattern of low recruitment in the long term (Fig 4b, 4d, Table S8). Recruitment limitation has been observed in other long-lived temperate invertebrates (Grigg 1988, Teixidó et al. 2011, Kersting et al. 2014) and can seriously hinder the ability of populations to recover after intense perturbations (Hughes & Tanner 2000).

For species with limited recruitment, adult survival becomes of paramount importance to ensure their persistence (Linares et al. 2007). We found extremely high survival for *C. rubrum* (Figs. 4a & 5a), with estimated mortality even lower than found in previous experimental studies (Garrabou & Harmelin 2002, Teixidó et al. 2011). In agreement with previous observations in long-lived gorgonians (Linares et al. 2007), perturbation analysis supported that survival is the vital rate

demonstrating the highest effect on population growth in *C. rubrum* (Fig. 5b). The absence of detectable senescence is common in sessile marine species and terrestrial plants (Tanner 2001, Baudisch et al. 2013), and may be related to high investments in structural tissues, although some authors have suggested that modularity may also play an important role (Tanner 2001). It should also be noted that our data, like that for most other studies of extremely long-lived species, lack the precision necessary to carefully test for senescence. Precious corals build an energetically demanding hard carbonate skeleton; while preventing a fast rate of colony growth, it may provide the structural basis that allows for an extremely high survival.

The shallow occurrences of the monitored populations of this long-lived species could seem to make it an outlier in the depth-longevity pattern. However, this coral is distributed from shallow to deep waters up to 1000m depth

(Knittweis et al. 2016) and its shallow occurrences are exclusively found in dim-light and dark habitats where it cannot be outcompeted by fast-growing organisms (Zabala & Ballesteros 1989). Given the overall pattern observed for marine species, the dynamics of red coral populations dwelling at deeper habitats are likely to be even slower and more dependent on adult survival. However, we must be cautious with these extrapolations and note that demographic data on deeper locations is not available due to logistical constraints. Exceptional shallow presence has been also observed in other important cold-water corals, such as *Lophelia pertusa*, which can occur at 25m in sediment laden fjords where light attenuation reduces the abundances of autotroph organisms, and thus competition; while at the other end of the scale this species may be found as deep as 3300m (Squires 1959).

Population-level consequences of extreme life-histories

As important as unraveling the drivers of extreme longevity are the implications of these strategies for dynamics of natural populations. We show that maximum longevity strongly predicts temporal stability in population growth across terrestrial and marine sessile species (Fig. 3). This finding agrees with previous studies suggesting that long-lived species can strongly buffer environmental stochasticity (Morris et al. 2008, García et al. 2008). As observed for the red coral *C. rubrum*, this buffering capacity relies on high adult survival and low reproductive success. While conferring stability under natural conditions, this extreme life history will also hinder the ability of long-lived species to overcome increased mortality rates, but whether these species will be able to cope with rapidly changing perturbation regimes driven by the ongoing global change is still unclear. For instance, while the red coral has persisted after millennia of historical

overharvesting due to the combination of consistent survival of partially harvested colonies (Montero-Serra et al. 2015) and small size at maturity (Torrents et al. 2005), new human-related stressors such as global warming have increased mortality rates and put some shallow populations at risk (Cerrano et al. 2000, Garrabou et al. 2001, 2009). Worryingly, although declines may already be driving populations of some long-lived species towards collapse, in some cases, these trajectories may be too subtle to be noticed by ecologists and managers (Hughes et al. 2013). The consistent depth relationship for longevity also implies greater sensitivity of species and communities occurring at greater depths to human perturbations, amplifying recent calls to better monitor and protect these vulnerable ecosystems (Danovaro et al. 2017).

ACKNOWLEDGEMENTS

We thank K. Kaplan, M. Pages, A. Griffith and one anonymous reviewer for their valuable comments on early versions of this manuscript. This study

was partially funded by the Spanish Ministry of Economy and Innovation Biorock project (CTM2009-08045), the Smart project (CGL2012-32194), the TOTAL Foundation Perfect Project and the European Union's Horizon 2020 research and innovation programme under grant agreement No 689518 (MERCES). This output reflects only the author's view and the European Union cannot be held responsible for any use that may be made of the information contained therein. IMS was supported by a FPI grant (BES-2013-066150), CL by a Ramon y Cajal (RyC-2011-08134), and JBL by a Postdoctoral grant (SFRH/BPD/74400/2010) from the Portuguese Foundation for Science and Technology. Support for DFD came from National Science Foundation awards 1340024 and 1242355. IMS, CL, JBL and JG are part of the Marine Conservation research group (2014 SGR 1297) from the Generalitat de Catalunya.

CHAPTER II

Harvesting effects, recovery mechanisms, and management strategies for a long-lived and structural precious coral

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INTRODUCTION

Overfishing is a major threat to the integrity of world's marine ecosystems. Historical exploitation of marine resources has resulted in drastic population declines, species extinctions and the general simplification of marine food webs (Pauly et al. 1988, Jackson et al. 2001). This structural impoverishment has also hindered the resilience of marine populations, lowering their ability to recover after recurrent natural and human disturbances (Hughes 1994, Hughes et al. 2003). Understanding the ecological consequences of different extractive practices on the structure and function of populations, especially of those organisms exhibiting a long life span and slow dynamics, is therefore one of the greatest challenges in conservation biology (Lotze et al. 2011).

Precious corals have been harvested and traded worldwide since ancient times due to the high economic value of their carbonate axial skeleton (Grigg 2008, Bruckner

2009, Tsounis et al 2010). These sessile invertebrates are considered habitat forming species with a keystone role on coastal systems because they provide structural complexity and host high levels of biodiversity (Gili & Coma 1998). Additionally, they can enhance fisheries by providing shelter during early stages for some commercially important species of rock fishes, shrimps and crabs (Husebø et al. 2002). The red coral *Corallium rubrum* is a precious octocoral endemic to the Mediterranean rocky bottoms and adjacent Atlantic waters. Available data showed recent large declines in the Mediterranean yields, suggesting that this fishery is unsustainably managed following boom and bust cycles (Tsounis et al. 2007, Bruckner 2009). Intensive harvesting has resulted in significant shifts in the size structure of current *C. rubrum* populations (Santangelo et al. 1993, Bruckner 2009) causing a decrease in biomass and mean and maximum colony size (Garrabou & Harmelin 2002, Linares et al. 2010).

Moreover, current warming and acidification trends associated to global change are major threats for shallow red coral populations (Garrabou et al. 2001, 2009, Bramanti et al. 2013, Cerrano et al. 2013). Consequently, there is a growing concern for the conservation of *C. rubrum* and the rest of the precious corals worldwide as evidenced by the recent struggle to include the Genus *Corallium* in the Appendix II of Cites (Bruckner 2009, 2010, 2013, Tsounis et al. 2013).

The need for a responsible fishery for *C. rubrum* and the negative effects of current practices are widely accepted (Tsounis et al. 2007, 2010, Cau et al. 2013). However, given the fragility and vulnerability to different threats of this species, all studies addressing harvesting effects and recovery have been carried out by comparing size-class distribution of different populations and periods (Tsounis et al. 2007, García-Rodríguez & Massó 1986, Abbiati et al. 1992,

Bavestrello et al. 2015). To date, biomass reduction has been only quantified once after a poaching event (Linares et al. 2012). Beyond the reported shifts in size structures of *C. rubrum* populations (Tsounis et al. 2007, Abbiati et al. 1992), sound field and experimental data on the effects of different extractive practices and the recovery process are essentially lacking in the literature.

Due to their low population dynamics and limited dispersal capacities (Ledoux et al. 2010a, 2010b) precious corals are considered to have low resilience to disturbances that cause high adult mortality such as harvesting. Paradoxically, *C. rubrum* has not undergone extinction despite being exposed to intensive harvesting since ancient times. In fact, the biological processes underlying coral populations' persistence are largely unknown. The early maturity of *C. rubrum* (2.4 cm in height) corresponding to an age of 6-10 yr (Torrents et al. 2005, Gallmetzer et

al. 2010), could partially explain this persistence. It allows the colonies to contribute to reproductive output before they reach the size of interest for harvesters. On the other hand, colonial organisms have the capacity to recover after events of partial mortality (e.g. breakage of a branch). Indeed, red coral colonies with signs of breakage and recent re-growth of new branches have been observed (Torrents et al. 2005, Hereu et al. 1999). The resilience to harvesting will therefore depend on two main mechanisms: (1) re-growth or clonal growth of colonies suffering partial mortality (i. e. when leaving the basal section) and (2) recruitment by sexual or asexual reproduction (i. e. when eradicating the whole colony). Yet, there is no assessment on the relative contribution to recovery of these two mechanisms from a long-term perspective. This assessment would provide a scientific basis for new management measures that enhance the sustainability of this natural resource.

In this study, we examined harvesting effects and the recovery processes of *C. rubrum* using 5-7 years photographic series on two populations located on the coast of Marseilles (France, NW Mediterranean). Fishermen unexpectedly harvested two *C. rubrum* populations that were already being studied by our research team and this represented a unique opportunity to establish before / after comparisons as well as to explore directly the recovery mechanisms by following individually affected as well as unaffected *C. rubrum* colonies. Our overall goal was to elucidate the demographic process underlying the recovery of *C. rubrum* after harvesting events with a particular focus on the relative contribution of re-growth of harvested colonies and reproduction. More precisely we quantified (1) survival and re-growth rates of harvested colonies, (2) survival of non-affected colonies and (3) recruitment rates and survival of recruits. Based on these data, we developed biomass

projections to compare population trajectories under different harvesting practices. We contend that the results provide meaningful insights to inform new legislation measures to promote resilience and ensure the conservation of this threatened species and other precious corals also at deeper depths, where most of fishing efforts are concentrated and this type of data is more difficult to obtain.

METHODS

Study Area

We monitored two Mediterranean red coral populations located along the rocky coast of Massif des Calanques in the SE of Marseilles, France. Due to the specific habitat characteristics of the area, such as submerged cavities and overhangs, along with instability of the water column during summer (Bensoussan et al. 2010), the development of red coral populations at shallow depths (15-22m) is favored. The studied populations are located in a vertical wall at Riou Island (43° 10' 23.47"

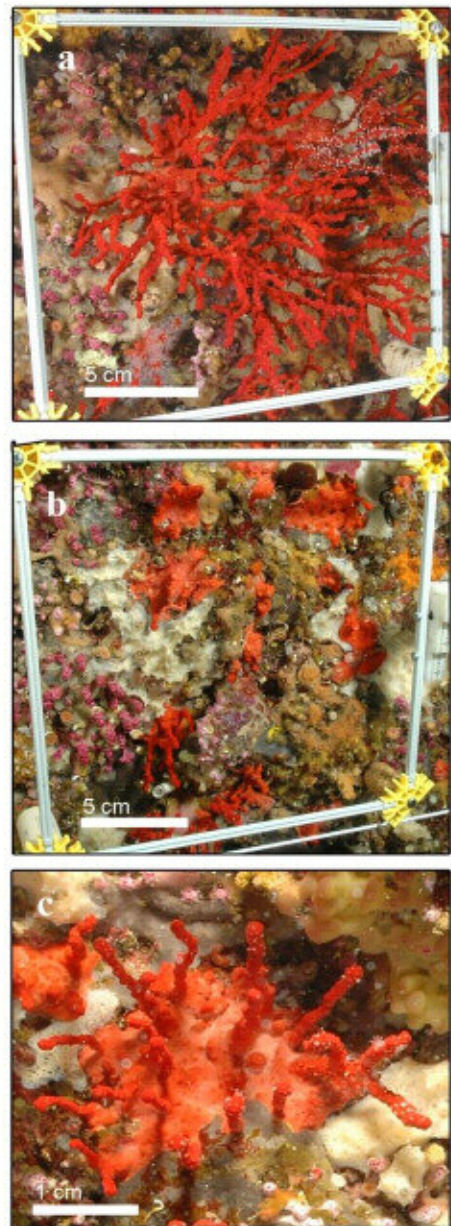


Figure 1. Harvesting effects on red coral populations. Partial mortality at Riou is shown a) before and b) after a harvesting event. c) Detail of re-growth of new branches on a partially harvested colony.

N i 5° 23' 13.24" E) (hereafter Riou) and in a cave-like tunnel at Maire Island (43°12'32.34"N; 5°20'14.01"E) (hereafter Maire).

Coral Population Monitoring

In each population we monitored 30 quadrants (20 x 20 cm) by setting up 2 permanent plots using PVC screws fixed to holes in the rocky substratum. Each plot was variable in length, depending on the complexity of the substratum, and 40 cm wide. In each sampling, a cord was deployed between the screws and quadrats assembled with a scale were sequentially positioned and photographed above and below the cord throughout the length of each transect. Two photographs from each quadrat (using 2 slightly different angles, ~30°) were used for analysis with photogrammetric techniques which allowed measuring the height of colonies (Drap et al. 2013). Transects were photographed using a NIKON D70 with a housing and 2 electronic strobes.

At Maire, permanent plots were installed at 15 m after a harvesting event was observed during June 2002 and monitored till 2009 but due to logistic constraints, pictures were not taken during 2007 and 2008. Colonies that suffered partial mortality were evident because fishermen remove the branches leaving a wide colony basis, short in height with injuries (naked tissue), that unequivocally belongs to an older red coral colony that was recently pruned. In contrast, at Riou the permanent plots were installed in 2005 just before the harvesting event. Between April and June 2006 fishermen were observed in the area and subsequent surveys clearly detected their impact in the permanent plots (Fig. 1).

Our study did not involve any sampling of or damage to red coral colonies. The authors had all the permits provided by the French authority *Affaire Maritimes* to perform scientific surveys in the study area.

Demographic parameters

From the photographic series, the harvested and non-harvested colonies were individually identified and compared year to year to estimate mortality rates, number of branches and colony height. We also estimated adult population density (colonies number per 400 cm²) and recruitment rates. We considered a recruit the new button like colonies observed ranging from a minimum detection size of 2 mm to 10 mm in height. Annual recruitment rates were estimated as the number of recruits appearing for the first time in a photoquadrat and post-recruitment mortality were estimated by following these recruits through the study period.

With this dataset we were able to assess:

3.1 Changes in size-structure. To assess population size-structure changes and the recovery process, we measured maximum height of all colonies and classified them into five size classes (0-30mm; 30-

60mm; 60-90mm; 90-120mm, >120mm). Riou population was measured before harvesting (2005), right after (2006) and four years after (2009). Maire population was measured right after harvesting (2002) and seven years after (2009). Biomass variations were estimated by applying a height-weight polynomial equation to height data. The equation was previously calculated from height and weight data measured on 300 dead red coral colonies collected in previous studies in the same region (Torrents 2007), and from different poaching events (Linares et al. 2012). The resulting curve was: $\text{Weight (g)} = 0.001(\text{Height, mm})^2 + 0.096(\text{Height, mm}) - 4.010$ ($R^2 = 0.868$, $P < 0.001$) (Fig. S1).

3.2 Recovery mechanisms.

We quantified the cumulative survival probability of non-harvested adults, harvested adults, and new recruits based on annual mortality rates, calculated as follows:

(1)

$$m = \frac{\left(1 - \frac{N_t}{N_0}\right)}{\Delta_t}$$

where m is the annual mortality rate of *C. rubrum* colonies, N is the total number of colonies, and Δt is the duration of the study period. In addition, we assessed the degree of recovery of affected colonies by quantifying the frequency of colonies showing new branches and the number of branches per colony during the study period.

4 Simulations of recovery process under different harvesting practices

To analyze whether recovery rates depend on harvesting practices in the long-term, we developed a set of basic stochastic demographic models under two different scenarios with contrasting levels of total (TM) and partial mortality (PM): a) removal of almost all colonies causing 90% TM and 10% PM; b) leaving the basal section in almost all colonies causing 10% TM

and 90% PM. The second scenario was based on the observed mortality values at Riou (see results). The two scenarios were calculated by setting initial population densities according to the corresponding levels of total mortality. To run the model we used demographic parameters (adult density, annual recruitment rates, adult and post-recruitment mortality, and biomass increase rates per colony) obtained from the two monitored populations during the study period. The two harvesting scenarios were projected by running 100 stochastic simulations during 30 years according to the following steps:

First, we estimated the changes in the number of colonies following the next formulation:

(2)

$$N_{T(t)} = N_{R(t)} + N_{A(t)}$$

$$N_{R(t)} = N_{R(t-1)} (1 - g) (1 - m_R) + r$$

$$N_{A(t)} = N_{A(t-1)} (1 - m_A) + N_{R(t-1)} g (1 - m_R)$$

where N_T is total population, N_R is total number of recruits, N_A is the total number of adults, m_R and m_A are annual mortality rate of recruits

and adults respectively, r is the annual recruitment rate and g is the proportion of recruits growing to the adult stage:

(3)

$$g = \frac{G}{h_{\max} - h_{\min}}$$

where G is the recruit growth rate ($2.5 \text{ mm} \cdot \text{yr}^{-1}$ in height, from Bramanti et al. (2005)), h_{\min} and h_{\max} are the lower and upper height intervals for recruits.

Secondly, we estimated the evolution of biomass recovery using the biomass increases (ΔB) per colony and abundance values from the demographic models previously described. Mean (SD) biomass increases were estimated according to the next equations applied to

colonies suffering partial mortality at Riou ($n = 57$) and Maire ($n = 67$) right after harvesting and 4 and 7 years after respectively:

(4)

$$\Delta B = \frac{1}{N} \sum_{i=1}^N \left(\frac{B_f - B_0}{\Delta t} \right)_i$$

(5)

$$B_{(t)} = NT_{(t)} \cdot \Delta B$$

where B_0 is the colony biomass right after harvesting, B_f is the total biomass at the end of the study period, Δt is the number of years, and NT is the total number of colonies in each population.

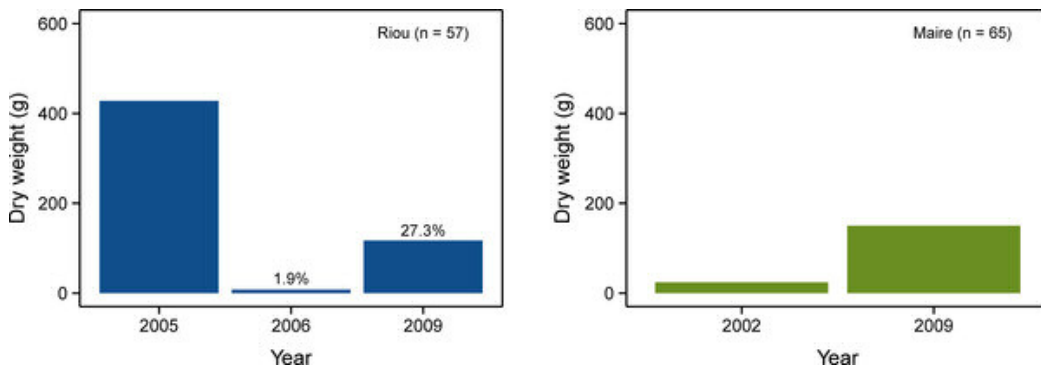


Figure 2. Biomass changes during the study period at Riou (left) and Maire (right).

RESULTS

Harvesting effects

Harvesting affected 30.1% of the colonies at Riou ($n = 209$) and 49.6% of the colonies at Maire ($n = 139$). At Riou, total mortality was around 9.5% and partial mortality was 90.5%, causing a large biomass loss with a 98.5% of reduction in total dry weight (Fig. 2). Harvesting targeted mainly medium and large

colonies, triggering a dramatic size-distribution shift toward populations characterized by a large proportion of small colonies (Fig. 1 and 3a). At Maire, because the monitoring started after the harvesting event, we could not determine the loss of biomass nor the size-class distribution shift. Nonetheless, the same size-distribution pattern was observed after the harvesting event with no large colonies present and a

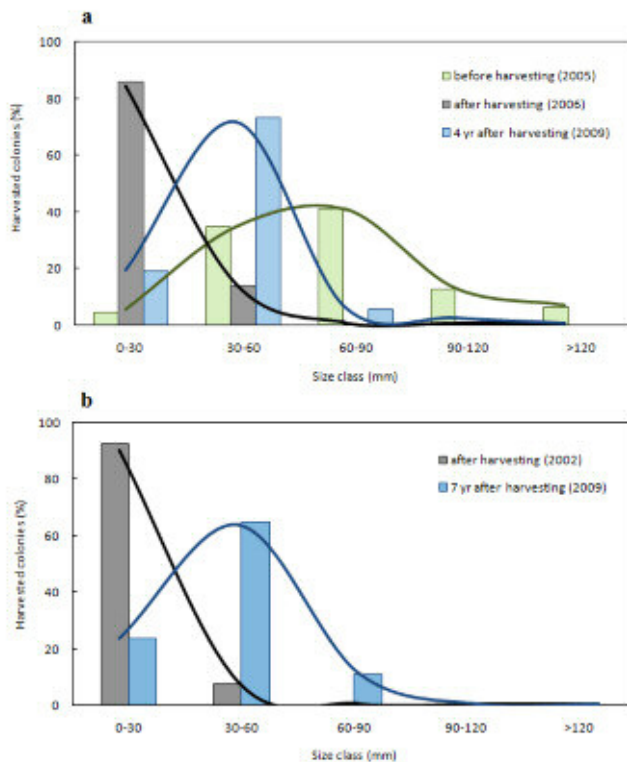


Figure 3. Size-distribution changes on harvested red coral populations. a) Size-distribution at Riou before harvesting, after harvesting, and four years after harvesting. b) Size-distribution at Maire after harvesting and seven years after harvesting.

great proportion of small-size colonies (Fig. 3b).

Temporal trends

Demographic parameters did not show large differences among sites.

Mean annual density was 12.26 ± 0.80 colonies \cdot 400 cm⁻² at Riou (n = 30) and 11.36 ± 0.52 colonies \cdot 400 cm⁻² at Maire (n = 30). Densities were quite stable during the study period (Fig. S2), and recruitment

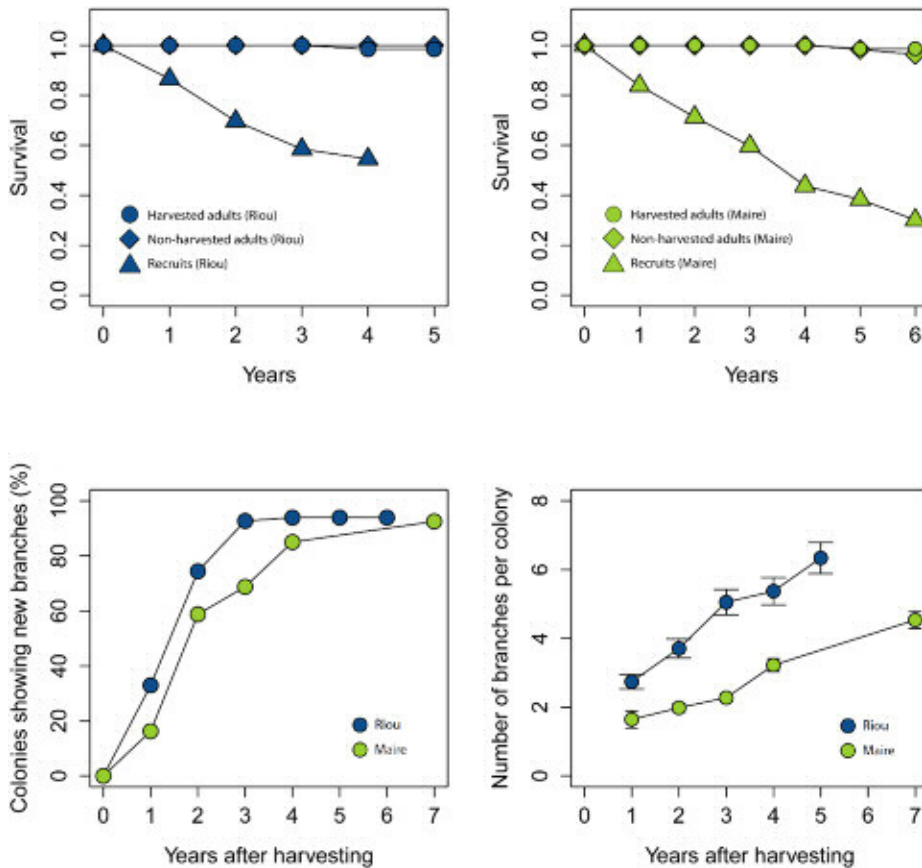


Figure 4. Demographic parameters estimated during the study period at Maire and Riou populations. Cumulative survival probability estimated at Riou (a) and Maire (b) for non-harvested colonies, harvested colonies, and recruits. (c) Temporal variation on the proportion of colonies showing new branches after fishing. (d) Temporal variation of mean (\pm SD) number of branches after harvesting.

rates were low at both sites: 0.98 ± 1.40 (SD) colonies $\cdot 400 \text{ cm}^{-2}$ at Maire and 0.21 ± 0.53 (SD) colonies $\cdot 400 \text{ cm}^{-2} \cdot \text{yr}^{-1}$ at Riou. Harvested colonies suffering from severe partial mortality did not show a

differential "post affection" survival as they showed similar high values than unaffected colonies (Fig. 4a, 4b). This could explain the observed stability in adult density in both populations throughout the study

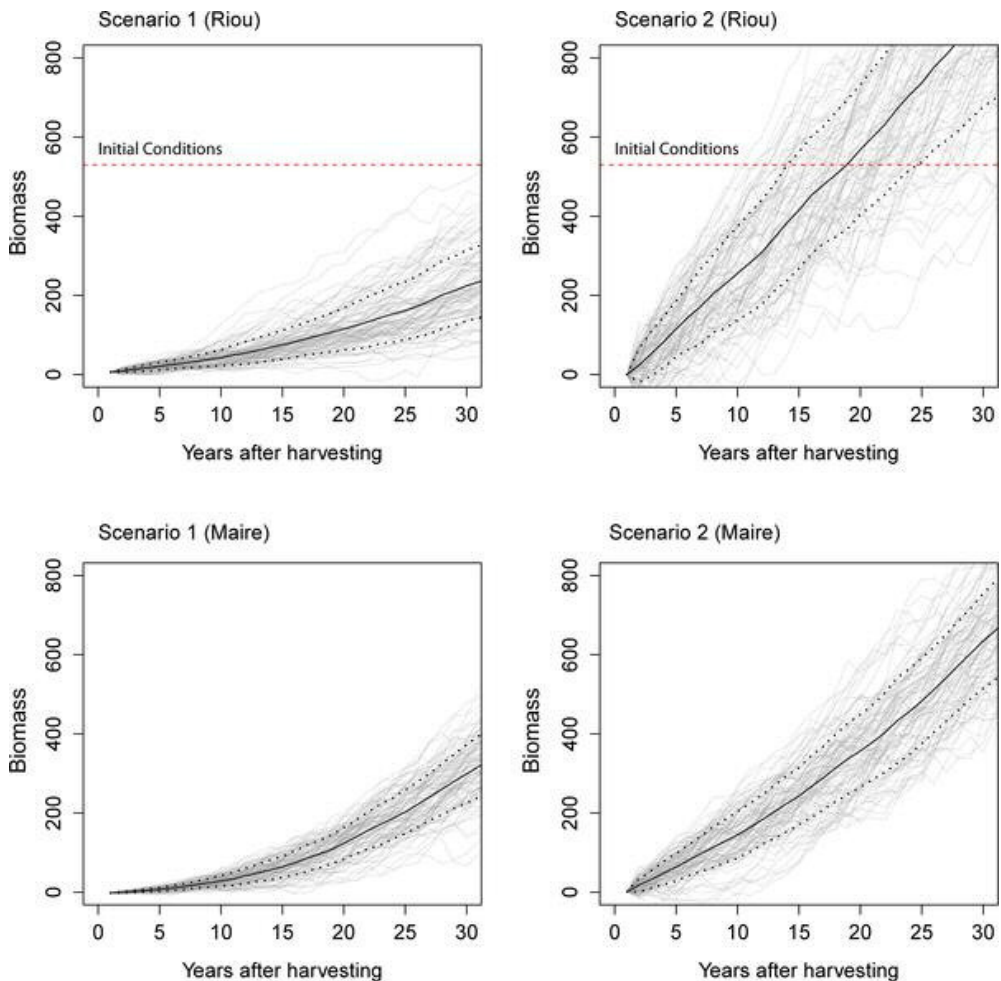


Figure 5. Biomass projections of red coral populations to compare different harvesting scenarios. Scenario 1) 90% total mortality and 10% partial mortality. Scenario 2) 10% total mortality and 90% partial mortality. Horizontal broken lines show initial conditions at Riou. Black lines represent the mean trend and dot lines the standard deviation ($n = 100$).

period (Fig. S1.2). Conversely, survival of recruits was very low and decreased rapidly at both populations (Fig. 4a and 4b).

Rate of recovery

The two studied red coral populations demonstrated very low rate of recovery. A small increase of biomass was observed at Riou after four years, representing 22% of the initial biomass recorded prior to the harvesting event (Fig. 2). A slight recovery was also observed by comparing size-class distribution from Riou and Maire populations after four and seven years respectively (Fig. 3). Despite a general increase in the proportion of medium size-classes, there was no recovery of the largest sizes (>100mm) and very little of colonies >80mm at Riou. At the end of the study, small colonies (0 - 60 mm) still represented a large proportion of the whole population (92% at Riou and 89% at Maire). Re-growth was observed in more than 75% of the harvested colonies after two years at Riou and four years at

Maire (Fig. 4c). Mean number of branches in harvested colonies also increased quickly in both populations (Fig. 4d).

Recovery process simulations

Biomass simulations based on demographic projections showed that harvesting practices strongly influence the recovery process of *C. rubrum* populations. Harvesting practices in which the colony basis remains attached showed recovery times from 15 to 25 yr, whereas events causing total mortality (eradicating the whole colony) slowed down the recovery process, preventing the populations from recovering to initial conditions even 30 years after being harvested (Fig. 5). Riou population showed a slightly higher recovery rate than Maire population. However, the influence of harvesting practices at slowing down the recovery rate when increasing total mortality was consistent for both populations (Fig. 5).

DISCUSSION

This study provides new insights into the recovery mechanisms of a heavily exploited precious coral and illustrates how different harvesting practices can strongly modulate the recovery of affected populations. Our results also highlighted the detrimental effects of harvesting and the low rate of recovery displayed by these long-lived invertebrates.

Harvesting heavily impacted red coral populations has led to drastic reductions of biomass and size-structure shifts towards populations dominated by small colonies. Although both populations still persisted at the end of the study period (after 4 and 7 years), they were far from initial conditions in terms of biomass and sizes. The slow recovery processes reported here confirms the long-lasting negative consequences of harvesting on slow-growing sessile organisms and indicates that extractive practices under the current legal framework are not sustainable from

a long-term perspective (Lotze et al. 2011).

We investigated the relative contribution of reproduction and re-growth of harvested colonies as drivers of recovery of red coral populations. The high survival of harvested colonies suffering partial mortality and fast growth of new branches demonstrated that re-growth plays a key role in recovery and persistence of *C. rubrum* (Fig. 4c, 4d). Similar to previous long-term studies conducted in French and Spanish localities (Garrabou & Harmelin 2002, Linares et al. 2012), the observed low recruitment rates and very low survival probability of new recruits suggest that recovery through reproduction is very limited (Fig. 4a, 4b). This demographic strategy is shared by other temperate gorgonians (Linares et al. 2007), and is concordant with life-history theory, which states that adult survival is inversely proportional to reproductive success (Heppell et al. 2000). Indeed, biomass simulations developed to compare different

harvesting practices showed that when the whole colony is removed and thus reproduction is the main recovery mechanism, the process is much slower and it can take much more than 30 years to return to initial conditions previous to harvesting. Conversely, when fishermen leave the basal section of the colonies recovery is enhanced through re-growth of new branches (Fig. 5). The biological reference point used in this study (size-class distributions at Riou prior to harvesting) does not represent an undisturbed population; therefore, recovery leading to pristine conditions would take longer periods than those reported here. Further, we must note that the models used here do not completely account for the complexity of demographic processes in long lived species. *C. rubrum* is characterized by a low dispersal that leads its populations to be generally self-seeding (Costantini et al. 2007, Ledoux et al. 2010a), and breeding units seem also to be restricted in space, suggesting that density may

play an important role in the reproduction of this species (Ledoux et al. 2010b). Thus, events causing total mortality on adult colonies could lower the potential for recovery even more through reproduction (Lasker 2013). Alternatively, negative density-dependent processes affecting recruitment and post-recruitment survival were also reported in corals due to intra-specific competition and resources limitation (i. e. availability of suitable substrate to settle) (Vermeij & Sandin 2008). Further research should therefore address these questions by applying more complex modeling techniques to long-term experimental data. However, the observed large and consistent differences on population trajectories depending on harvesting procedures revealed by our study represent an important first step in the characterization of the ecological consequences of different harvesting practices on *C. rubrum* populations.

Reported recent declines on the Mediterranean yields sparked a wide

debate about the conservation status of *C. rubrum*. This discussion raised questions about the sustainability of this fishery and if this species should be categorized as threatened (Bruckner 2009, 2010, 2014, Santangelo & Bramanti 2010). Our results show that populations demonstrate a high degree of persistence in terms of density due to high survival of affected colonies. This may explain the existence of red coral populations despite intensive harvesting during the past few centuries. Nevertheless, we emphasize the large reduction of biomass and the size-distribution shifts toward populations dominated by small colonies likely hindered their structural function. A general simplification of benthic communities may also have negative consequences for multiple trophic levels that use these habitats as shelter during early life-stages (Jones et al. 1994, Husebø et al. 2002). This finding should be especially relevant for policymakers. Indeed, abundance is a parameter commonly used by international

organizations such as IUCN or CITES to categorize vulnerability levels in commercial species (Bruckner 2009, UICN 2014). However, our results challenge the suitability of abundance-based metrics to assess the conservation status of clonal (modular) organisms by showing that large declines in biomass due to partial mortality of colonies can remain masked. Furthermore, abundance measures do not reflect the reduction in reproductive potential caused by harvesting colonial organisms, which exhibit a size-based exponential increase in reproductive output (Santangelo et al. 2003). The development of a new metric focused on size and biomass parameters is crucial to assess the conservation status of precious corals and to develop sustainable fisheries management plans.

On the other hand, benthic communities are currently facing multiple perturbations derived from the ongoing global change such as warming and ocean acidification

(Garrabou et al. 2001, 2009, Coma et al. 2009, Cerrano et al. 2013). The precautionary principle accounting for potential synergistic effects of fishing and climate warming should thus be considered in these new regulations (Garrabou et al. 2001, 2009, Torrents et al. 2008). While climatic perturbations are generally diffuse and difficult to minimize, fishing restrictions on no-take areas can be useful tools at a local scale to enhance the resilience of sessile invertebrates through increased larval production of large individuals (Micheli et al. 2013). MPAs also enhance the recovery of exploited populations in terms of biomass and size-structure (Linares et al. 2010, 2012). However, given the low recruitment rates and limited dispersal of gorgonian larvae (Ledoux et al. 2010a), it is unlikely that protected populations act as a source to exploited populations ensuring their long-term persistence. Further, poaching and diving activities hinder the enhancement of coral populations' recovery within MPAs when regulations are not

enforced (Hereu et al. 1999). Therefore, we should not just consider improving the actual network of MPAs but also moving towards a more restrictive legal framework outside of the protected areas. Several new regulations have been proposed during the last decades to improve *C. rubrum* fisheries, including the establishment of annual quotas and minimum harvesting sizes and the ban on using dredges. However, these current guidelines are still largely based on untested assumptions and have failed to improve the sustainability of coral fisheries (Bruckner 2014). For instance, a significant step was the recent recommendation on total ban on harvesting of *C. rubrum* populations under 50m depth (Cau et al. 2013), although poaching in shallow waters is still widespread and individual countries may avoid this prohibition by developing specific management plans. Rotating systems were also proposed for precious corals fisheries in the Pacific and the Mediterranean

though there still is a large uncertainty around the recovery periods of affected populations (Bruckner 2014). Here, we show how red coral populations may take much more than three decades to recover and that total removal of colonies can significantly slow down this process further. Our approximations are consistent with previous studies conducted within French and Spanish MPAs where populations showed only partial recovery after 20 to 30 yrs of protection (Tsounis et al. 2007, Linares et al. 2010). Additionally, there has been a recent development of new jewelry manufacturing processes that allow using small pieces of coral that can be ground to powder and mixed with epoxy or other substances (Cau et al. 2013). This development worsens the effects of harvesting by targeting whole size ranges and including entire colonies. Recovery of populations affected by these new practices might be even more jeopardized since re-growth mechanisms are dramatically

limited. Finally, this study highlights the importance of long-term ecological monitoring programs because they provide reliable data on key ecological processes such as the recovery patterns of natural populations after human-induced disturbances and enable the improvement of conservation strategies.

ACKNOWLEDGEMENTS

We acknowledge the support of all the people that helped us with the photographic surveys during all the years. The authors are part of the Marine Conservation Research Group (2009 SGR 1174) from the Generalitat de Catalunya.

CHAPTER III

**Marine protected areas enhance structural complexity
but do not buffer the detrimental consequences of ocean
warming for an overexploited precious coral**

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INTRODUCTION

The cumulative effects of multiple stressors acting at both local and global scales threaten the integrity and resilience of virtually all ecosystems, and have been particularly of concern for marine ecosystems and their associated services (Halpern et al. 2008, Worm et al. 2006, Ainsworth et al. 2016). The rapid rate of marine biodiversity loss because of overexploitation and climate change is raising major concerns about the effectiveness of current conservation strategies (Hughes et al., 2017). Global policy agreements to reduce greenhouse gas emissions are urgently needed to slow down the current rate of warming and other associated impacts of global change such as ocean acidification (Kennedy et al., 2013, Anthony et al. 2015). Nonetheless, at local scales, conservation management tools such as fishing restrictions and marine protected areas (MPAs) may enhance ecosystem function and resilience to global change (Bozec et

al. 2016, Mumby & Harbone 2010, Mellin et al. 2016, Wolff et al. 2018). Given this dichotomy, there are multiple calls for the use of enhanced local conservation efforts to compensate for delayed and ineffective efforts to slow or reverse climate change impacts (Hughes et al. 2010, Roberts et al. 2017).

At a global scale, climate-induced mass bleaching and mortality events are among the greatest threats faced by tropical and temperate marine benthic ecosystems (Garrabou et al., 2009, Wernberg et al., 2016, Hughes et al., 2018). Several comprehensive studies have reported the immediate detrimental impacts of positive thermal anomalies on habitat-forming species such as corals and sponges (Garrabou et al. 2009, Hughes et al. 2017), but much less is known about the long-term consequences of these recurrent events.

Individual-based demographic data and modeling tools such as structured population models can improve predictions of the effects of global warming, as well as provide a framework to

understand how these effects interact with other stressors.

In a context of rapid environmental change, a critical uncertainty is how effective local management tools such as no-take marine protected areas (MPAs) are in enhancing the resilience of benthic ecosystems to recurrent warming impacts. Some studies suggest positive effects of MPAs on coral cover and

complexity (Linares et al. 2010, Selig & Bruno 2010), recruitment (Mumby et al. 2007) and resilience (Michelli et al. 2012, Olds et al 2014, Mellin et al. 2016). But null (Toth et al. 2014, Bruno & Valdivia 2016, Cox et al. 2016) or even negative relationships between local protection and resistance to climatic disturbances have also been reported (Graham et al. 2008, McClanahan 2008, Hughes et al. 2017).

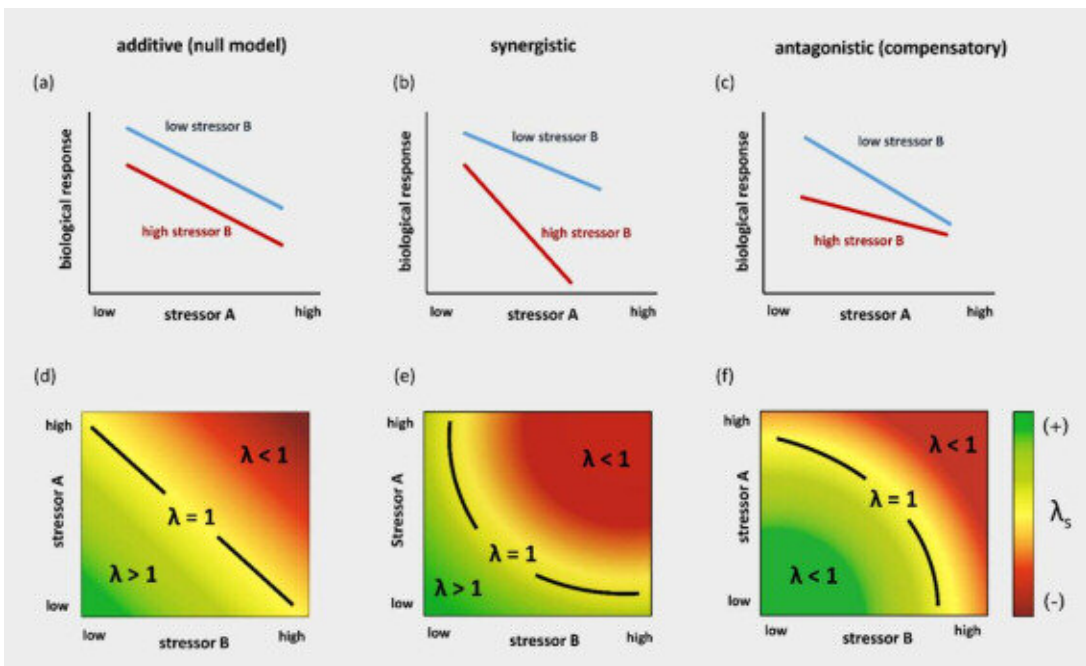


Figure 1. Conceptual models for two interacting stressors. Potential interactions can be additive (left), in which the total impact is equivalent to the sum of the two individual stressors, synergistic (center), when the combined effect leads to a higher impact than the expected addition of the two individual stressors, and antagonistic or compensatory (right), when the combined effects of the two stressors lead to a lower impact than the expected sum of the two individual stressors.

Developing better methods to assess how local management shapes the stochastic dynamics and structural complexity of habitat-forming species is thus a key research need.

The success of local conservation efforts in effectively mitigating for climate change effects will depend on the extent to which different impacts are additive, synergistic or compensatory in their combined effects (Fig. 1). In marine coastal systems, ecological synergies between fishing and climate change have been commonly hypothesized, but meta-analyses and empirical evaluations have suggested that they are less frequent than expected (Darling et al. 2008, Côté et al., 2016). However, most prior work has been carried out through correlational approaches, preventing a clear dissection of the effects of individual drivers. For example, losses of percent cover in corals, or many other easy to monitor measures of impact, can arise from different processes and lead to counter-intuitive interactions between stressors (Darling et al.

2010). Thus, we still need better mechanistic frameworks for understanding how multiple stressors interact to mitigate or exacerbate their negative consequences on vulnerable ecosystems (Côté et al. 2016). While there are different levels of mechanistic analysis, one that can uncover at least some synergies or other interactions is demographic modeling, which tries to tie stressors to impacts on particular vital rates. Unfortunately, obtaining individual-based demographic data can be methodologically challenging in the marine realm, and there are few long-term datasets that allow for an in-depth exploration of the interactive effects of multiple drivers. Here, we apply a demographic analysis to look at the impact of two major threats – ocean warming and fishing– to populations of a long-lived octocoral. We show that, because of their contrasting effects on different vital rates, these threats do not have the strong synergies or compensatory effects that might in theory be expected.

Focusing on the Mediterranean red coral, *Corallium rubrum*, which is a precious long-lived octocoral with an important structural role in Mediterranean hard-bottom communities, we first (1) quantify the effects of fishing (local stressor) and warming-driven mass mortality events (global stressor) to specific vital rates. Then, we (2) predict the long-term consequences of these two different threats. Finally, we (3) evaluate the interactive effects of global change and local protection in the fate of red coral populations. Our results reveal the detrimental effects of climate change on red coral dynamics, and also challenge previous assumptions of MPA capacity to buffer climatic impacts. We also provide insights into how local protection can shape the structural complexity and resilience patterns of temperate reefs. Our approach highlights how population models can be used to integrate individual-based demographic data on different impacts to predict the stochastic dynamics of populations threatened by multiple stressors,

especially when data sets are incomplete.

METHODS

Study System

The Mediterranean red coral, *Corallium rubrum*, is an iconic and long-lived precious octocoral with an important structural role in Mediterranean hard-bottom communities (Ballesteros 2006, Garrabou et al. 2017). During the last several decades, there has been a growing concern about the conservation of Mediterranean red coral and other precious corals worldwide due to unsustainable fishing practices (Tsounis et al. 2010, Brukner 2013, Montero-Serra et al. 2015), and multiple mortality events associated to recent warming in the Mediterranean (Cerrano et al., 2000, Garrabou et al. 2001, 2009, Coma et al. 2009). As a result, this species has recently been listed as an endangered species in the IUCN Mediterranean Red List of Threatened Species due to the risks

associated with overexploitation and climate change (Otero et al. 2017).

Study area and coral monitoring

Eight red coral populations located in different locations in the NW Mediterranean Sea at relatively shallow depths (15 to 25m), were monitored during periods ranging from 3 to 13 years (Table S1). Among the studied coral populations, five were located within MPAs and three were non-protected and subjected to several fishing events during the monitoring (Montero-Serra et al. 2015). One of the protected populations, located in the Réserve Naturelle de Scandola (Corsica, France), has been recurrently affected by positive thermal anomalies since 2003, which have triggered coral mass mortality events (Garrabou et al. 2009, Coma et al. 2009, for a detailed description of study sites see Table S1).

In each population from 2 to 4 replicate transects were marked using plastic screws attached to the substratum. Colonies were

monitored once a year using photographic series, which allow the individual identification of colonies as well as an accurate assessment of life-history processes such as colony growth, partial mortality (e.g. shrinkage and necrosis), mortality and recruitment (for more details see Montero-Serra et al. 2017, Drap et al. 2017).

Demographic analyses

To estimate the spatial and temporal variability in demography for our study species, we fit a set of annual- and population-level models for size-specific vital rates. These vital rates models included colony size and year as fixed effects and were first fit separately for each population and vital rate: logistic GLMs described probabilities of annual survival and extreme shrinkage (described below) rates, a normally distributed GLM described normal growth data, and a Poisson GLM described fecundity.

As for many other clonal species, red coral can both grow and shrink over time. Height data from all

annual photographs were not available, and growth is so slow in this species that even 7 to 8-year time periods only result in mean growth on the order of 10 to 40mm in maximum height (2.74 ± 0.19 mm·yr⁻¹, Montero-Serra et al. 2017). Thus, growth data from two populations over variable length periods (7 and 8 years) were used to fit a single model characterizing mean annual growth rate depending on size. Shrinkage or negative growth is a common process in plants and sessile invertebrates that is usually accounted for by assuming normally distributed variation in size changes around a mean rate (e.g., Salguero-Gomez & Casper 2010). However, for many species assuming normally distributed growth variance is problematic, because a few individuals experience sporadic extreme shrinkage events and these results in a highly skewed distribution of growth rates (Shriver et al. 2012). We accounted for these two types of growth following the procedures proposed by Shriver and colleagues

(2012), describing normal growth and extreme shrinkage as a two-part process (Montero-Serra et al. 2017). Size-dependent extreme shrinkage was fit as two parameters (annual probability of shrinking, and then mean and variance if it occurs). Unlike normal growth, extreme shrinkage probabilities were fit to each population and time-period.

To directly assess the mean effect of different disturbances on coral survival and extreme shrinkage rates, we also fit a set of Generalized Linear Mixed Models (GLMM) including colony size and disturbance level (protected, unprotected, warming) as fixed factors and population and year as random effects (Table S2 & S3).

Integral Projection Models

Based on the linear models characterizing each vital rate, we derived a set of annual size-structured integral projection models for each population (see Montero-Serra et al. 2017 for details on vital rates estimation and model construction). These sets of models

were used to compute the long-term stochastic growth rate at each population, as well as growth rates assuming different combinations of local management and climate effects, as described below.

We computed long-term stochastic population growth rate (λ_s) for each red coral population, based on their observed variation in demography, by running 1000 Monte Carlo simulations over relevant time horizon of 100 years, choosing each of the 3 to 13 annual transition matrices estimated for each population with equal probability. This methodology is commonly used in population viability analysis (Genovart et al. 2017). In the simulations, we randomly drew each of the observed annual matrices with equal probability. Simulations were initiated with a population at the stable stage distribution of the mean matrix for the population. λ_s was estimated as:

$$\lambda_s = \frac{1}{1000} \sum_{i=1}^{1000} \frac{\ln(N_{i,t=100}) - \ln(N_{i,t=0})}{100}$$

Where $N_{total,t}$ is the summed population number across all sizes at time t.

Climate-induced mortality: demographic effects of extreme heat waves

The effects of heat waves were modeled under current conditions using 13yrs of demographic monitoring data in an affected red coral population located in the Scandola Natural Park (Corsica, France). This population has experienced several warming events that led to significant demographic effects. A total of 13 annual-level IPMs were derived from the demographic dataset and were used to forecast the long-term dynamics of this population under current trends. We compared the observed number of coral colonies from 2003 to 2016 to the projected trajectories by starting the simulations at 2003 with the observed initial abundance ($n = 143$ colonies) and size distribution. In addition, to estimate the effects of warming events on size-dependent survival rates (the

main vital rate influenced by these events), we fit a logistic GLM to the survival data of 2016, when the highest intensity Mass Mortality Event was recorded.

Local protection effects on population size and colony-size structure dynamics

We tested the effect of local management on the dynamics of coral populations by simulating 1000 trajectories using 36 annual- and population-level IPMs derived from the demographic data on 4 protected red coral populations within MPAs (21 matrices) and 3 unprotected coral populations (15 matrices) (Table S1). As a starting population vector, we set 100 individuals at the stable stage distribution, which corresponded to the right eigenvector of a mean matrix that included both treatments. In each simulation, we randomly picked one annual transition matrix from the suite of matrices estimated across all years and populations for either protected or unprotected sites.

Structural complexity is an important indicator to assess the health status and functional role of key marine habitat-forming organisms such as corals and gorgonians. Thus, we quantified the proportion of large coral colonies (>100mm) as a proxy for population structural complexity (Linares et al. 2010, Montero-Serra et al. 2017).

Modeling the combined effect of fishing and warming-induced mass mortality events

To explore the interactive effects of local and global stressors (fishing and warming-driven mass mortality events) on the long-term stochastic growth rate and mean time to extinction of red coral populations, we simulated dynamics under 100 scenarios of fishing and warming impacts. To present stable mean results, in these simulations, we only included demographic variation due to local protection and warming events, but not population or annual variation in vital rates. Local management showed no significant effects on the survival rate of coral

colonies but it revealed a strong impact on extreme shrinkage probability. Therefore, we used the predicted mean values from GLMMs for extreme shrinkage probability in protected vs. unprotected coral populations to explore the effects of MPAs. To provide a continuous gradient of local disturbances, we multiplied the effect size estimate of MPA effect by 10 levels (Table S3 and Figure S1). The maximum/high local disturbance intensity corresponds to shrinkage probability outside MPAs (where fishing is allowed), while the minimum/low probability corresponds to shrinkage rates observed in populations within MPAs (where fishing is completely forbidden). On the other hand, a gradient of 10 levels of probability of occurrence of high intensity MME (from 0.05 to 0.95) were set using mortality rates quantified during a high intensity mortality event occurred in 2016 in Palazzu (Fig. 4b), and background mortality rates quantified using data from all monitored populations (Table S1).

These two gradients of local and global stressors resulted in 100 demographic combinations. We compared these different scenarios by computing the long-term stochastic population growth rate for 1000 simulations of the 100 demographic combinations using the same methodology described above. We also used stochastic projections to computed mean time to extinction for each combination of stressors by setting an initial population size of $N_0 = 1000$ individuals, and a quasi-extinction threshold of 10% that corresponded to 100 individuals.

RESULTS

Demographic consequences of fishing and no-take marine protected areas

All seven red coral populations not subject to mass mortality events had very low extinction risk, with long-term stochastic growth rates near at those for a stable population ($X \pm SE$; $\log(\lambda_s) = 0.0108 \pm 0.0044$); and this was consistent across protected and unprotected populations

(ANOVA; $F = 2.033$; $P = 0.2130$; Fig. 2a). Size-dependent mortality rates also did not differ between unprotected sites and protected red coral populations (Logistic-GLMM; $P = 0.6810$; Fig. 2b, Table S2). Projected dynamics using IPMs fitted to data from protected and unprotected populations showed very similar patterns regardless of protection status, with no significant differences on the first 30 years of simulations (Fig. 2c).

In contrast, colony shrinkage rates depended strongly on local management status. The best fitted model for shrinkage probability included a significant interaction between coral colony size and protection status, with much higher shrinkage rates in large and unprotected coral colonies (Figure 2d, Logistic GLMM; $P < 0.001$; Table S3).

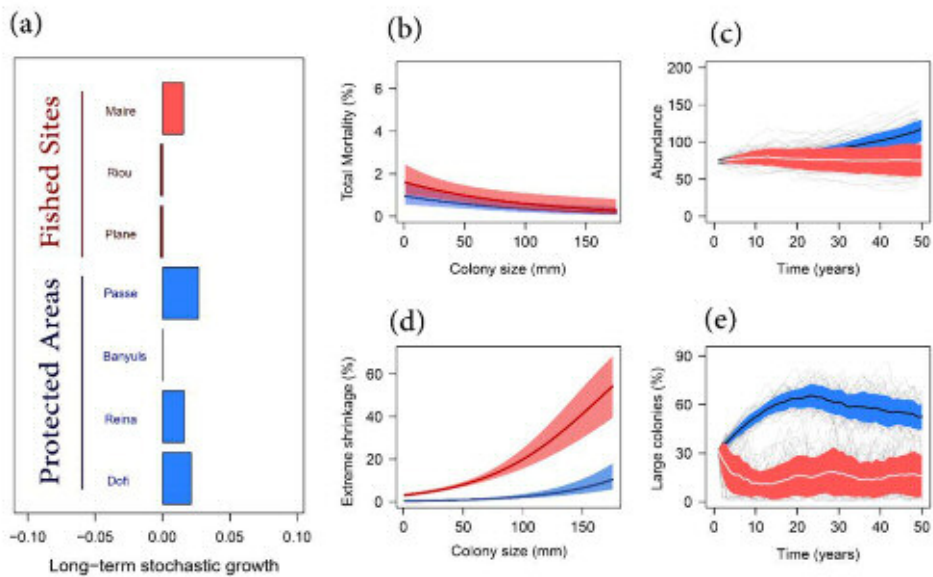


Figure 2. Local protection effects on coral population dynamics and structural complexity. Unprotected sites affected by recurrent fishing events (red) and protected protection within MPAs where fishing is not allowed (blue). (a) Long-term stochastic population growth rate at each site. (b, d) MPA effects in vital rates: (b) total mortality (% of annual dead colonies depending on maximum colony size) and (d) partial mortality (annual shrinkage probability depending on colony size). (c, e) Simulated stochastic dynamics in abundance (c) and structural complexity (e), represented as % of large colonies (>100mm).

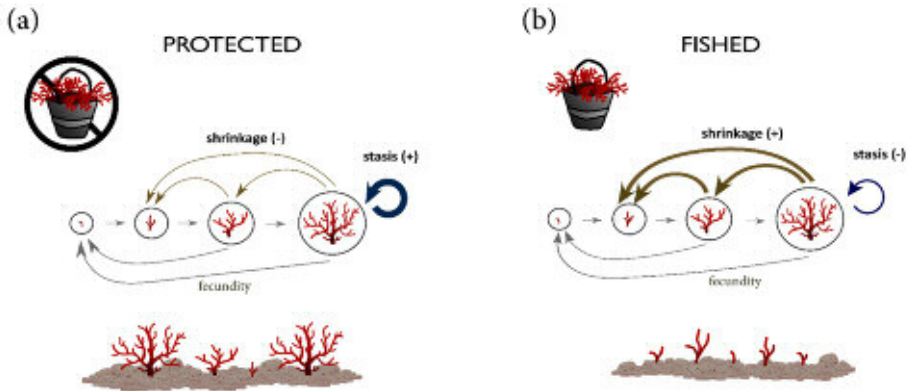


Figure 3. Effects of local protection and overfishing on demographic processes and structural complexity of red coral populations. The conceptual diagram shows how shrinkage rates and stasis of large colonies are shaped by local protection (a) and fishing (b). Protection from fishing results in more complex red coral populations with presence of large coral colonies, whereas recurrent fishing events outside MPAs cause a structural simplification of populations which are dominated by small to medium size coral colonies. (exact values are shown in Figure S2)

When considering the proportion of large colonies as proxy of population structural complexity, projected trajectories of coral populations within and outside MPAs showed divergent trajectories: starting at the stable-stage distribution, protected populations increased their structural complexity while unprotected populations showed a rapid and strong decrease in the presence of large coral colonies (Fig. 1e). This caused a change in the relative importance of life-history processes

depending on local management schemes. In protected populations, stasis was the dominant life-history process, with large colonies remaining within the same state and resulting in high structural complexity. On the contrary, recurrent shrinkage events coupled to slow colony growth drive a structural simplification in unprotected populations, which are dominated by small and medium-size coral colonies (Fig. 3 & S2). Across colony sizes, stasis is the dominant demographic process in both protected and un protected

sites. However, in large colonies, summed stasis probabilities shift from 0.991 in protected populations to 0.802 in unprotected ones (Figure S2).

Demographic consequences of warming-driven mass mortality events

Recurrent positive temperature anomalies caused several mass-mortality events in the red coral population at Palazzu (Corsica)

during the 13 years of demographic monitoring (Fig. 4). These recurrent mass mortality events resulted in a negative long-term stochastic growth ($\lambda_s = 0.98$), and drove a steady decline that may result in the local extinction of the warming-affected population (Fig. 4a). The most severe event was observed during the summer of 2016, and caused whole-colony mortality rates of approximately 25% of the total coral colonies (Fig. 4b).

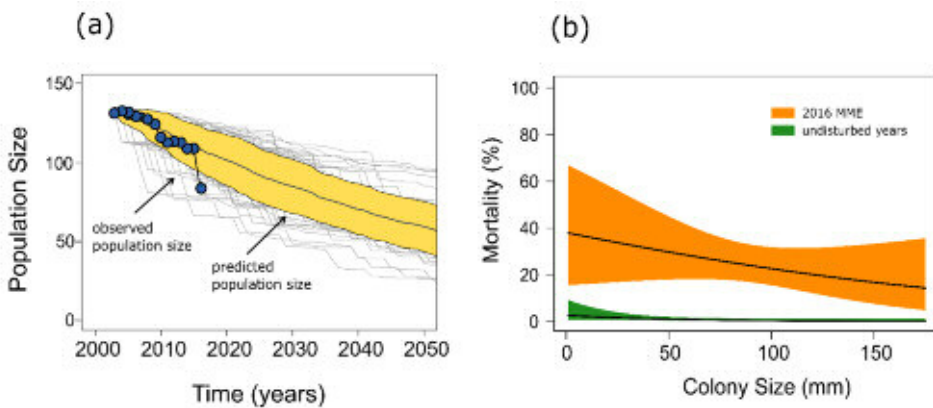


Figure 4. Demographics effects of warming-driven mass mortality events. (A) Long-term stochastic dynamics on a warming-affected red coral population. Dots show the observed abundance during the 13yrs of monitoring. Grey lines represent 100 stochastic projections using IPMs and yellow area represents the 95% CI around the predicted trajectories. (B) Logistic GLMMs describe undisturbed mortality rates across all populations and years (excluding MME) (green) and a GLM describe mortality rates during the high intensity MME recorded in 2016 at Palazzu population (orange).

Interactive effects of local and global stressors

Simulated trajectories under multiple scenarios of differing intensities of global warming and fishing revealed that red coral long-term stochastic population growth rate is highly sensitive to changes in frequency of climate-driven mass mortality events (MME) (Fig. 5 & 6). Local protection was an important factor enhancing population growth under low MME probability. However, its effects were negligible under hypothetical scenarios of increased frequency of mortality events. The interaction between these local and global scale stressors was mainly additive, with only very weak compensatory effects (Fig. 5 & 6).

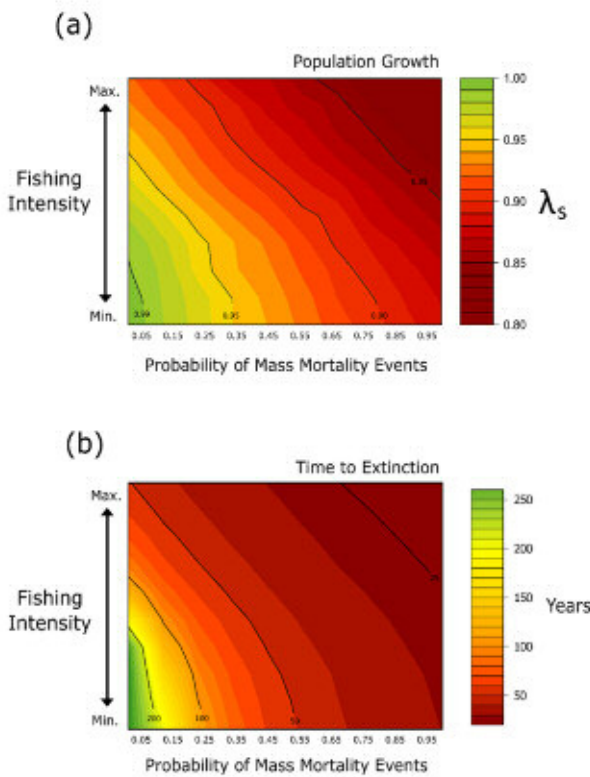


Figure 5. Interactive effects of local and global stressors on the stochastic population dynamics and extinction risk of Mediterranean red coral populations. (A) Long-term stochastic growth rate (λ_s) and (B) shows the mean time to extinction. Vertical axis represents ten levels of local disturbance intensity (see methods and figure S2) and horizontal axis represents ten levels of probability of mass mortality events.

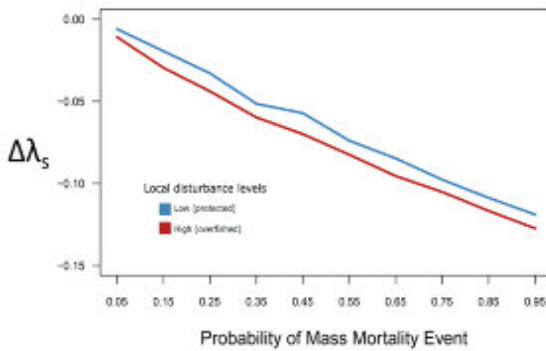


Figure 6. Interaction between local disturbances (fishing) and global warming (probability of Mass Mortality Event) on the long-term stochastic growth rate of Mediterranean red coral populations.

DISCUSSION

Predictions of future ecological change are currently challenged by a poor understanding of how multiple stressors interact to shape the dynamics and structure of populations and communities. Here, we show the strong but divergent demographic consequences of warming-induced mortality events and overfishing on a long-lived octocoral. Recurrent heat waves raise whole-colony mortality rates and compromise the long-term viability of coral populations. By contrast, banning fishing in MPAs

did not affect coral mortality patterns but did reduce shrinkage rates, allowing colonies to attain large sizes and this strongly increased the structural complexity of populations. However, our results suggest that these local and global scale stressors are mostly additive, with MPAs having only a weak compensatory effect on the population-level effects of warming events.

Effects of global warming on red coral populations

In the Mediterranean, several mass mortality events have impacted multiple organisms, most of them slow-growing species, across vast regions (Cerrano et al. 2001, Garrabou et al. 2009). The immediate destructive effects of these widespread climate-driven mortality events in corals and other habitat-forming marine invertebrates are well known (Garrabou et al. 2009, Hughes et al. 2017). However, predictions of the long-term, population-level consequences of these events are still relatively

scarce. In this study, stochastic projections based on individual-level demographic data over a 13-year period in a red coral population subjected to recurrent climatic disturbances revealed clear detrimental consequences for population viability (Fig. 4). Although there still is a large uncertainty around the expected future climatic impacts, current evidences suggest that the frequency and intensity of heat waves has already increased and is likely to continue follow this path during this century (Hughes et al. 2017b, 2018a). This is especially worrisome for the Mediterranean Sea, where warming is occurring at faster rates than in larger ocean basins (Vargas-Yanez et al. 2008, Macías et al. 2013). Under these trends, local extirpation of shallow red coral populations appears likely in a near future. The predicted declines can be even more adverse for long-lived species such as the red coral, in which rescue effects are very unlikely at ecological time-scales due to highly restricted larval

dispersal (Ledoux et al. 2010a, b). For long-lived species with slow population dynamics, catastrophic consequences could go unnoticed by managers when looking at short-term trends (Hughes et al. 2013). Some authors have suggested that adaptive responses can lead to increased thermal resistance of red coral colonies to warming (Ledoux et al. 2015). However, observed increases on the severity of mortality events and their long-lasting consequences provide little empirical support to any adaptive process that could counterbalance these climate change impacts. Furthermore, long generation times for habitat-forming species such as the red coral make rapid genetic change unlikely. The largest survey of impacts of the mass bleaching event occurred in the Great Barrier Reef in 2016 also refutes any optimistic expectations that corals exposed to previous thermal stress may show increased resistance to future warming impacts (Ainsworth et al. 2016, Hughes et al. 2017).

Local stressors and the role of Marine Protected Areas

In the current context of rapid environmental change, understanding the effects of MPAs on population dynamics and extinction risk in marine sessile invertebrates is an essential but unresolved question. Our results show that local protection did not significantly alter red coral numbers or overall dynamics. Recurrently fished populations outside of MPAs displayed low extinction risk and their long-term stochastic growth was nearly at equilibrium (Figure 2a). This counterintuitive result could be explained by the clonal nature of the red coral and its capacity to survive and undergo regrowth of new branches after being partially harvested, which is common when fishers leave the colony bases attached to the substratum (Montero-Serra et al. 2015). It is important to note that highly destructive red coral fishing practices are still observed in some areas of the Mediterranean and they could lead to much worse ecological

effects than those reported here (Cattaneo-Vietti et al. 2017). A slight increase in total mortality rates due to fishing would result in a dramatic increase on the population extinction risk outside MPAs and result on an enhanced role of local protection. Broad-scale quantification on the extent of these unsustainable practices is essential to improve current management strategies for the conservation of *Corallium rubrum* and other precious corals.

In contrast to the lack of effects on numbers, the absence of coral fishing in no-take areas had a strong positive effect on the structural complexity of coral populations by reducing the probability of colony breakage and allowing coral colonies to reach larger sizes (Figure 2d). Outside MPAs, structural simplification was observed with absence of large colonies, driven by both the natural slow colony growth of the species and recurrent colony breakage events due to fishing. Thus, local management can shape the relative importance of the

demographic processes underlying red coral dynamics, enhancing the structural complexity and functionality of red coral populations (Fig. 3 & S2). These results are consistent with previous observations that found that populations within MPAs have larger biomass than unprotected sites, which can harbor more than 200-fold biomass as observed in a recently discovered pristine red coral population (Bavestrello et al. 2014, Garrabou et al. 2017). Here, we provide a mechanistic understanding on how differential shrinkage but similar whole colony mortality rates shape the consistent broad scale patterns observed in colony-size structures and population persistence of the red coral across the Mediterranean Sea (Linares et al. 2010, Garrabou et al. 2017, Fig. 2 & 3).

Interactive effect of local management and global warming on the viability of red coral populations

It remains unclear how effective local management tools such as Marine Protected Areas (MPAs) are in building resilience of benthic ecosystems to climate change. Local protection enhanced the size-structure of red coral populations and, since climate-driven whole colony mortality is size-dependent in the red coral (with smaller colonies being more susceptible, Fig. 4b), a compensatory effect of MPAs against climatic perturbations could be expected. However, our results show that MPAs provided only a very weak compensatory effect. This slows down the rate of population declines along a thermal stress gradient, but does so only under low to moderate frequencies of mass mortality events (MME) (Figure 5 & 6). When recurrent MME occurred at high frequencies, MPAs had a little buffering effect on the long-term persistence of coral populations. To date, only few

studies provide data over relevant time-periods to assess the potential role of MPAs at buffering climatic impacts. Michelli and colleagues (2012) showed that, in a temperate abalone species, MPAs enhanced population resilience to climatic disturbances by increasing the presence of larger individuals that drove a major reproductive output as well as larvae export potential. Similarly, in tropical coral reefs, Mellin and colleagues (2016) reported a strong effect of MPAs at enhancing resilience of benthic communities to external perturbations, although the mechanisms underlying these patterns were not demonstrated.

In contrast, Graham and colleagues (2008) found that reefs under local protection had been more impacted by El Niño bleaching event in 1998. More recently, an unprecedented mass bleaching event occurred at the Great Barrier Reef, with more than 90% of the reefs affected irrespective of their level of local protection (Hughes et al., 2017). Our results add support to the latter

findings in showing that local protection could be effective at lower levels of thermal stress, but have weak to nonexistent compensatory effects under potential future scenarios of increased ocean warming. Also, this study highlights the role of demographic approaches to provide a mechanistic understanding on how multiscale stressors interact to shape the dynamics of reef corals and other habitat-forming species, which is urgently needed to anticipate and manage threats driven by a rapid environmental change.

Conclusions

Long-term studies provide a unique window to understand and predict the responses of marine populations and communities to multiple stressors. This study demonstrates that MPAs are a good tool to enhance the structural dynamics of economically and ecologically important species such as the red coral. However, warming represents a new threat originated at global scale that can drive either subtle or

more abrupt declines that will result in the “extinction debt” of these key habitat-forming species (Hughes et al. 2013). While the red coral and other long-lived habitat-forming species have persisted millennia of human pressures, novel climatic threats may lead to unanticipated and massive consequences for the integrity of temperate reefs. Unfortunately, local protection may not be enough to ensure the persistence of shallow populations under future scenarios of higher frequency and intensity of mass mortality events. A rapid and drastic reduction of greenhouse gas emissions seems the only effective way to prevent the widespread extinction of shallow red coral populations and other long-lived marine invertebrates. In the context of a variety of multi-scale stressors, any conservation action aimed to preserve the functional role and ensure the persistence of benthic communities will need to consider conservation actions at both local and global scales.

ACKNOWLEDGEMENTS

This study was partially funded by the Spanish Ministry of Economy and Innovation Biorock project (CTM2009-08045), Smart project (CGL2012-32194), TOTAL Foundation Perfect Project, Prince Albert II of Monaco Foundation MIMOSA project and the European Union’s Horizon 2020 research and innovation programme under grant agreement No 689518 (MERCES). This output reflects only the author’s view and the European Union cannot be held responsible for any use that may be made of the information contained therein. IMS was supported by a FPI grant (BES-2013-066150), CL by a Ramon y Cajal (RyC-2011-08134), and JBL by a Postdoctoral grant (SFRH/BPD/74400/2010) from the Portuguese Foundation for Science and Technology. Support for DFD came from National Science Foundation awards 1340024 and 1242355. IMS, CL, JBL and JG are part of the Marine Conservation research group (2017 SGR 1521) from the Generalitat de Catalunya.

CHAPTER IV

Accounting for life-history strategies and timescales in marine restoration

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INTRODUCTION

Marine coastal ecosystems host high levels of biodiversity and provide goods and services to a large proportion of the world's human population (Palumbi et al. 2008). The cumulative effects of multiple stressors such as overfishing, habitat destruction, and pollution together with new global threats (i.e. climate change, biological invasions) have driven compositional changes, local extinctions, and wholesale destruction of many benthic communities (Jackson et al. 2001, Airoidi & Beck 2007). To face this challenge, actions at both global (i. e. reduction of greenhouse gas emissions) and local levels are urgently needed (Kennedy et al. 2013). At local scales, fishery regulations and marine protected areas can help to reduce or remove threats (Edgar et al. 2014). Even so, when the resilience of natural systems has been seriously diminished, active restoration may be necessary as a complementary tool to restore damaged populations

and communities (Possingham et al. 2015).

Over the last few decades, the success of ecological restoration efforts in terrestrial landscapes has improved dramatically, with successful examples of enhancing ecosystem structure, function, and the provision of ecosystem services (Benayas et al. 2009, Perrings et al. 2015). In the marine realm, however, restoration approaches have generally been successful only at very small spatial scales and continue to present many challenges (Edwards & Gomez 2007). Strikingly, the degree of success in marine restoration actions is not generally related to the underlying costs of the project (Bayraktarov et al. 2016). This is partly due to the high methodological constraints, but also to relatively poor understanding of the drivers underlying successful actions.

Habitat-forming species such as corals and seagrasses have been the primary targets of marine restoration activities, and transplanting

asexually produced units (i.e. coral fragments or seagrass shoots) has been proposed as the tool of choice for recovering habitats by bypassing sensitive early life stages (Edwards & Clark 1998). Recently, research has shown that corals display high survival rates after transplanting when compared to the dominant organisms found in seagrass beds, oyster reefs, or saltmarsh ecosystems (Bayraktarov et al. 2016). However, this generalization ignores the high diversity in life-history strategies of the dominant species in these benthic communities (Darling et al. 2012; Madin et al. 2014). Indeed, life-history tradeoffs between demographic rates have been observed in hard coral species, suggesting potential effects on short-term and long-term restoration success (Edwards & Clark 1998; Dizon & Yap 2006, Glassom & Chadwick 2006). Yet, quantitative evidence of how the life history of target species shapes restoration outcomes is lacking in the scientific literature. To advance the theoretical

framework of marine restoration and provide tools to enhance the effectiveness of transplantation efforts, we need to go beyond habitat type towards a fuller quantitative analysis of how life-history strategies determine the best strategies or allow better prediction of the speed and eventual success of restoration efforts.

To date, most studies of transplant success in marine systems have focused on survival rates of transplanted individuals over relatively short monitoring periods (usually less than 2 years) (Bayraktarov et al. 2016). However, a broad goal of restoration efforts is to recover structural complexity that can provide ecosystem services at rates similar to natural ones. Thus, when planning restoration actions, managers should consider the factors affecting the time required from any transplantation action to reach the restoration goals for the target species and habitat. Long-term monitoring programs can provide suitable data to inform this

issue, but funding often constrains the duration of monitoring after restoration actions and experiments (Lindenmayer & Likens 2009, Precht & Robbart 2006). Demographic modeling methods such as matrix and integral projection models (Morris & Doak 2002, Ellner & Rees 2006) can be used to synthesize individual data into predictions of the longer-term development of transplanted populations (Linares et al. 2008).

In the present study, we combined demographic monitoring of transplanted and natural of colonies of a temperate coral species, a comprehensive literature review of tradeoffs in the life histories of sessile marine species, and the use of population projection models to explore the dynamics of transplant efforts targeting species with different life-histories. Our results support the utility of explicitly linking life-history theory to marine restoration and provide an illustrative example of anticipating

the expected dynamics and timescales of restored ecosystems.

METHODS

Study system

The precious red coral *Corallium rubrum* is a structural octocoral of a highly diverse coralligenous assemblage of the Mediterranean Sea and also possesses important cultural and economic value. Due to historical overexploitation, most shallow populations of *C. rubrum* can be considered functionally impaired and many are ecologically extinct (Bruckner 2009, Tsounis et al. 2010). To reverse this situation, an international agreement urged Mediterranean countries to strengthen their *C. rubrum* fishery regulations during the last decade (Cau 2013). Unfortunately, the lack of enforcement of regulations on coral harvesting along with poaching is widespread across the Mediterranean basin and represents a major problem for the management of the species,

hindering the effectiveness of its conservation (Linares et al. 2012).

Study area and transplant experiment

In 2011, the Catalan authorities intercepted 14.5 kg of illegally harvested *C. rubrum* along the Montgrí Coast (Catalonia, Spain). About 300 red coral colonies, a small portion of the intercepted colonies, were selected for a transplant experiment. These colonies were initially kept at 16 °C and fed in aquarium facilities at the Institute of Marine Sciences in Barcelona (Spain). After one week, the colonies were transported in coolers to the Parc Natural del

Montgrí, Illes Medes i Baix Ter in the NW Mediterranean and transplanted onto a rocky wall ranging from 15 to 17m depth using a two-component epoxy putty as glue. The site was chosen because some sparse red colonies were found in the vicinity, indicating its suitability for the species (Fig. 1).

Demographic traits

Four transects were established within the transplanted population and surveyed through photographic sampling after transplantation, in May 2011, and again in May 2015 (Fig. 1). Survival rates of the transplanted colonies were quantified by individually

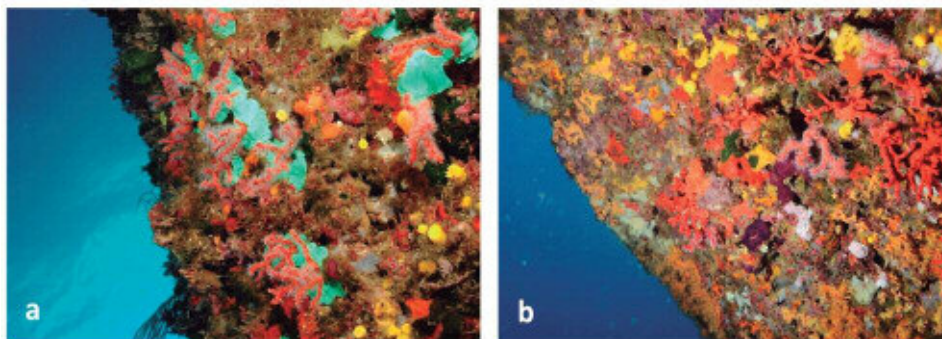


Figure 1. Restoration of *Corallium rubrum* populations. (a) A new population was transplanted in 2011; (b) most transplanted colonies survived in 2015, after four years of transplantation.

identifying coral colonies from the photographic series from 2011 and 2015. Natural survival rates of *C. rubrum* colonies were calculated from long-term data on 8 natural populations (see Supplementary Methods for a complete description of surveys). Reproductive potential of colonies was estimated for a sample of transplanted colonies ($n = 35$) outside the monitored transects and from a natural adjacent population ($n = 35$) in late June of 2015 by counting *C. rubrum* larvae found inside the polyps of the fertile female colonies (Tsounis *et al.* 2006). Samples were collected by SCUBA diving and fixed in 4% formaldehyde. At the lab, 15 polyps per sample were dissected and larvae found inside the polyps were counted.

Literature review

We explored life history tradeoffs in marine restoration experiments following two steps. First, we systematically reviewed all transplantation experiments of marine sessile species we could

identify in a search of the literature up to November 2015. Using *Google Scholar* we searched for a combination of the terms “restoration”, “transplantation”, or “rehabilitation” with a second term related to marine sessile taxa: “coral”, “gorgonian”, “sponge”, “macroalga” or “seagrass”. We then selected those studies that conducted experimental transplants as a restoration technique and reported survival rates during at least one year after transplanting.

We also compiled data on growth rates of sessile marine species since this vital, or demographic, rate is highly correlated to overall life history (Darling *et al.* 2012). We searched available studies reporting standard data on linear extension rates to approximate average species-specific growth rates for corals (Madin *et al.* 2016). In seagrasses, mean horizontal rhizome elongation rates were used as an indicator of the species growth rate (see Marbà & Duarte 1998), thus,

seagrass and sessile invertebrates were analyzed separately.

Demographic projections

Red Coral. To synthesize data on growth, survival and reproduction into predictions of population growth and increasing sizes within populations, we used Integral Projection Models (IPMs) parameterized with long-term demographic data from several natural red coral populations (full description data analysis and model construction are in the Supplementary Material S1). Based on annual IPMs, we computed 1000 stochastic projections assuming that all annual models can occur with equal probability at each time-step. Maximum height of transplanted colonies was measured in 2015 using photogrammetric techniques (Drap et al. 2013) and the distribution of heights was used to establish the starting population vector for the projections.

Linares et al. (2010) argue that the structural complexity of *C. rubrum*

populations can be assessed by quantifying the proportion of large colonies (>100 mm), since these larger colonies provide structural complexity. Based on this parameter, we compared the outputs from our population projections to the proportion of large colonies in three relatively un-impacted *C. rubrum* populations that are located within old and well-enforced Mediterranean MPAs (Fig. 1 & S1, Linares et al. 2010).

Comparative Analyses. We also searched in the literature for published matrix population models of other marine sessile species. We then used these models to perform deterministic population projections of 100 individuals starting at the smaller size class and computing time periods until the population reached a proportion of large individuals (largest size class) equivalent to the 20% and 80% of the expected proportion when reaching the Stable Stage Distribution (SSD).

RESULTS

Demographic traits of *C. rubrum* transplanted colonies

After four years, 99.1% of transplanted *C. rubrum* colonies were still alive. Annual survival rates of transplanted colonies did not show significant differences from control populations (Fig. 2a). Transplanted colonies also had similar reproductive potential to colonies in natural populations, considering both the proportion of fertile colonies as well as the frequency of larvae per polyp (Fig. 2b, 2c).

Comparative survival and growth in transplant experiments/actions

We found 50 studies that allow calculation of mean annual survival rates after at least one year following transplanting for a total of 59 marine structural species (Fig. 3a). These included 40 species of hexacorals, which have a mean annual survival of 60.8% (range of 6.8 to 98.6%); 5 species of

gorgonians, including the present study, with mean annual survival of 48.1 (range of 30.0 to 99.1%); 1 species of sponge, with mean annual survival of 85.7%; 11 species of seagrasses, with mean annual

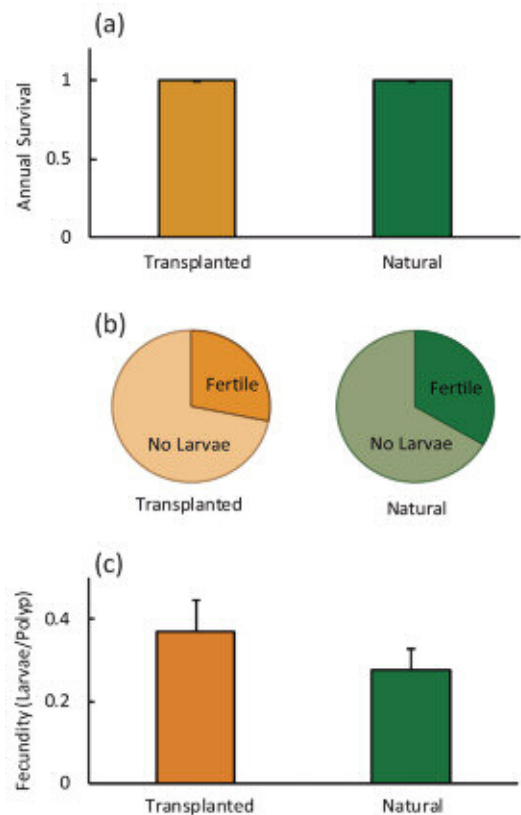


Figure 2. Demographic traits in transplanted and natural *C. rubrum* populations. (a) Mean annual survival rates; (b) Proportion of fertile colonies; (c) Mean polyp fecundity, calculated as the frequency of larvae found per polyp within fertile colonies.

survival of 42.5% (range of 28.9 to 69.2%) and 2 seaweeds, with mean annual survival of 43.1% (range of 25.1% to 80.0%). We observed a significant negative correlation between survival after transplantation and the species mean

growth rates measured under natural conditions in marine sessile invertebrates (Fig. 3b; $n = 35$; Pearson's $r = 0.47$, $P = 0.005$; Spearman $\rho = 0.37$, $P = 0.046$). Seagrass species revealed a parallel pattern (Fig. 3c; $n = 8$; Spearman

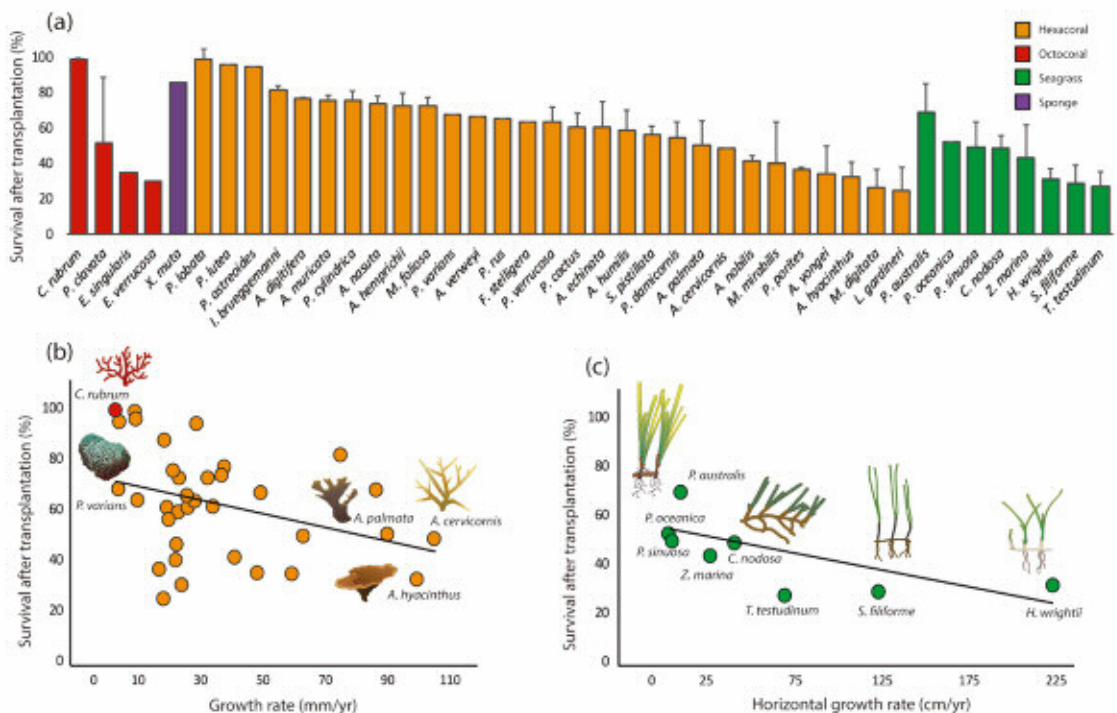


Figure 3. Survival rate of marine sessile species in transplant experiments. (a) Mean annual survival rates (Mean \pm SE). (b) Life history tradeoff between survival after transplantation and growth rates in 35 marine sessile invertebrate species and (c) Life history tradeoff between survival after transplantation and growth rates in 8 seagrass species. Each dot represents a species for which mean annual survival after transplantation and mean growth rate could be calculated from a range of published studies (see Table S1). In seagrass species, growth represents mean horizontal rhizome elongation rate (see Marbà & Duarte 1998). Images: Integration and Application Network, University of Maryland Center for Environmental Science (ian.umces.edu/symbols/).

$\rho = 0.81$, $P = 0.022$), although the relationship was only marginally significant according to Pearson's correlation (Pearson's $r = -0.69$; $P = 0.059$). Growth data measured in natural and transplanted colonies for coral species were also highly correlated (Fig S3, $n = 17$; Pearson's $r = 0.85$; $P < 0.001$).

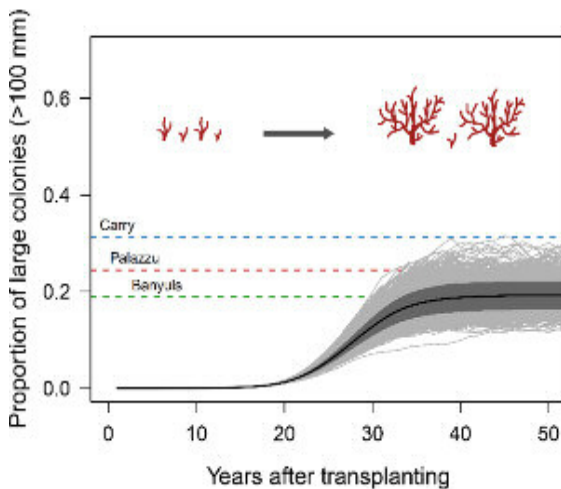


Figure 4. (a) Predicted temporal dynamics of the *C. rubrum* population size frequency distribution. Black line represents the mean and shaded area the confidence intervals of 1000 stochastic projections. The size frequency distribution of three natural and well protected *C. rubrum* populations (dotted lines) were used as an ecological baseline (Linares *et al.* 2010).

Demographic projections and recovery periods

Red coral. The transplanted population in 2015 was dominated by small individuals (most red coral colonies were < 35 mm in height, Fig. 1 & S1), while natural red coral colonies had extremely low growth rates (Fig. S2). The stochastic integral projection models incorporating these traits showed that a period ranging from 30 to 40 years after transplanting is needed for populations to have a proportion of large colonies comparable to that seen in the well-preserved *C. rubrum* populations used as an ecological reference (Fig 4).

Comparative analyses. The simulated recovery periods for 41 marine sessile species were highly variable in length, ranging from years to several decades (Fig 5, Table S3). The expected recovery length was strongly and positive associated with the species' mean survival rate regardless of the conservation goal ($n = 41$; 20% threshold: $R^2 = 0.419$; $P < 0.001$;

80% threshold: $R^2 = 0.495$; $P < 0.001$). After accounting for potential artifacts due to different matrix dimensions, mean survival rates were still a strong predictor of the expected recovery periods (Table S1).

DISCUSSION

Marine restoration is a relatively young discipline with most efforts only operating at very small spatial scales (Bayraktarov et al. 2016). Filling knowledge gaps on the processes underlying restoration success is therefore crucial to help

further develop this field and ensure meaningful planning and success over larger spatial and temporal scales. In this study, we quantify the role of life history in shaping restoration outcomes and demonstrate a consistent tradeoff between survival and growth across different taxa with contrasting life-history and functional traits, which in turn drives a tradeoff between required minimal transplantation effort at the start of a project and the maximum possible speed of ecosystem recovery.

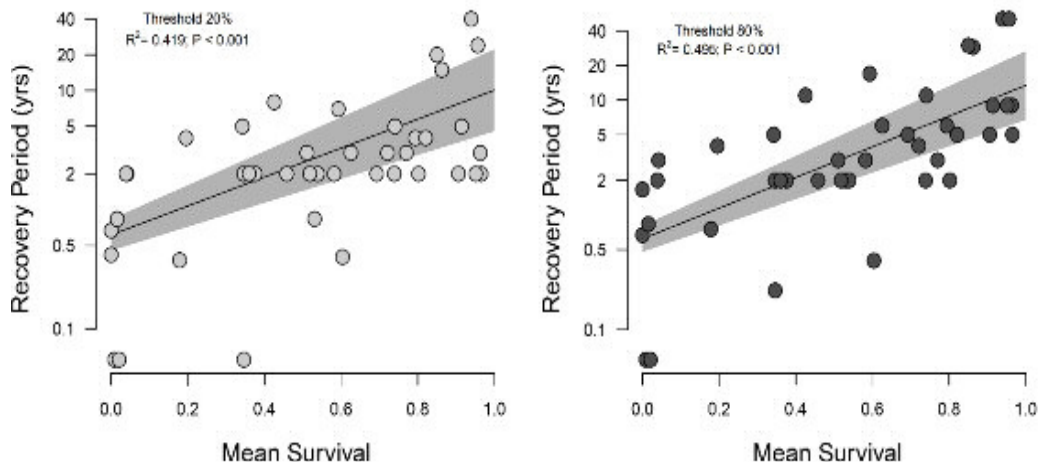


Figure 5. Projected recovery times for 41 marine sessile invertebrate species using published matrix populations models (Table S3) and setting a recovery threshold of (a) 20% and (b) 80% of the number of large colonies expected in a population at the Stable Stage Distribution. The black lines and the shaded areas correspond to the mean and standard errors of the linear models (Table S1).

Anticipating mortality patterns after transplantation is central to the design of any restoration action since it may determine the initial attaching effort required to achieve specific conservation goals. Here, a systematic review of transplantation experiments from tropical and temperate habitat-forming species revealed a negative tradeoff between growth and survival after transplantation that was supported despite differences in experimental techniques and physical properties of the environment that were not explored (Fig. 3b). Slow growing massive hard corals such as *Porites astreoides* and *P. lutea* and the sponge *X. muta* showed the highest survival after transplantation, with rates ranging from 86% to 98%. On the contrary, fast-growing corals such as *Acropora cervicornis*, *A. yongei* and *A. palmata* had survival rates that ranged from 35% to 44%. These results were consistent with previous transplant experiments in tropical coral species with contrasting life histories (Edward & Clark 1989, Dizon & Yap 2006).

Similarly, among seagrasses, the three slow-growing *Posidonia* species showed higher shoot survival after transplantation (from 49% to 69%) compared to faster-growing *Syringodium filiforme* (29%) and *Halodule wrightii* (27%) (Fig. 3c). Our findings agree with allocation theory, which predicts that tradeoffs between vital rates such as growth, reproduction and survival may arise from energetic constraints acting at physiological levels (Stearns 1989). Further, branching morphologies associated to faster life histories may increase exposure to physical damage and result in higher mortality rates (Madin et al. 2014). There are also a number of external drivers that can strongly influence restoration success such as predation and herbivory, density of transplants, and catastrophic events (Shaish et al. 2010, Gomez et al. 2014). In spite of the clear importance of these effects, our results show that species' life histories can still provide strong predictive power concerning the outcome of transplantation projects.

Better understanding of both the intrinsic and extrinsic drivers of mortality patterns after transplantation would be ideal and could lead to the implementation of more successful restoration designs, since this combined approach can better define both anticipated time periods for restoration and also the relative benefits of direct transplantation effort.

There have been major calls worldwide to ban the international trade in precious coral and to implement management regulation aimed to ensure the conservation of these species (Bruckner 2014). Yet, the feasibility of restoration actions for these emblematic species has remained uncertain and this may hinder the potential for development of future restoration plans. In the present study, we observed that the colonies of the octocoral *Corallium rubrum* were extremely resistant to the stress of transplantation, displaying high survival rates similar to those in natural populations (Fig. 2a, Garrabou &

Harmelin 2002). It is remarkable to observe this high survival rate in transplanted *C. rubrum* colonies that were subject to the stresses of being harvested, kept out of the water in the poachers' nets, transported, maintained in aquaria for one week, and then transplanted back into natural habitat. Yet these transplanted *C. rubrum* colonies had a similar proportion of fertile colonies and even higher frequency of larvae per polyp after 4 years than observed for colonies in natural populations (Fig. 2). Assessing reproductive potential is also critical when working with most marine sessile species which, like *C. rubrum*, show limited larval dispersal and high self-recruitment rates (Ledoux et al. 2010). Indeed, to effectively recuperate populations through a single transplantation effort, newly restored populations must also be viable in the long term, with reproduction reaching natural rates. Here, the high survival and reproductive potential displayed by transplanted *C. rubrum* confirmed the potential success of this

restoration action and strongly support the feasibility of these techniques, at least at local spatial scales, with potential applications for other long-lived precious coral species.

As important as choosing a suitable species and restoration method is considering the appropriate time scale and ecological baselines over which to evaluate restoration outcomes or to expect the restoration of ecological functions (Bull et al. 2014). Stochastic projections developed revealed that periods ranging from 30 to 40 years may be necessary for newly established *C. rubrum* populations to show a colony size distribution comparable to those observed in well-preserved natural populations (Linares et al. 2010). These results suggest that, similar to relatively fast-growing terrestrial forest systems (Vesk et al. 2008), long-lived coral stands can take up to several decades to recover their functionality and to allow the development of associated organisms, such as fish and

invertebrates, as may occur in tropical coral reefs (e.g., Cabaitan et al. 2008). More interestingly, we found that potential recovery periods can be accurately predicted by the specific mean survival, demonstrating the strong influence of the species' life histories on the temporal scales associated to restoration actions (Fig. 5).

Overall, this study demonstrates a tradeoff between initial transplantation effort needed to achieve a target density of individuals and the speed of recover that may be achieved in a restoration action. For instance, targeting fast-growing species such as *A. cervicornis* or *A. hyacinthus* (with survival rates ranging from 40 to 50%), will require a two-fold to three-fold initial amount of attached colonies to obtain the same density of survivors compared to actions targeting slow-growing resistant species such as the red coral *C. rubrum* or the massive coral *Porites lutea*. On the other hand, life histories of the target species will

also have a strong effect on the expected recovery periods that may vary as much as 20 to 30 yrs (Fig. 5). Finally, because life-history and functional traits are highly correlated (Adler et al. 2014), favoring specific strategies can have long-term consequences for habitat complexity and ecosystem responses to global change (Ortiz et al. 2014).

ACKNOWLEDGEMENTS

The authors thank the Agents Rurals de Catalunya for their invaluable work against red coral poaching. We are indebted to A Lorente for his support. We thank E Aspillaga and L Navarro for field assistance. P Capdevila, A Gori, K Kaplan, A Medrano, and two anonymous reviewers provided valuable comments on the manuscript. Funding was provided by the Spanish MINECO (CTM2009-08045, CGL2012-32194), the Oak Foundation, the TOTAL Foundation Perfect Project and the European Union's Horizon 2020 research and innovation programme under grant agreement No 689518 (MERCES). This output reflects only the author's view and the European Union cannot be held responsible for any use that may be made of the

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CHAPTER V

Predicting depth-mediated impacts of ocean warming at the end 21st century in Mediterranean coastal ecosystems

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INTRODUCTION

Anthropogenic global change threatens the structure, function and resilience of world's marine ecosystems (Hoegh-Guldberg & Bruno 2010, Ainsworth et al. 2016). Widespread changes in the distribution and population dynamics of marine organisms have been linked to recent warming (Perry et al. 2005, Pinsky et al. 2013), with negative consequences for both marine biodiversity and the provision of ecosystem services (Barange et al. 2014). In this context of rapid environmental change, understanding what shapes distribution ranges and, more importantly, how climate change will affect the biogeography and vertical range dynamics of structural marine species is central to designing sound adaptive conservation strategies.

Poleward distribution and elevation range shifts are well documented in terrestrial landscapes (Chen et al. 2011). In the marine realm, increasing evidences show how

mobile organisms such as fish and invertebrates are tracking temperature trends over broad spatial scales (Pinsky et al. 2013, Montero-Serra et al. 2015). Importantly, strength and speed of responses to warming are mediated by functional traits (Sunday et al. 2015) and habitat requirements (Rutteford et al. 2015). Despite having a paramount role in coastal ecosystems, there still is a large uncertainty on whether structural marine species will track changing thermal habitats. Besides the lack of motility and long generation times, several environmental factors may constrain poleward movements for sessile species, including light limitation (Coma & Gili 2000, Muir et al. 2015), availability of suitable habitats (Rutteford et al. 2015), and large-scale geographic barriers (for example in enclosed seas such is the Mediterranean Sea). When environmental barriers prevent these poleward movements (Muir et al. 2015), widespread mortalities at shallow habitats may drive bathymetrical range contractions as a response to warming impacts

(Bridge et al. 2013, Linares et al. 2014, Muir et al. 2017).

Species distribution models (SDM) are a powerful tool to understand and predict the effects of global change on species geographic ranges (Pearson & Dawson 2003, Hijman et al. 2006, Rutteford et al. 2015). Yet, this field is still in its infancy for marine species, partly due to methodological constraints that lead to a relative scarcity of suitable data (Robinson et al. 2011). Oceans are inherently three-dimensional complex systems with strong environmental gradients at fine spatial scales. In spite of these features, a common assumption when modeling marine species distributions is that environmental parameters remain constant across depths (Sbrocco et al. 2013, Parravicini et al. 2015). Surface environmental conditions may be meaningful for shallow occurring species. However, they can be misleading when modeling potential warming impacts because they are often depth-dependent (Garrabou et al. 2001, Mumby et al. 2001, Bridge

et. 2013, Muir et al. 2017). Exploring how depth mediates potential impacts of ongoing climate change in benthic ecosystems is thus a current challenge in conservation. The recent development of novel oceanographic models that include the third dimension (e. g. Macías et al. 2015), opens new opportunities to advance our predicting ability to explore potential warming impacts along depth gradients.

Temperate benthic ecosystems host great levels of marine biodiversity and provide goods and services to dense populated coastal communities. However, temperate regions are warming at faster rates than global averages and have been considered hotspots of present and future climate change (Oliver et al. 2018). The Mediterranean Sea is an example of such overlapping patterns: hosts high levels of biodiversity and provides ecosystem services to millions of people from 21 countries (Coll et al. 2010; NOAA 2015). Mediterranean coastal habitats have been historically damaged by multiple

local and regional stressors such as overfishing, habitat destruction, pollution and invasive species (Lejeusne et al. 2010, Claudet & Fraschetti 2010). Recently, several heat waves linked to ongoing global change impacted benthic communities causing catastrophic mass mortalities over broad spatial scales (Cerrano et al. 2001, Coma et al. 2009, Garrabou et al. 2001, 2009). The frequency and intensity of these extreme climatic events is predicted to increase in the near future (Oliver et al. 2018). It is thus essential to fully understand and predict potential warming impacts on Mediterranean benthic communities to advance towards large-scale and cross-boundary sound adaptive conservation strategies (Game et al. 2008, Michelli et al. 2013, Mazor et al. 2015).

Here, we took advantage of a recently developed 3D hydrodynamic-biogeochemical coupled model of the entire Mediterranean Sea that provides predictions at different depth layers

(Macías et al. 2015). We explored how present and future climate shapes the bathymetrical dynamics of two emblematic habitat-forming species across the Mediterranean Sea: the red gorgonian, *Paramuricea clavata*, and the red coral, *Corallium rubrum*. Firstly, we assessed the role of local thermal regimens and biogeochemical factors in shaping the upper range limits of gorgonian populations. Then, we applied logistic generalized additive models at different depths (from 5 to 55m) to predict present and future vertical patterns of habitat suitability. Our results revealed large-scale negative impacts of climate change and challenge current predictive modeling approaches based on surface temperature data by revealing complex patterns of differential warming impacts along depth gradients across the Mediterranean Sea.

METHODS

Model species

We focused on two dominant habitat-forming gorgonians of Mediterranean hard bottoms: the red gorgonian *Paramuricea clavata* and the precious red coral *Corallium rubrum*. They are slow-growing and long-lived suspension feeders with an important ecological role providing structural complexity in Mediterranean mesophotic reefs (Ponti et al. 2014). They display slow population dynamics driven by high adult survival and low reproduction success (Linares et al. 2007, Montero-Serra et al. 2018), which results in a high vulnerability to human-driven perturbation such as overfishing and climate change (Garrabou et al. 2009, Linares & Doak 2010). The precious *C. rubrum* is threatened by legal and illegal fisheries (Tsounis et al. 2010, Linares et al. 2012, Montero-Serra et al. 2015). During the past three decades, several mass-mortality events linked to climate change have impacted benthic communities

over broad spatial scales (Cerrano et al. 2001, Garrabou et al. 2001, 2009). These events seem depth-dependent (Garrabou et al. 2001) and are putting at risk some shallow gorgonian populations (Linares & Doak 2010).

Modeling philosophy

Mediterranean gorgonians distribution and abundance are largely influenced by the availability of hard bottoms and other environmental conditions such as the irradiance levels, temperature, exposure to flow and food availability (Weinberg 1979, 1980, Zabala & Ballesteros 1989). Unfortunately, we still lack of comprehensive fine-scale cartography for Mediterranean seafloors features. Thus, exploring the biogeography of these species without considering the availability of rocky bottoms can seriously compromise the validity of predictions. Therefore, rather than predicting the distribution of our model species across geographic space, we restricted our predictive

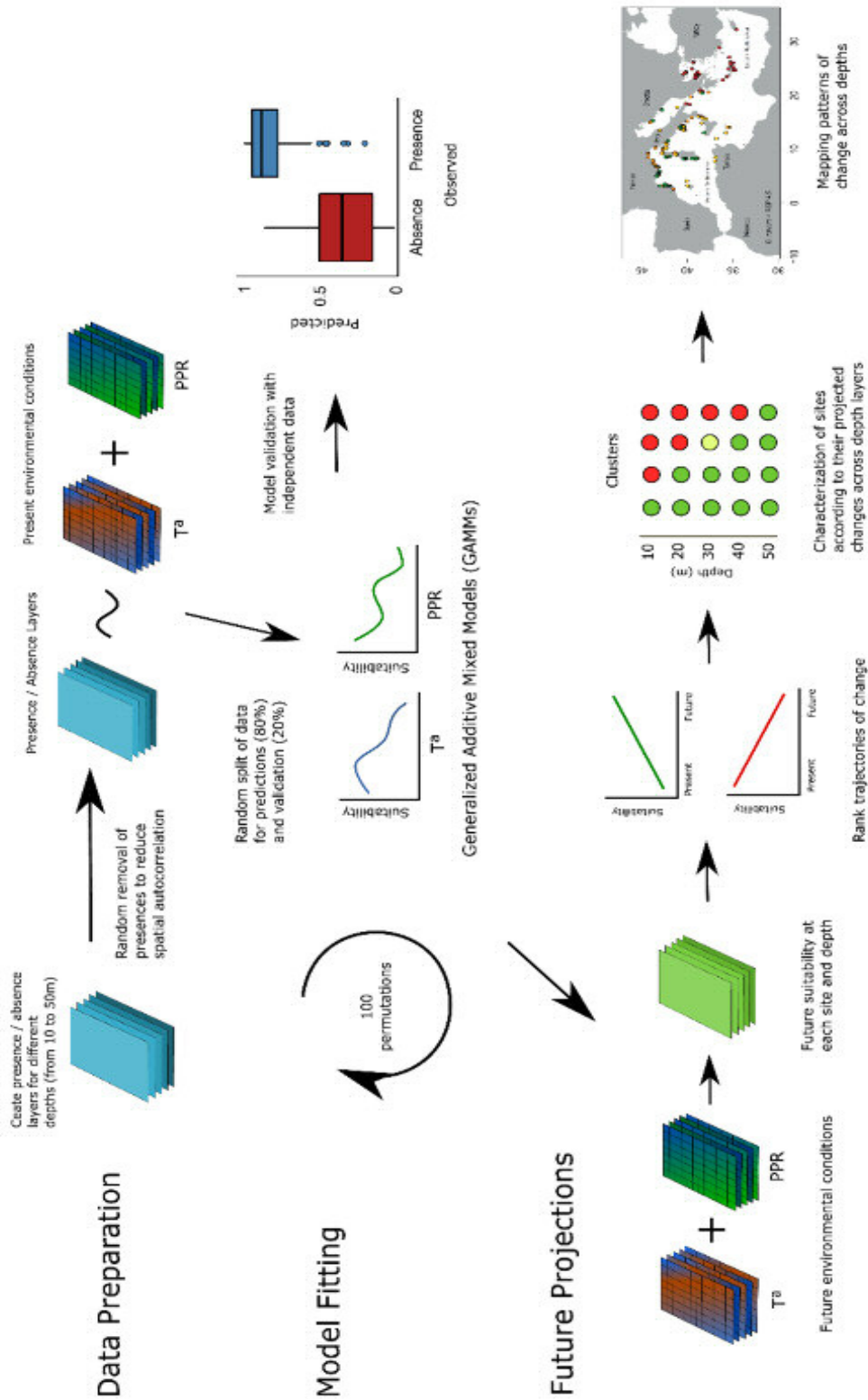


Figure 1. Modeling framework to predict the effects of climate change at different depths.

modeling approach to known locations. Based on a large dataset of reported occurrences and absences at different depth layers, from 5 to 55m, we fitted Generalized Additive Mixed Models (GAMMs) to link bathymetrical distribution of gorgonians to current environmental conditions (water temperature, organic and inorganic nutrients). We then used these trained models to forecast changes in habitat suitability conditions across depth layers within locations where gorgonians are present (Fig. 1). Additive models make no a priori assumptions about the nature of associations between predictors and response variables and have been used to assess the importance of different environmental drivers on patterns of distributions and relative abundance in marine ecosystems (Rutteford et al. 2015).

Biological data

Data on the horizontal and bathymetrical distribution of *Corallium rubrum* and *Paramuricea clavata* were obtained from surveys

carried out by two citizen science programs: *SeaWatchers*, *ReefCheck Italy* and *Cigesmed*. Briefly, each survey included name and geographic coordinates, date and time, underwater visibility, survey depth range (min and max), and observation effort in terms of dedicated time. Additionally, a comprehensive literature review was conducted to compile all available occurrence data from published papers of *C. rubrum* and *P. clavata* (Table S1).

Ocean Model

A 3D General Estuarine Transport Model (GETM) was used to simulate the hydrodynamics in the Mediterranean Sea (Macías et al. 2015, Macías et al. 2018). A detailed description of the GETM equations could be found in Stips et al. (2004). The predictive ocean model has a horizontal resolution of 5'×5' and includes 25 vertical layers. The bathymetric grid used was constructed based on topographic data from ETOPO1 (NOAA,

www.ngdc.noaa.gov/mgg/global/). From this model, monthly temperatures, nutrient (phosphate and nitrate) concentrations and primary productivity were extracted. In the present study, we used two long-term model runs covering the periods 2000–2013 and 2085–2099 were used. Model runs were based on two different emissions scenarios according to the Fifth Assessment Report of the Intergovernmental Panel on Climate Change (IPCC AR5): the business as usual RCP8.5, in which CO₂ emissions will continue to grow at current rates, and the more optimistic RCP4.5, which includes a strong slow-down in the current CO₂ emissions' rate.

Data filtering and pre-analyses

Geographic distribution data was classified into three categories. We first removed duplicated data by randomly deleting all but one overlapping geographic data points. To account for data gaps on environmental parameters along the coastlines, we performed a spatial interpolation called “Inverse

Distance Weighted (IDW)”. The IDW technique computes an average value for unsampled locations using values from nearby weighted locations. The weights are proportional to the proximity of the sampled points to the unsampled location and can be specified by the IDW power coefficient (*idw()* function from the package “gstat” in R).

Exploring environmental drivers of upper bathymetrical limits

We subset the dataset to only include sites were the upper range limit of the population was known. Then, a set of linear models were fit to explore correlations with local environmental variables and upper range limits of both *C. rubrum* and *P. clavata*. Among the predictors, we included temperature and biogeochemical variables (phosphate, nitrate and primary productivity) at different depths (from 0 to 50m). Best fit models were selected using Akaike Information Criterion corrected for small sample sizes (AICc).

Modeling the relative probability of occurrence at different depth layers

A set of logistic generalized additive mixed effects models (GAMMs) were fitted to presence and absence data of *P. clavata* and *C. rubrum* populations at different depth layers from 5 to 55m deep (5-15m; 15-25m; 25-35m; 35-45m; 45-55m). These models included potential combinations of environmental predictors of both biotic and abiotic variables under current conditions (extracted from the ocean model during the period 1997 to 2012), topographic features and depth layer as fixed factor. In addition, site-specific coordinates were included in the models as a random effect in order to account for repeated measures within each site (non-independence of samples).

In a first exploratory analysis, we tested a set of potential environmental variables (temperature, nitrates, phosphates, primary productivity and topographic features) at driving the

observed presences at each depth. To fit the models under present environmental conditions, we included the mean values of environmental predictors for the period from 1997 to 2012 at different temporal scales (annual mean, annual maximum, August maximum, September maximum). We also included topographic variables (slope, concavity, NS and EW aspect) extracted from MARSPEC (www.marspec.org) (Sbrocco & Barber 2013). Then, we kept only the most explanatory variables (annual mean temperature, August temperature, mean annual primary productivity, and slope) (Table S6).

We retained the best predictors (annual and august mean water temperatures and annual mean primary productivity) to fit GAMMs and predict the current and future occurrence of gorgonians at different depth layers. We performed 100 permutations as follows: first, to account for spatial autocorrelation we randomly selected only one data point for each

cell in a pre-defined grid of $0.1^\circ \times 0.1^\circ$ (Longitude / Latitude). Second, the dataset was further split into randomly selected 80% of data for model training and 20% for independent validations. Third, we fitted GAMMs using the training data and environmental variables under present conditions (see fitted GAMMs in Fig. 3), and we used these models to predict present and future probability of occurrence at each depth layer for 2100 according to two socio-economic scenarios (RCP4.5 and 8.5). Model outputs of the 100 permutations were averaged to obtain a single present and future suitability value for each site and depth. Finally, we used the remaining 20% of data for independent validations, comparing predicted occurrences at different depth layers to observed presences and absences (see schematic workflow in Fig. 1).

Classification of sites according to their predicted trajectories of change

To summarize the expected patterns of change across depths at each location, we ranked expected trajectories of change using a numerical index framework that indicates sign and strength of expected change across depths. Firstly, we categorized the present and future habitat suitability for each depth layer within locations in three levels: low (from 0% to 33%), medium (from 33% to 66%), or high (from 66% to 100%) (Fig. 4). Based on this classification, we then characterized the sign and strength of expected change from present to future conditions at each depth within sites. We rank suitability changes with numerical indexes from (-3) for large decreases to (+3) for large increases (Fig. 5). Briefly, sites that went from bad to good conditions received positive values from (+2 or +3); sites that went from good to bad conditions received negative values (-2 or -3), and sites that kept constant

conditions received (+1) if they were good, (0) if they kept constantly medium, or (-1) if they showed constantly low probability of occurrence (see Fig. 5). Finally, a k-means cluster analysis was

applied to the indexes of change for the 5 depth layers at all sites to identify the most common and informative patterns of change (Fig. 6).

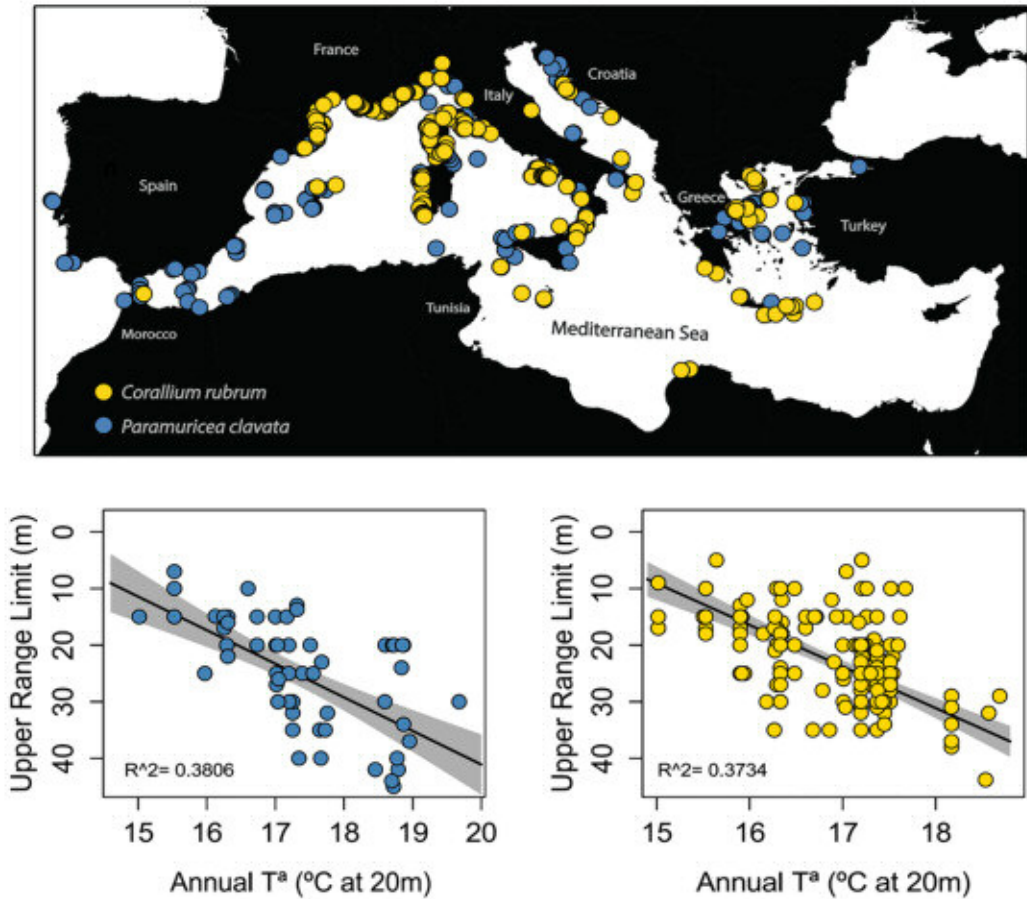


Figure 2. Study area and environmental drivers of the upper range dynamics for *P. clavata* and *C. rubrum*. (Top) Geographic distribution of *Paramuricea clavata* (blue) and *Corallium rubrum* (yellow). (Bottom) Relationship between upper range limits and local temperatures for *P. clavata* (left) and *C. rubrum* (right).

RESULTS

Current drivers of upper bathymetric limits

The red gorgonian *Paramuricea clavata* and precious red coral *Corallium rubrum* are extensively distributed across the entire Mediterranean Sea and adjacent Atlantic waters. More interestingly, upper range limit (i. e. shallowest occurrence) of both gorgonians were negatively correlated to local water temperature, showing a general pattern across the Mediterranean Sea, from frequent shallow presences in the generally cold-warm water NW Mediterranean Sea to deeper upper ranges in the warmer Eastern basin. A set of linear models including multiple potential environmental drivers

(temperature and biogeochemical variables) revealed that mean annual temperature at 20m was the best predictor of the upper range limit for both *C. rubrum* and *P. clavata* (Fig. 2; Table 5). On the other hand, primary productivity also showed a significant positive correlation to upper range of both species (Table S5). Some exceptions were observed in the Western basin such as the warmer and more oligotrophic Balearic Islands, where gorgonian populations are only found from 35m or deeper waters. Overall, upper range limits of both species were strongly driven by local thermal regimes.

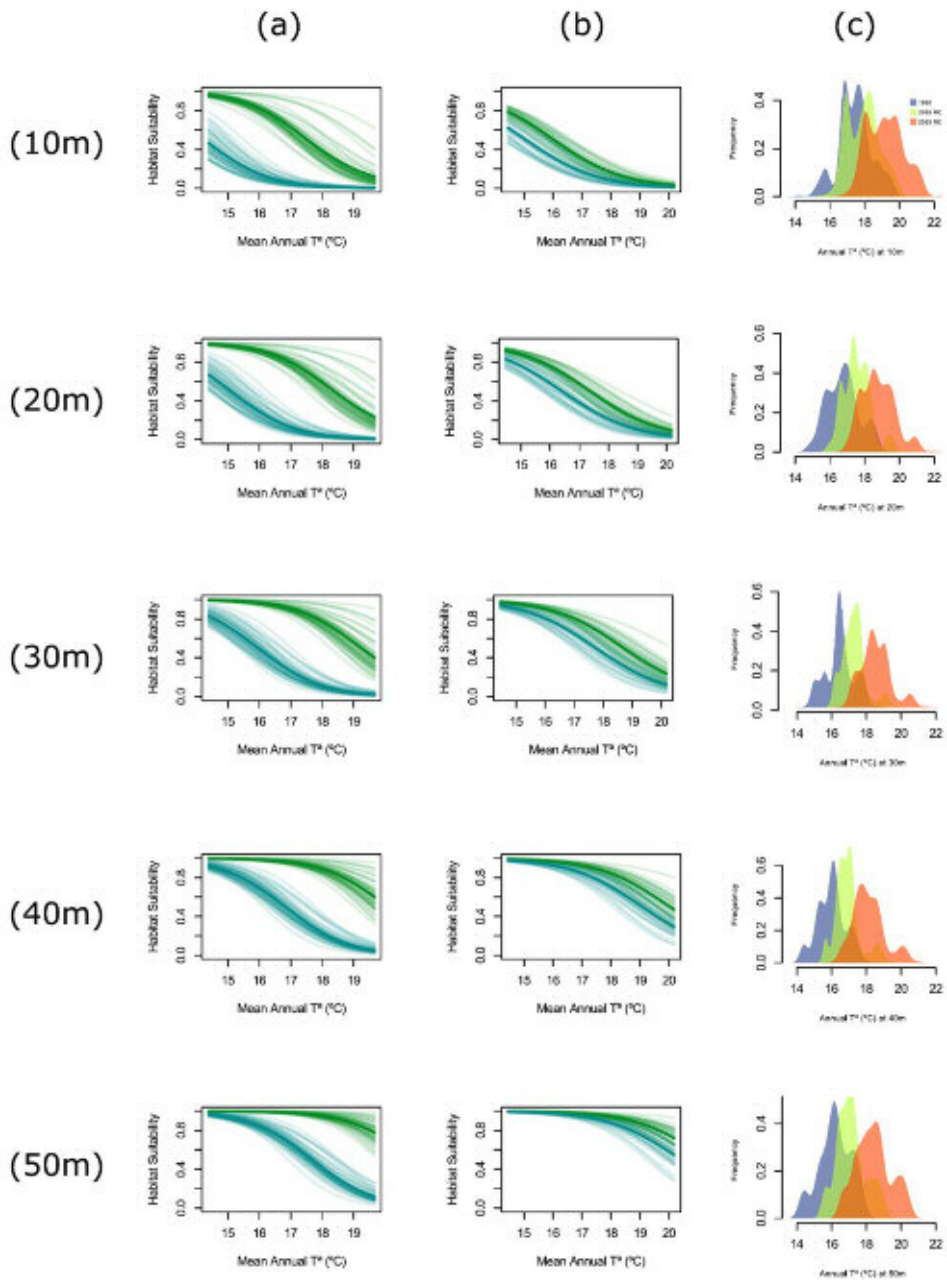


Figure 3. Logistic generalized additive mixed models (GAMMs) describing gorgonian's probability of occurrence at different depth layers depending on local mean annual temperature and primary productivity. Each row indicates a depth layer (from 5-15m, 15-25m; 25-35m; 35-45m; 45-55m). First and second columns show the modeled presence probability according to local temperature under high and low primary productivity conditions for *Corallium rubrum* (left) and *Paramuricea clavata* (center). The third column (right) show the density function of local temperature at the study sites under present conditions (blue), future optimistic RCP 4.5 scenario (green), and future business-as-usual RCP 8.5 scenario (red).

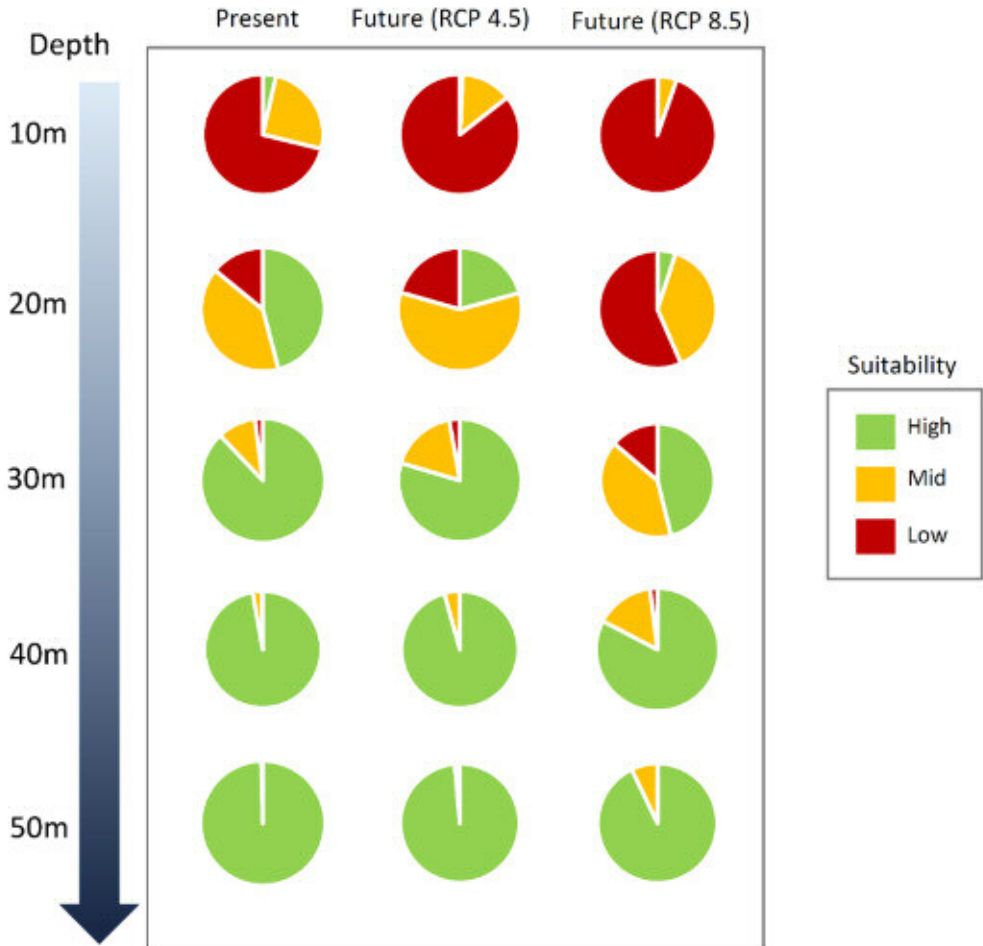


Figure 4. Present and future predicted probability of occurrence of gorgonians at the study sites across the Mediterranean Sea at different depths. Probability of occurrence was classified as low (from 0 to 33%), medium (from 33 to 66%), and high (from 66 to 100%).

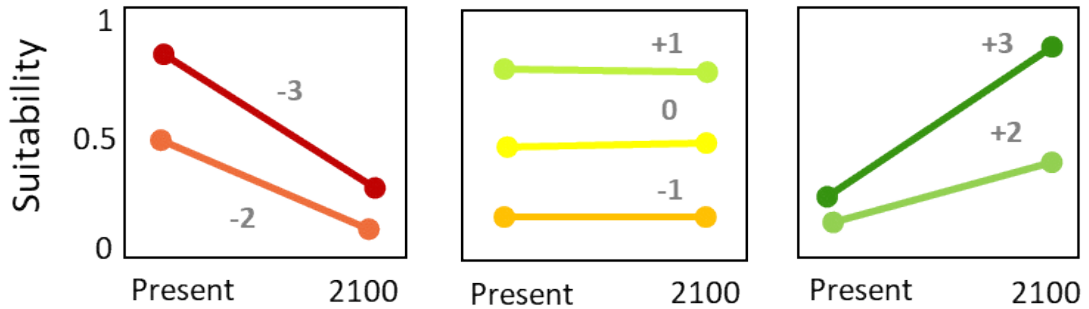


Figure 5. Potential trajectories of change according to present and future climate conditions. We classified each site and depth according to their expected rate of change. The first plot (left) shows positive trajectories; the second (center) show sites where no change is expected; the third plot (right) show sites that will experience negative changes (from good to medium or bad conditions).

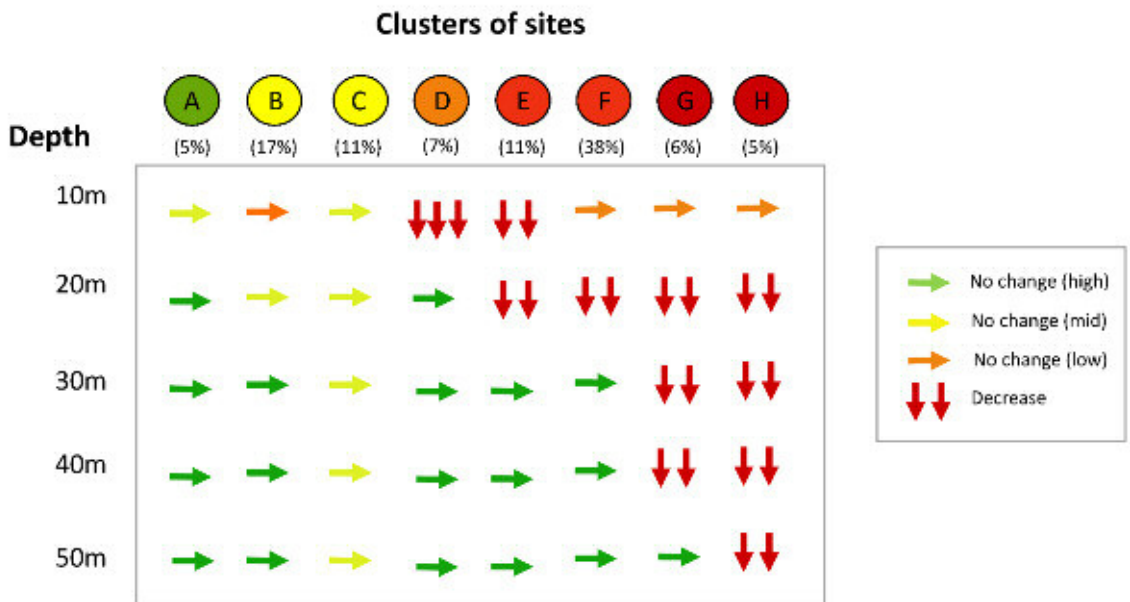


Figure 6. Patterns of expected warming impacts along a depth gradient. (A) K-means cluster analysis for sites according to their expected patterns of change across depth layers. Horizontal arrows indicate no change and down-pointing arrows indicate negative changes.

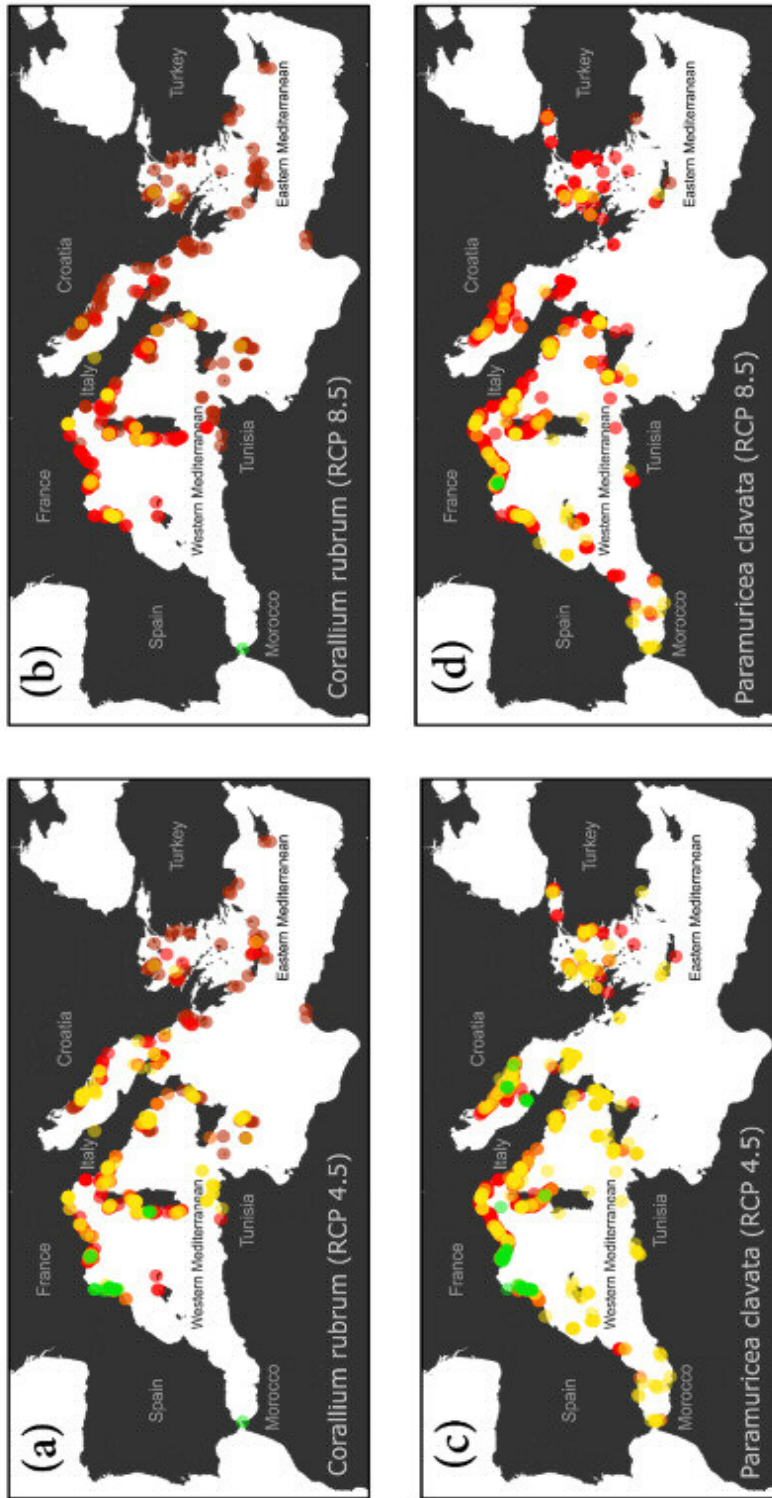


Figure 7. Geographic distribution of sites according to their expected patterns of change across depth layers. Colors correspond to clusters described in the top panel. Maps represent sites with known presence of *Corallium rubrum* (top) and *Paramuricea clavata* (bottom) under RCP4.5 (left) and RCP8.5 (right) emissions scenarios.

Projected impacts of climate change

Predictions for the end of the 21st century revealed extensive negative changes in probability of occurrence across the Mediterranean Sea. Overall, the frequency and intensity of changes appears strongly dependent on the CO₂ emission scenario and depth. Under the “business as usual” emission scenario (RCP8.5), almost all high suitability sites disappeared at shallow waters. Although less extensive, warming will cause impacts at deeper waters. We also observe broad regional differences: the proportion of sites that are predicted to become unsuitable at the end of the century was lower at deeper depths in the Western basin, but extensive in the Eastern Mediterranean Sea.

When looking at trajectories of change across sites, we found different patterns of change across depths (Fig. 6). Overall 33% of sites are expected to keep similar thermal conditions. From these, relatively

few will keep high thermal suitability (Cluster A), and the frequency of these sites sharply decreases depending of emissions scenario (from 149 sites in RCP4.5 to only 5 sites in RCP8.5). Some sites that had already low suitability in shallow waters (up to 25m) under present conditions will kept consistent in the future (Cluster B and C). Other clusters included sites that experienced negative impacts of warming but differed among them on the bathymetrical extend of the expected impacts. Sites included in clusters D were strongly impacted only in very shallow waters (up to 15m). Clusters E and F included sites in which shallow waters were already non-suitable under present conditions but will experience further decreases in thermal suitability in deeper habitats up to 35m. Finally, clusters G and H are expected to experience the deepest impacts of climate change, including sites that had bad conditions at shallow waters under the present climate and will experience further negative changes

in the thermal suitability up to 55m (Fig. 6 & Table S7).

When looking at the spatial arrangement of these clusters, we observe a gradient from frequent deep impacts at the Eastern Mediterranean Sea and heterogeneous patterns of warming impacts in the Western Mediterranean. There, future predictions reveal widespread decreases in habitat suitability due to warming, but these impacts will strongly depend on depth, basin (Western vs. Eastern), and IPCC emissions scenario. In the warmer Eastern Mediterranean basin, climatic impacts will generally reach deeper areas, while in the Western Mediterranean we expect highly heterogeneous patterns of changes across space and depth. Some areas that may experience negative changes at shallow depths, other will be affected up to 45m and 55m. On the other hand, there will be few climatic refugees, defined here by sites predicted to preserve high suitability at all depths. These sites are mainly found along in the

NW Mediterranean along the coastline of Catalonia and France, in proximity of the generally cold-water masses of Gulf of Lions.

DISCUSSION

Understanding and predicting how warming impacts will spread across horizontal and vertical distributions of marine species is a major challenge in conservation yet partially unsolved. Here, using a novel three-dimension ocean model and extensive vertical distribution data of two dominant habitat-forming gorgonians across the Mediterranean Sea, we demonstrate a crucial role of considering depth when predicting future impacts of climate change in coastal marine ecosystems.

Current drivers of upper bathymetric limits

Our results demonstrate that upper bathymetric limits of *Paramuricea clavata* and *Corallium rubrum* are shaped by local environment across the Mediterranean Sea (Fig. 2). Upper edges were negatively

correlated to local temperature and positively to primary productivity over broad spatial scales (Table S5). Previous physiological microcosm experiments provide mechanistic support of these results by showing that warm temperatures combined to shortages in food supply can be detrimental for the survival of both *P. clavata* and *C. rubrum* under controlled conditions (Torrents *et al.* 2005, Coma *et al.* 2009, Crisci *et al.* 2017). Therefore, populations may be strongly limited by warm and nutrient-depleted waters that remain stratified above the thermocline at the end of summer periods (Bensoussan *et al.* 2014). Indeed, several field studies have linked water temperatures above approximately 24-25°C for a several days with catastrophic mass mortality events on both gorgonians up to certain depths (Cerrano *et al.* 2000, Garrabou *et al.* 2001, 2009; Linares *et al.* 2014). Despite having less explanatory power in our models, primary productivity was also positively correlated to the presence of shallow occurring populations (Table S5). Besides the

role in food supply for the survival of these suspension feeders, primary productivity also controls light penetration across depth gradients, and this plays a key role in competition for substrate between benthic macroalgae and suspension feeders in Mediterranean rocky reefs (Zabala & Ballesteros 1989, Ballesteros 2006). Contrarily to what is generally observed in tropical coral reefs, temperate gorgonians commonly inhabit semi-dark habitats and their presence is enhanced when light availability is low due to higher primary productivity that reduces light penetration and allows outcompete macroalgae species that otherwise dominate benthic communities. Both parameters clearly play a role on the distribution of the bathymetrical dynamics of our model species. However, disentangling their relative contribution is challenging because of their high correlation across large spatial scales and will require further investigation through finer-scale analyses and manipulative experiments. Overall, these results

provide new insights into environmental controls of the bathymetrical dynamics for two dominant habitat-forming species in temperate reefs with important implications for our general understanding of potential effects of rapid environmental changes.

Predicting potential deepening of gorgonian populations

In this study, we used a novel predictive framework to forecast how changing temperatures and primary productivity patterns will impact Mediterranean gorgonian populations across depth gradients. Our models predict extensive impacts of 21st climate change that will strongly depend on emissions' scenario and depth. Under RCP8.5, most gorgonian populations are likely to experience shallow extinctions and bathymetrical range contractions (deepening of populations) (Fig. 4 & 7). The deepening of *P. clavata* population from 30 to 45m has been documented after recent warming events, suggesting that our

predictions of potential extinction of shallow ranges may be already occurring (Linares et al. 2014). In addition, our predictions agree with more general observations on decreasing warming-driven mortality with depth in Mediterranean gorgonians as well as in tropical reef corals (Garrabou et al. 2001, Mumby et al. 2001, Bridge et al. 2013, Muir et al. 2017). A general deepening of shallow occurring gorgonians would be catastrophic for benthic ecosystems and coastal societies in the Mediterranean Sea. On one hand, gorgonians enhance biodiversity (Ponti et al 2014) and, although not yet fully understood, may provide thermal buffering effects to other benthic species (Jurgens & Gaylord 2018). But also, in cases in which shallow-water gorgonian forests deepen their distribution beyond the recreational scuba diving limits (~40m), small local economies dependent on tourism can be negatively affected. Scuba diving and related tourism activities represent a growing economic sector in the Mediterranean Sea and

worldwide and have a key role in marine conservation by promoting the enthusiasm and support of great part of coastal societies that benefit either directly or indirectly from tourism (Badalamenti et al. 2000, Sorice et al. 2007).

Climatic refugees and complex depth-mediated impacts of warming

Identifying warming hotspots and potential climatic refugees is a challenging endeavor but essential to incorporate the global change dimension into spatial conservation planning. Here, we provide strong evidences that warming can differentially impact marine biodiversity at different depths, and these potential depth-mediated impacts should be considered when adapting the configuration of MPA networks to a warming ocean. There have been recent calls to include depth-related biodiversity gradients into large-scale spatial prioritizations (Levin et al. 2017, Venegas-Li et al. 2018). While ecologists often use surface temperatures for predicting

the negative consequences of future climate change, our results call for a carefully reconsideration of the vertical dynamics of habitat-forming species as well as the importance of three-dimension patterns of warming, since they may not be homogenous across large geographical areas (Levin et al. 2017). This shifting paradigm is essential anticipation of these complex patterns of warming will have major implications for the conservation of mesophotic reefs and may lead some important socio-economic negative consequences for coastal areas that depend upon the tourism industry.

Finally, major differences in extend and severity of warming impacts were predicted depending on the CO₂ emission scenario (Fig. 4 and 7). For instance, the observed climatic refugees in the NW Mediterranean Sea became drastically reduced under RCP8.5. these results highlight the importance of global political actions to drastically reduce emissions to ensure the conservation

of benthic marine ecosystems. While the role of local actions to increase the resilience of marine communities to global change is still unclear, our results clearly demonstrate a strong role of global emissions trends on the conservation of Mediterranean gorgonian forests.

Limitations

This study represents a crucial step forward in the way we conceive predictions of climate change impacts for coastal marine ecosystems. However, there still are important limitations of our modeling framework that must be acknowledged. First, we believe that the spatial resolution of the model (5 arc-minutes), despite being much finer than most of the environmental data currently used in predictive spatial models (Nadeu et al. 2017), is still too coarse to accurately reproduce the micro-climatic conditions experienced by individual gorgonian corals. Finer-scale predictive models validated and corrected by local temperature records are needed to improve our

predictive capacity. Indeed, the best fitted GAMMs for both species included annual mean temperatures. While annual averages are suitable for representing broad trends and highly correlate to frequency and severity of marine heat waves (Oliver et al. 2018), they could underestimate short-term high intense warming events. On the other hand, envelope models are a powerful tool to understand and project realized niches from a holistic perspective, but they fail to link environmental drivers to demographic mechanisms and interspecific interactions that drive the distributions of marine organisms, and this can potentially lead to misleading correlations between environmental predictors and the presence of the species (Fordham et al. 2018). These limitations can be addressed by a deep understanding of the species biology and natural history and a critical evaluation of environmental predictors used and their relationship to geographic data. Coupling correlative approaches to mechanistic modeling frameworks

that can explicitly link patterns to underlying processes would also aid improving model accuracy. For instance, hybrid models combining envelop models and thermal physiological thresholds from controlled experiments would represent a step forward to obtain more robust estimates and increase our predictive capacity of marine benthic dynamics in a changing world (Talluto et al. 2016). Despite these limitations, our models showed a relatively high predictive capacity in validations against independent. Overall, the projections presented here are unique in three aspects: (I) they have higher spatial resolution of the climate projections (5 arc min is a 20 times higher spatial resolution than most of previous works that commonly use $1 \times 1^\circ$ (van Hooijdonk et al. 2013, Couce et al. 2013). (II) The inclusion of the third dimension, which is inherent to marine ecosystems, provides a much more accurate and detailed picture of the ongoing and future warming-driven impacts to marine biota, and this is relevant to most coastal

marine ecosystems. (III) A novel use of multiple abiotic and biotic environmental variables, including primary productivity, which is very relevant for heterotrophic octocorals, and may play a crucial role in predictions of future global change scenarios for other dominant habitat-forming species such as seagrass meadows or kelp forests.

Conclusions

Our study predicts large-scale negative impacts of climate change and challenge current modeling approaches based on surface temperature data by revealing complex patterns of differential warming impacts depending on depth. These results should have important implications for the design of adaptive management strategies in the Mediterranean Sea. The dramatic change in the expected impacts depending on different socio-economic scenarios highlights the importance of global political decisions on the conservation of Mediterranean gorgonian populations and hence, all the

associated biodiversity. Finally, this study provides a novel approximation to better understand the consequences of warming along depth gradients with potential applications to other coastal marine ecosystems worldwide.

ACKNOWLEDGEMENTS

We thank the citizen science programs Reef Check Italy, SeaWatchers, and CIGESEMED for providing valuable geographic data. Comments by YM Bozek, JC Ortiz, N. Kruek, and K. Kaplan helped improving this manuscript. Funding was provided by the European Commission (MERCES), the Prince Alberto II Foundation (MIMOSA) and the Spanish Minister of Innovation (SMART).

General discussion

A current challenge for policymakers, managers and society is to counterbalance multiple threats to marine ecosystems and ensure the preservation of biodiversity and ecosystem functions for next generations. Habitat-forming species play a crucial role in benthic communities; unfortunately, they are experiencing large declines all over the world. Understanding how multiple stressors impair their resilience requires field monitoring efforts over large temporal and spatial scales, but gathering long-term data in marine ecosystems is challenging and often needs to involve large collaborative networks of scientists that can persist beyond the typical short time windows of most PhDs and funded research projects. Consequently, we generally lack high quality field data on benthic species at relevant scales, and this can hinder decision-making and the implementation of effective management strategies. This is

especially valid for the management of long-lived species, such as our model species, where the scale of the studies is far to be representative of their long lifespans. In the present thesis, we benefit from unique long-term demographic and geographic distribution datasets which we combined with innovative approaches using quantitative tools, principles of life-history theory, and meta-analyses to advance in the field of marine conservation in a context of rapid environmental change.

Through this thesis, we used the red coral, *Corallium rubrum*, as a study case to explore resilience patterns and drivers for long-lived octocorals, with the final goal of guiding conservation strategies for Mediterranean benthic habitats. We documented the dramatic impacts of fishing and climate change on different demographic processes and predicted their long-term consequences, estimated periods required to recover functional complexity of coral populations, and revealed potential benefits as well as limitations of local conservation

tools such as marine protected areas (MPAs) and active restorations actions. More generally, I provided novel applications of population and spatial modeling for conservation and climate change research and used meta-analyses to go beyond the model species to show general drivers and consequences of high longevity in marine benthic ecosystems.

Life-history theory: from evolutionary ecology to conservation

Life-history evolution

The role of the environment in shaping the evolution of life histories is a long-standing question in ecology and evolution that remains partially unresolved. In Chapter 1, we explored longevity patterns of marine habitat-forming

species, asking how predictable longevity is across depths and whether there are consistent consequences of individual longevity for population dynamics. We found strong positive relationships between depth and maximum lifespan across multiple sessile marine taxa, including corals, bivalves, sponges and macroalgae. Depth-related environmental gradients may exert an external forcing to the evolution of sessile life histories, which lead to highly predictable general distribution patterns. These findings have implications for our general understanding on drivers of life-history evolution in the marine realm but also provide some important insights into potential vulnerability of less known deep-sea benthic communities.

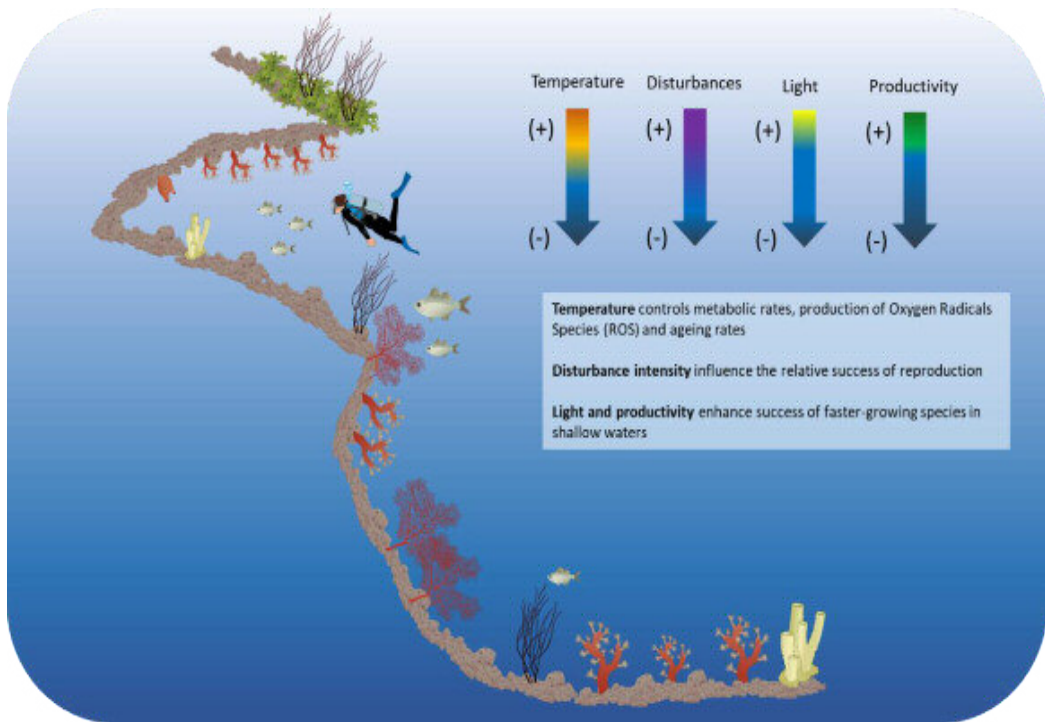


Figure 1. Conceptual diagram showing environmental drivers generally shaped by depth that may play a role in the dominance of long-lived species in deeper benthic habitats.

On one hand, our results suggest that local environmental variables that generally decrease along depth gradients - including temperature (both mean and variance), light penetration, physical disturbances (i. e. storms) and primary productivity - shapes evolutionary success of sessile species (Fig.1). Among several metrics commonly used in comparative demography, longevity is one of the most powerful

descriptors of life-history strategies: it positively correlates to survival and negatively to growth and reproductive rates (Salguero-Gomez et al. 2016, Fig. 2). A successful life-history strategy of long-lived species inhabiting deeper benthic habitats (which tend to be colder, less exposed to physical disturbances, and less productive) may thus be based on a disproportional allocation of

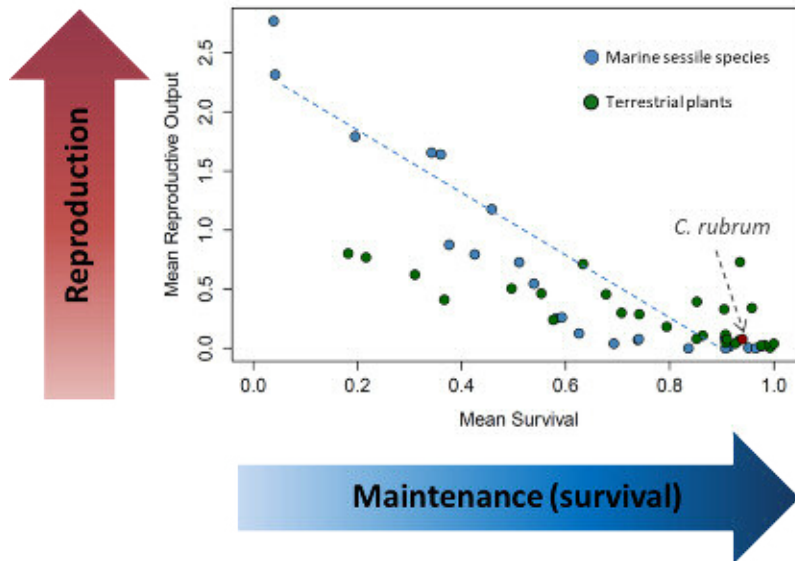


Figure 2. Life-history tradeoff between mean survival over an individual lifetime and mean reproductive output. Data were computed using matrix models from published demographic studies of a range of marine sessile species and terrestrial plants (see Table S10 in Chapter 1).

resources to physiological maintenance of cells and tissues, favoring individual survival over growth or reproduction. It is important to consider some important limitations of our global but basic comparison. Firstly, we used maximum depth occurrence as a metric to assess the overall bathymetrical range of the species, but it could be inaccurate for some species, especially for those dwelling in deep sea habitats. Secondly, intra-specific longevities can vary across latitudes as well as depth ranges. For instance, Pratched et al. (2015) showed that reef

building corals grow slower at deeper sites, but he also found strong correlations between coral growth and latitude. Thirdly, some notable exceptions must be noted to this general pattern of longevity and depth and they may provide interesting questions for further consideration. For instance, some highly longevous organisms such as seagrasses (p. e. *Posidonia oceanica*), inhabit shallow waters. Disproportionally large body-sizes and a great capacity to modify local habitats may aid these organisms to scape environmental pressures that generally favor short-lived species.

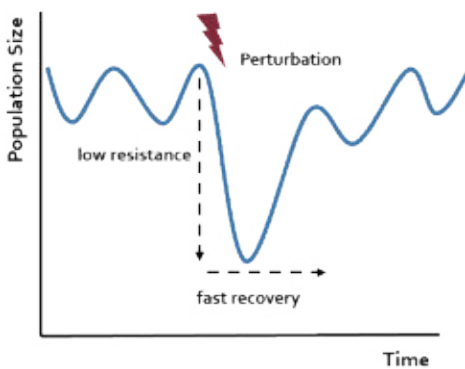
Finally, while this study represents an important step forward in comparative demography, our correlative approach does not allow for a mechanistic understanding of the observed patterns nor assessing the relative role and potential interactive effects of different environmental features at enhancing longevity of benthic species. These important questions should be addressed in detail by conducting finer-scale comparative studies or manipulative experiments that allow separating the effects of the different

potential drivers.

Life-history theory and conservation

In Chapter 1, we also demonstrate that high longevity drives high demographic stability across a range of marine sessile species and terrestrial plants. The underlying mechanism of this pattern is likely to be the general tradeoff between survival (which positive correlates to longevity) and reproductive success (Fig. 2).

Short-lived



Long-lived

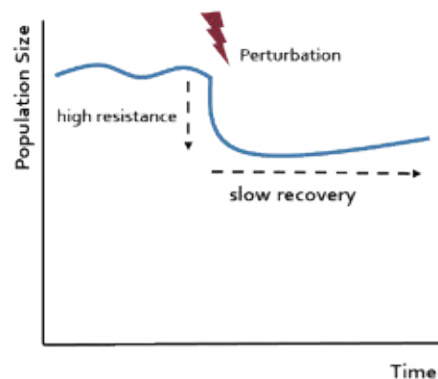


Figure 3. Conceptual model showing potential effects of life-history tradeoffs in two components of resilience at the population-level: resistance and recovery.

Life-history tradeoffs can play an important role underpinning resilience of populations and communities to recurrent high intensity disturbances. Resilience is a popular concept in natural and social sciences defined as the capacity of a system to withstand recurrent disturbances by resisting (avoiding change) or recovering (Côté & Darling 2010, Bernhardt & Leslie 2013, Darling & Cotte 2018). Theoretically, longer-lived species should show higher resistance to external perturbations and lower recovery capability due to a more limited reproductive success compared to shorter-lived species (Fig. 3). Our comparative analysis revealed low temporal demographic variability and low reproductive success in long-lived species, suggesting a relatively high resistance but low recovery potential due to limited recruitment. Overall, long-lived sessile species will tend to be low resilient to any external perturbation that causes high mortality events. However, resilience includes ecological processes at different levels of

biological organization, from populations and communities to ecosystems. In the present thesis, we narrowed this definition and focused on population-level resilience for long-lived species with highly restricted dispersal, in which recovery depends on local reproductive success (also called self-recruitment). For systems with higher dispersal potential, connectivity patterns and potential rescue effects will be crucial processes to consider when exploring resilience (Mumby & Hasting 2008).

From an applied perspective, our results predict a high vulnerability of long-lived species, which tend to be dominant in deep benthic communities, to external stressors that increase mortality such as direct and indirect human impacts of bottom fishing, deep-sea mining, oil spills, etc. (Van Dover et al. 2017, Danovaro et al. 2017). A recent study exploring recovery of deep-sea gorgonians after the Oil Spill of the Gulf of Mexico, supports these conclusions by showing that long-

lived gorgonians are highly vulnerable and their recovery can take hundreds of years (Girard et al. 2018). Their vulnerability arises from a naturally low resilience of these demographic stable communities. Therefore, evaluations of potential impacts of human activities that can damage deep-sea habitats should always include the precautionary principle, even in cases in which the affected benthic communities are not fully known (Danovaro et al. 2017).

On the other hand, in Chapter 4, we revealed that the tradeoffs among vital rates predicted by life-history theory are reflected in ecological restoration actions for habitat-forming species. A global meta-analysis demonstrated that slow-growing species tend to show higher survival after transplantation compared to fast-growing species. Thus, targeting faster-growing species may take larger transplantation efforts due to higher mortality of the manipulated individuals. On the other hand, using population models, we showed that

long-lived species with higher survival rates after transplantation may need much longer recovery periods to form fully functional populations. These results highlight the crucial role of life-history tradeoffs on the response of benthic species to transplantation and have implications for better designing and enhancing the overall effectiveness of ecological restoration efforts for marine benthic communities.

These examples add compelling evidence supporting the important role of life-history classifications to understand and predict the response of benthic communities to multiple stressors and conservation actions (Darling et al. 2012).

Quantitative tools in marine conservation: opportunities and limitations

Through the work presented in this thesis, we combined life-history classifications and quantitative tools to address conservation issues: anticipating expected recovery times (see Chapter 1) and the overall timescales that can be expected after

restoration actions for species with contrasted life-histories (Chapter 4), or exploring potential impacts of fishing and climate change at different temporal and spatial scales. Novel applications of quantitative tools such as integral projection models (IPMs) and general additive mixed models (GAMMs, for spatial analyses) have been successfully applied yielding new insights into the conservation of Mediterranean gorgonians such as assessing the role of MPAs and climate change in shaping population dynamics or predicting geographic and bathymetrical patterns of future climate change impacts. However, there are several limitations that must be considered when using modeling tools. In Table 1, we highlighted relevant benefits as well as limitations of the quantitative tools included in the present thesis: population models, spatial models and meta-analyses. One of the most important aspects to consider when using quantitative tools is the inherent tradeoff that exists between model complexity and transferability (Johnson & Lidström 2018). Under

some circumstances, basic models can represent a powerful tool to find generalities for a broad range of systems and provide important advances for the field. Simple models and processes can also be easier to communicate to a broader audience of non-specialists, including policymakers and society. However, it is essential to recognize that simplifying study systems comes to a cost of ignoring ecological complexity that exist in the real world. Indeed, in some cases, avoiding complexity could prevent to unravel the underlying drivers of observed patterns. For instance, in this dissertation, we used data from eight populations over relatively long-term periods to explore general drivers and resilience patterns for the red coral as a species. We tried to find broad generalities that can inform large-scale conservation initiatives, but it is important to keep in mind that these findings are somehow a simplification of real dynamics. The red coral is widely distributed along the Mediterranean Sea and living from shallow to deep habitats, likely

to experience slightly different environmental conditions than the populations studied here. In addition, our longest demographic dataset comprises 13 years of individual-level data. Although only rarely we find temporal scales in current marine ecological studies, it represents a tiny time window for a long-lived species such as the red coral that can live several hundreds of years. This means that, while the present work represents an important advance into sound principles for the conservation of this species, we must be cautious and recognize that there still some knowledge gaps and processes that may vary between and even within red coral populations. For instance, the responses to global warming can be shaped by adaptive processes that occur at the population-level (Ledoux et al. 2015) and may result in different sensitivity levels across geographic scales and depths (Torrents et al. 2005). Expanding demographic studies to cover broader spatial scales, including deeper habitats, is crucial to assess conservation status and trends of the

red coral and over its entire bathymetric range. Importantly, according to our findings, deeper dwelling red coral populations may be even more vulnerable to external stressors. The fast development of Remotely Operated Underwater Vehicles (ROVs) provides new opportunities to investigate deeper habitats (Rossi et al. 2008). Unfortunately, using these technologies is still very expensive and they rarely allow gathering high-quality individual-level demographic data (but see Girard et al. 2018 for an illustrative example of a fine demographic analysis on gorgonian populations dwelling at more than 1000m depth). Account for indirect or delayed effects of warming using more complex models is another aspect that should be addressed to advance our predictive capacity in the context of global change. For instance, when analyzing the recurrent impacts of climate change on coral populations, for simplicity, ecologists often consider only immediate whole-colony mortality rates due to acute thermal stress, ignoring any

potential delayed effect of these events. However, in the case of the red coral, we observed that warming also leads to increased mortality probability of affected colonies some years after the event. Thus, more accurate demographic analysis should model not only the immediate mortality due to warming but also any potential delayed effect on mortality of colonies that were partially affected. Increased complexity in demographic models may allow addressing novel questions but it can also make studies overly data-demanding or hinder their general transferability. Balancing these two aspects is a critical decision that must be thoughtfully taken considering all potential implications (logistics and data availability, temporal and spatial variance in measured traits and specific ecological questions).

Table 1. Benefits and limitations of quantitative tools used in the present thesis.

Quantitative Tool	Benefits	Limitations
Population Models	<p>Projections beyond the temporal scales covered by the field period to unravel subtle trends and to estimate extinction risk.</p> <p>Quantify relative contribution of each vital rate to the population growth, which informs about evolutionary drivers and conservation priorities.</p> <p>Explore responses of populations under different management scenarios and hypothetical changes in exposure to stressors.</p> <p>Estimate recovery periods after management interventions such as MPAs designation, fishing closures or restoration actions.</p> <p>Integrate field data into a standardized model that allows deriving life-history descriptors and systematic demographic comparisons.</p>	<p>Assume that field data is representative of current temporal and spatial variability. This is challenging for long-lived species.</p> <p>Assumes environmental conditions to remain constant in the future. Unlike under the present context of climate change.</p> <p>Ignores potential inter- and intra-population variability in life-history traits and vulnerability</p> <p>Accurately quantifying key processes such as larvae production, settlement and recruitment is challenging in marine ecosystems.</p> <p>In high dispersal species, modeling recruitment is more challenging and may require increased complexity and more assumptions.</p>
Spatial Models	<p>Explore species distribution ranges and realized niches.</p> <p>Identify environmental parameters that control species distribution boundaries.</p> <p>Forecast current and future habitat suitability according to different climatic scenarios.</p>	<p>Lack of fine-scale topographic and climatic data that can accurately reproduce micro-habitat features experienced by individuals.</p> <p>Potential biases in data collection and lack of systematic surveys.</p> <p>Correlative approaches may not capture real drivers such as interspecific interactions.</p>
Meta-analyses	<p>Compare species-specific responses of species with divergent life-histories to management interventions to find generalities and advance in our predictive power.</p> <p>Describe life-history changes along ecological gradients to infer potential evolutionary drivers.</p> <p>Assess global patterns in conservation issues.</p>	<p>Data quality can be variable and difficult to assess.</p> <p>Potential biases in reporting results can lead to misleading conclusions.</p> <p>Phylogenies and non-independent sampling issues when comparing species.</p>

Box 1 Global perspectives on MPAs and habitat-forming species

In this thesis, I assessed the role of local protection and climate change for an emblematic and overexploited coral in the Mediterranean Sea. MPAs can be effective at enhancing the functionality of red coral populations but our models suggest a limited role in buffering climatic impacts. Nonetheless, from a broader perspective, some crucial questions remain unresolved. Can we anticipate when and how MPAs will be most effective at protecting benthic habitats and enhancing their resilience to climate change? This crucial topic for marine conservation still generates intense debates in peer-reviewed journals, scientific conferences, and even in social media. In an exercise to synthesize current scientific evidences of MPA effects on habitat-forming marine species, I did a systematic review on Web of Knowledge using the keywords (“Marine Protected Areas” or “Marine Reserves” or “Local Protection” and “Corals” or “Benthic” or “Gorgonians” or Macroalgae” or “Seagrass”) and a deep search on GoogleScholar selecting the first 100 references. I found 78 relevant studies and classified them according to their observed effects (positive, null or negative) on different ecological parameters (benthic cover, density, size-structure, and species richness) to assess conservation status of habitat-forming species.

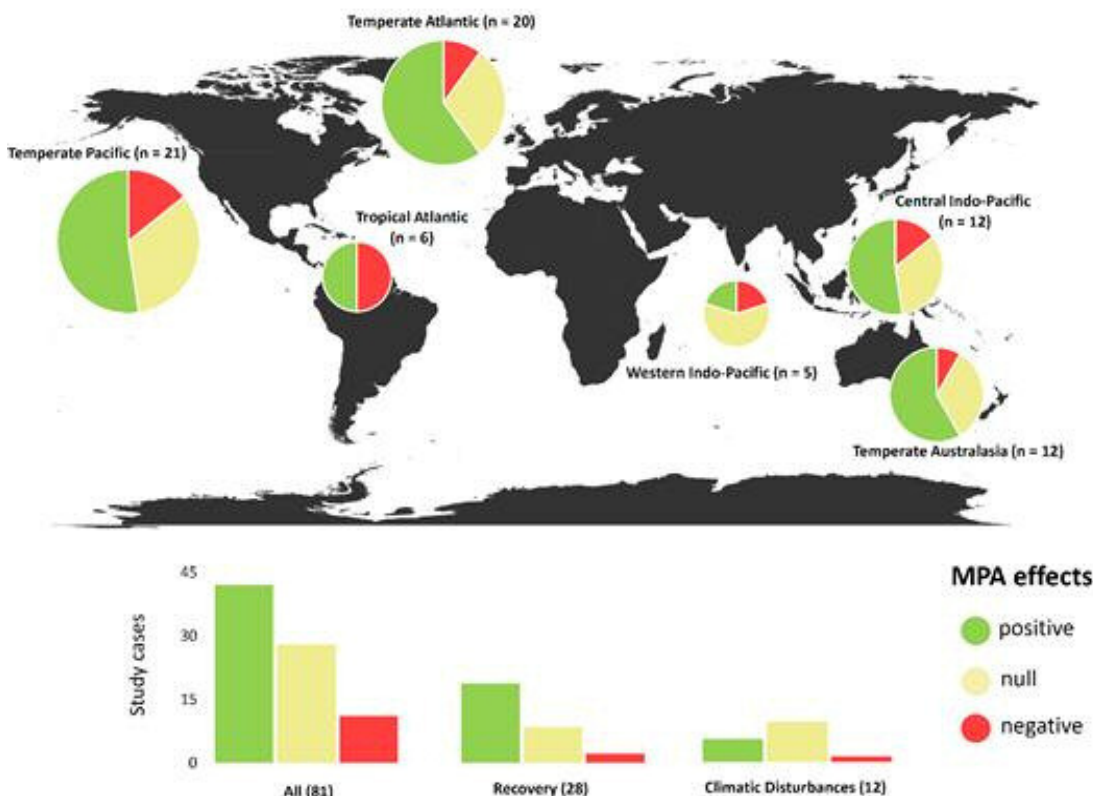


Figure 4. Global empirical evidences of marine protected areas effects on habitat-forming benthic species. A. Effects represented by global ecoregion according to classification by Spalding et al. (2007). B – D Show proportion of positive, null and negative effects in: all cases (B), only studies that look at recovery patterns (C), and only studies that assessed responses to climatic perturbations (D).

Overall, in approximately half of the relevant studies, MPAs provided positive effects on benthic communities, and these broad patterns were consistent across different ecoregions (Fig. 4). Interestingly, when looking only at studies that assessed recovery processes, the proportion of positive effects becomes much higher. In contrast, when looking at studies that have assessed responses to climatic perturbations, positive effects are less frequent (Fig. 4). This shows that MPAs will not always benefit habitat-forming species and suggest that benthic community responses to local protection can be much more complex than what it is often assumed. They are likely to depend on the species-specific processes that are affected by local protection and complex inter-specific interactions (Fig. 5). For instance, while evidences for enhanced resistance in protected sites against climatic perturbations appears not well supported (Hughes et al. 2017a); a key process in ecosystem resilience such as recovery seems much more sensitive to MPA implementation (Mumby et al. 2007, Micheli et al. 2010). For this reason, when studying effectiveness of MPAs for habitat-forming species, we need to clearly define priory expectations and metrics of conservation success, linking them to specific ecological process that can be affected by protection. Although this is a basic preliminary exploration that will require finer analyses to account for potential confounding factors, it provides a novel overview of a highly relevant topic and opens new questions on the overall role of local protection at enhancing different processes involved in benthic ecosystems resilience.

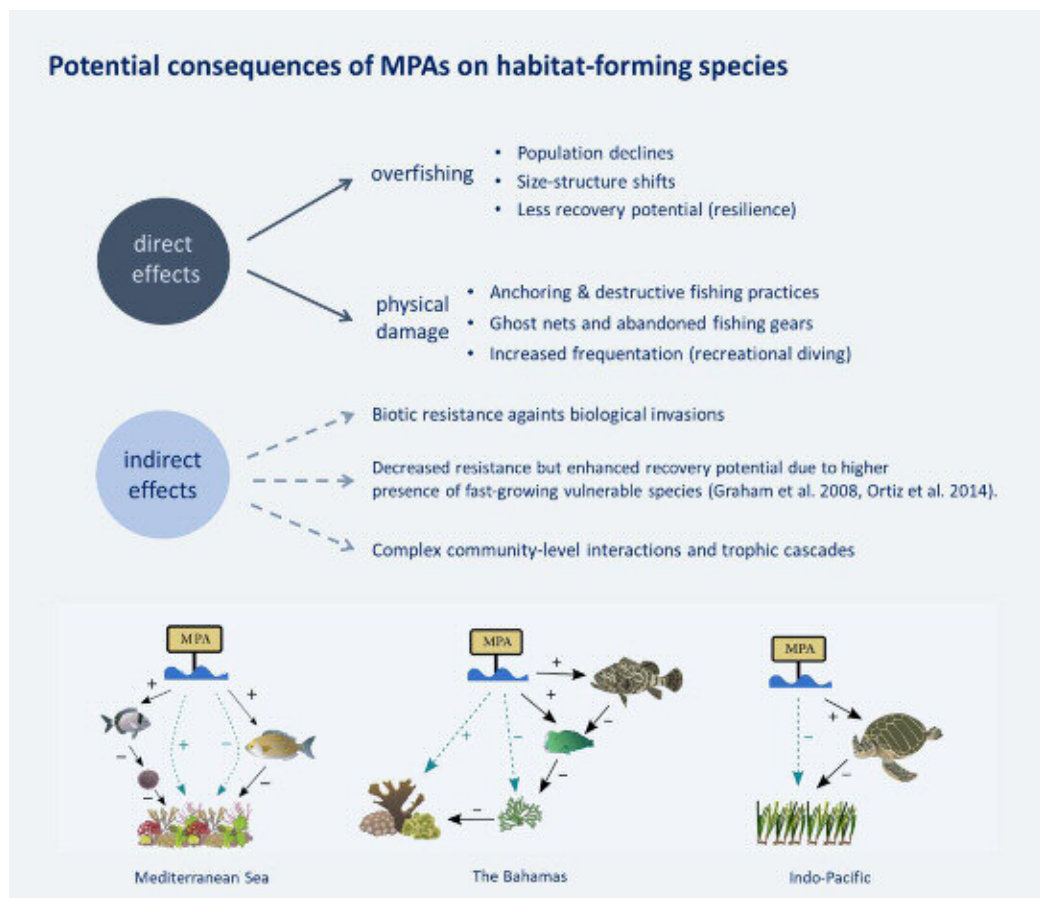


Figure 5. Processes involved in potential direct and indirect effects of marine protected areas (MPAs) on habitat-forming species.

Study case: Resilience of Mediterranean red coral populations to fishing and climate change

Precious corals are found in several seas across the world and have been collected for thousands of years for use as jewelry, homeopathic medicines, and cultural artifacts. They are vulnerable to depletion because of their slow life history, small and patchy populations, and have been subjected to “boom and bust” fishing strategies with a variety of fishing gears (Bruckner 2016). Today, there is a growing concern on their conservation due to historical unsustainable fisheries and novel threats associated global change (Bruckner 2013; Garrabou et al. 2009). In the Mediterranean, the red coral *Corallium rubrum* has been recently enlisted as Endangered in the Mediterranean Red List of the International Union for the Conservation of Nature (IUCN), and several European countries are increasing their commitment and efforts to ensure the persistence of this charismatic

species (Otero et al. 2017; Cau et al. 2013). In Chapter 2, we explored the short and long-term consequences of fishing based on data in two affected populations on the coast of Marseille (France). Our results reveal dramatic reductions of coral biomass and minimal recovery after 7yrs. We then predicted that full recovery would take several decades. Interestingly, we unraveled an important recovery mechanism for this species: when fishers remove coral branches but leave the colony basis attached to the substratum, most colonies can survive and regrow new branches through asexual reproduction. This recovery mechanism had not been quantified before and may explain the paradoxal persistence of this extremely slow-growing species that has been intensive overexploited since ancient times. These results highlight that not only minimum colony-size or harvesting quotes are important to enhance the sustainability of red coral fisheries, but also regulating fishing methods can aid enhancing population recovery. We must also note that,

although in our studied populations open to fishing we observed relatively low impact practices, highly destructive fishing gears are still used in Mediterranean waters (Cattaneo-Vietti et al. 2017), and historical records and local ecological knowledge clearly indicate that several red coral populations have gone locally extinct due to overfishing or remain functionally impaired (Garrabou et al. 2017b, Linares et al. 2017).

Based on the findings presented here and evidences from several scientists across the Mediterranean (Garrabou et al. 2017b), minimum requirements for improving the sustainability of red coral fisheries may include: (1) banning on all red coral fishing for a long recovery period of at least two decades until populations can recover its structural role and detailed management plans can be developed; (2) gathering enough information on the distribution, genetic pool and size-distribution of potentially target populations to evaluate vulnerability levels and assess feasible quotes; (3)

a long-term rotatory harvesting scheme coupled to restrictions on minimum sizes and the obligation of leaving all colony bases attached to the substratum. These would be minimum regulations needed to ensure a more sustainable fishery. However, considering that red coral products are a non-locally marketed and only provide few jobs, while causing disproportional damage to benthic ecosystems and long-lasting ecological effects, we advocate for a fully protecting of the red coral as a cultural and ecologically important species, abandoning any type of extractive practice, and develop a re-insertion plan for artisanal red coral fishers into education tasks or alternative community services. In this sense, a recent new legislation approved by the Government of Catalonia in 2018, which will totally ban red coral extractions for a period of ten years, provides a glimpse of hope for this species and the overall conservation of coralligenous habitats. Besides the case of the Mediterranean red coral, our findings can inform fishery management on other precious coral

species that share similar life-history traits such as the Japanese red coral *Paracorallium japonicum* or pink coral *Corallium elatius*, in which the sustainability of current practices is also at concern (Bruckner 2013, 2016).

Red coral poaching is still widespread across European and African Mediterranean coasts. Thus, if not properly enforced, new and more restrictive fishery regulations can be *dead papers* and remain ineffective at meeting conservation goals. In an increasingly globalized world, growing Asian demands for goods are putting at risk several vulnerable species, including precious corals (Bruckner et al. 2016). Banning external trade for endangered species such as the red coral will therefore be an essential step to prevent illegal fishing.

The Mediterranean Sea is a socially complex multi-state ecosystem that currently faces great politically challenges, including large income inequality among different regions and high immigration pressure of refugees escaping from wars and

hanger. Reversing historical depletion of coastal habitats and buffering novel global threats will be especially challenging in this context. The European Union is embracing several conservation initiatives that aim to protect and restore vulnerable species and habitats in EU coastal waters. However, the success of any large-scale conservation plan will require cooperation between non-EU Mediterranean states and those along Nord-African coasts as well as global agreements to regulate trade of endangered species.

Marine Protected Areas

An alternative conservation tool to recover fish stocks and enhance the conservation of benthic habitats is the implementation of spatially-explicit fishery closures or Marine Protected Areas (MPAs). Previous studies demonstrated that MPAs can recover size-structure of red coral populations after several decades of proper enforcement (Linares et al. 2010, 2012, Bavestrello et al. 2015). Whether MPAs also shape their resilience in the context of climate

change is an important further question yet to be addressed. In Chapter 3, we used Integral Projection Models (IPMs) to explore the demographic consequences of MPAs and climate change. Contrary to priority expectations, our results showed that there was no high extinction risk for fished red coral populations, nor significant differences in long-term growth between fished and protected populations. These observed patterns could be explained by similar survival rates of fished colonies due to low impact fishing practices in the studied populations (see Chapter 2 for a detailed analysis). However, colony shrinkage rates were strongly shaped by local protection: much higher in unprotected than protected populations. This drives a pattern of similar population size dynamics regardless of protection status but strongly divergent dynamics in structural complexity, with unprotected populations remaining highly simplified and dominated by small colonies. This allows for a mechanistic understanding of the

broad-scale patterns that have been reported for Mediterranean waters, which showed unprotected populations to be heavily depleted in terms structural complexity but found no consistent patterns in population densities (Garrabou et al. 2017a, 2017b) and highlight the role of MPAs at enhancing the functionality of benthic habitats. More broadly, these results call for a reconsideration of traditional measures of extinction-risk that have been largely based on population sizes or densities. For clonal marine habitat forming-species, we need to move towards a more holistic characterization of conservation status for benthic populations and communities that includes trends in population sizes but also take into account their structural complexity and functional role.

Climate Change

Ongoing climate change is a novel threat to the conservation of most habitat-forming species inhabiting shallow waters. In the NW Mediterranean, red coral populations have been impacted by several heat

waves during past decades, and there is compiling evidence from field studies and aquaria experiments that warm temperatures can cause their partial and whole-colony mortality (Garrabou et al. 2001, 2009, Torrents et al. 2008). Here, by integrating observed warming-driven mortality rates into demographic models, we show the detrimental population-level consequences of climate change in the long-term. While *C. rubrum* has persisted after historical overexploitation due to a high resistance, several shallow populations may be now at risk due to recurrent warming events that sharply decrease survival rates of coral colonies. Even if declines seem subtle in the short-term, and may go unnoticed for managers, our demographic projections reveal consistent declines and potential local extinctions in the long-term with low uncertainty (Hughes et al. 2013).

To predict potential consequences of future increases of thermal stress in coral populations, we used

demographic and spatial models. On one hand, in Chapter 3, we tested whether MPAs enhance resilience of coral population to current and future warming scenarios. Because our demographic data series did not include the interaction between fishing and warming (we did not have unprotected sites open to fishing also affected by warming), we used IPMs to simulate their stochastic dynamics into different levels of local impacts (fishing) and warming-driven mortality frequencies. Our simulations reveal that MPAs may provide a slight buffering effect under low to moderate warming scenarios, but their effect become much weaker and almost negligible at higher frequencies of warming. These results agree with recent large-scale surveys in tropical coral reefs, which demonstrated that local protection had no influence on coral resistance to high intensity heat waves (Hughes et al. 2017).

On the other hand, in Chapter 5, we used spatial models to explore geographic and depth-related

patterns of expected impacts of climate change for 2100 under two emissions scenarios (RCP4.5 and RCP8.5). Our models predict broad-scale decreases in the probability of occurrence for *C. rubrum* as well as for another long-lived gorgonian, *Paramuricea clavata*. This modeling exercise revealed a crucial role of depth in shaping the extent and severity of thermal impacts across the Mediterranean Sea. Shallow populations (up to 25m depth) may experience extensive thermal stress across the Mediterranean Sea, suggesting the warming is likely to drive a regional-scale deepening of gorgonian populations. At deeper sites, however, we expect heterogeneous spatial patterns with some regions highly affected up to 45 and 55m depth.

Earth's oceans are inherently three-dimensional. Many environmental and biotic processes vary widely across depths. However, most current predictive modeling approaches and management decisions and policies do not incorporate the 3D patterns of

expected impacts into consideration. Our results provide a characterization of potential impacts of climate change across depth layers and call for considering this third dimension when analyzing broad scale conservation area priorities (Levin et al. 2017, Venegas-Li et al. 2018). Although they represent an important step forward in our understanding of the geographic and bathymetric dynamics of future climate, we are aware of some potential limitations of the climatic data, distribution data and the correlational nature of our modeling framework (see Quantitative Approaches section for a detailed discussion).

Future directions

Coping with damage to marine ecosystems from multiple interacting stressors will require major efforts from managers, policymakers, and society. Science should also play a crucial role in this endeavor by providing effective tools and evidence-based frameworks for optimal decision-making in conservation. I strongly believe that new scientific paradigms and large steps forwards can only be accomplished if we broaden our perspectives on how natural systems work. It is critically to advance our current understanding on how benthic systems will respond to changing environments and, more importantly, when and how can we use local management to enhance ecosystem resilience to global change.

To address this important question, we need to work towards to a more inclusive and interdisciplinary science, enlarge collaborative networks, and bridge traditionally isolated disciplines. For instance,

since marine reserves manage people and the uses they make of the ocean, we cannot keep isolating marine ecosystems from human dynamics when projecting trends and responses to multiple drivers. Thus, one of my future goals is to include the social dimensions into ecological models to provide more realistic projections of social-ecological change.

My personal experience tells me that combining fieldwork and quantitative tools is a powerful approach to explore patterns and unravel mechanisms driving the dynamics of ecosystems and I would like this to keep playing a major role in my future career. However, exploring alternative perspectives using theoretical models could guide novel research lines and feed parallel empirical studies. Theoretical ecology forces an in-deep evaluation of how natural systems work, allowing to test a wider range of hypotheses and evaluate the sensitivity of the overall system to different components. I believe a hybrid approach using

fieldwork and theoretical models could provide new insights into expected responses of vulnerable ecosystems to global change. Also, by comparing theoretical predictions against empirical observations, we may be able to identify crucial knowledge gaps, misperceptions, and challenge current paradigms.

Finally, I would like to explicitly bridge molecular and population ecology to assess the relevance of potential adaptive responses of corals to climate change. Recent studies suggest a potential critical role of adaptive responses such as evolutionary genetic adaptation and acclimatization in the future persistence of coral reefs. Most of these studies, however, have looked at molecular responses at the individual level. I would like to use novel statistical approaches and complex ecosystem models as a platform to scale-up from the observed adaptive responses of individuals to explore the broad implications of adaptation and acclimatization for the resilience of

coral populations under present and future climatic scenarios.

General conclusions

Chapter I

There are strong positive relationships between depth and longevity across multiple sessile marine taxa.

Extreme longevity leads to stable population dynamics.

The red coral *Corallium rubrum* is an extreme long-lived species with low reproduction success, stable population dynamics, and high vulnerability to increased mortality sources.

Chapter II

Red coral fishing causes large decreases in biomass and strong size-class distribution shifts towards populations dominated by small colonies.

Recovery after fishing is a very slow process that can take decades and relies on regrowth of new branches of colonies that survive.

High survival and regrowth capacity explain the persistence of this species after overexploitation during millennia.

Chapter III

Climate change has dramatic population consequences for the red coral in the long-term.

MPAs are an effective local conservation tool to enhance the structural complexity of red coral populations.

However, MPAs may not be enough to ensure their persistence under future scenarios of increased thermal stress.

Chapter IV

Life-history strategies affect the outcomes and timescales of active restoration actions.

Transplantation of slow-growing species will tend to require lower initial effort due to higher survival after transplanting, but the period required to fully recover habitat complexity will tend to be far longer.

Chapter V

Temperature and primary productivity shape the upper bathymetric range of *Corallium rubrum* and *Paramuricea clavata*.

At the end of the present century, climate change will strongly impact gorgonian populations across the Mediterranean Sea.

The severity of warming impacts will be strongly mediated by depth and global greenhouse gas emissions.

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Annexes

Supplementary Materials Chapter I

Figure S1. Long-term population trends at nine red coral populations in the NW Mediterranean.

Figure S2. *Corallium rubrum* mean and maximum longevity estimates depending on matrix dimensions.

Figure S3. *Corallium rubrum* normalized age-dependent vital rates and size-dependent elasticity patterns.

Table S1. Red coral populations in the NW Mediterranean Sea studied in this work.

Table S2. Correlation coefficients between maximum depth and maximum lifespan in marine sessile species.

Table S3. Best fit models for different longevity estimates based on matrix models of octocorals, hexacorals and sponges.

Table S4. Summary statistics of best supported multiple linear regression model between maximum depth occurrence and maximum lifespan marine sessile species

Table S5. Summary statistics of best-supported multiple linear models for the effect of maximum depth occurrence on maximum lifespan of marine sessile species, including the potential effects of different ageing methods (annual ring counts, radiocarbon dating, demographic simulations and growth curves) and taxonomic levels (Phylum, Class, Order). Model support was based on Akaike Information Criteria corrected for small sample sizes (AICc).

Table S6. Summary statistics of quantile linear regression models between maximum depth occurrence and maximum lifespan across all marine sessile species (n = 223).

Table S7. Multiple linear regression models used to assess the effects of maximum longevity on temporal variance in population size in terrestrial plant species (n=25) and marine sessile invertebrates (n=9).

Table S8. Mean fecundity values of marine sessile species ($n = 28$). Fecundity values correspond to the mean annual recruitment rate per adult and were computed based on the F values of each size-class of the population matrix (first row), weighed by the proportion of each class at the Stable Stage Distribution. We only included species in which the first stage corresponded to recruits.

Table S9. Source studies of longevity estimates and maximum occurrence depth for marine sessile species included in the comparative analysis.

Table S10. Source studies of long-term demographic variability estimates included in the comparative analysis.

Supplementary Methods (I): Model Construction

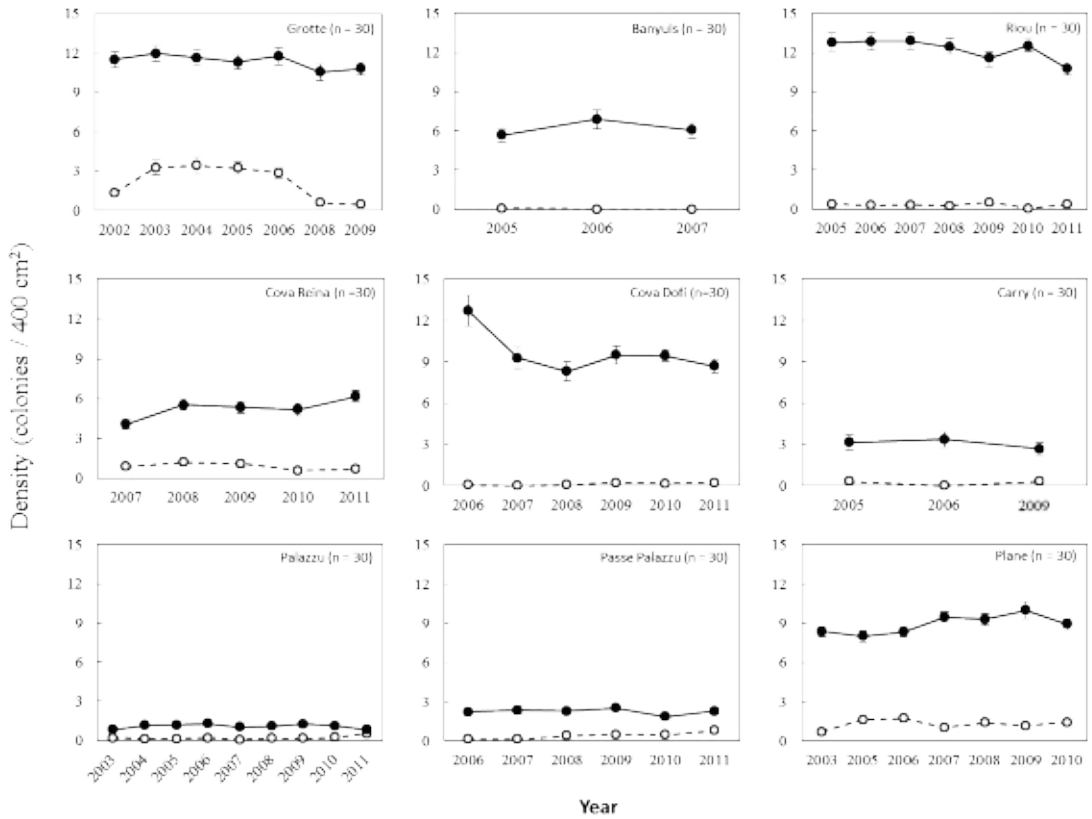


Figure S1. Long-term population trends at nine red coral populations in the NW Mediterranean. Black dots indicate adult and white dots recruit densities.

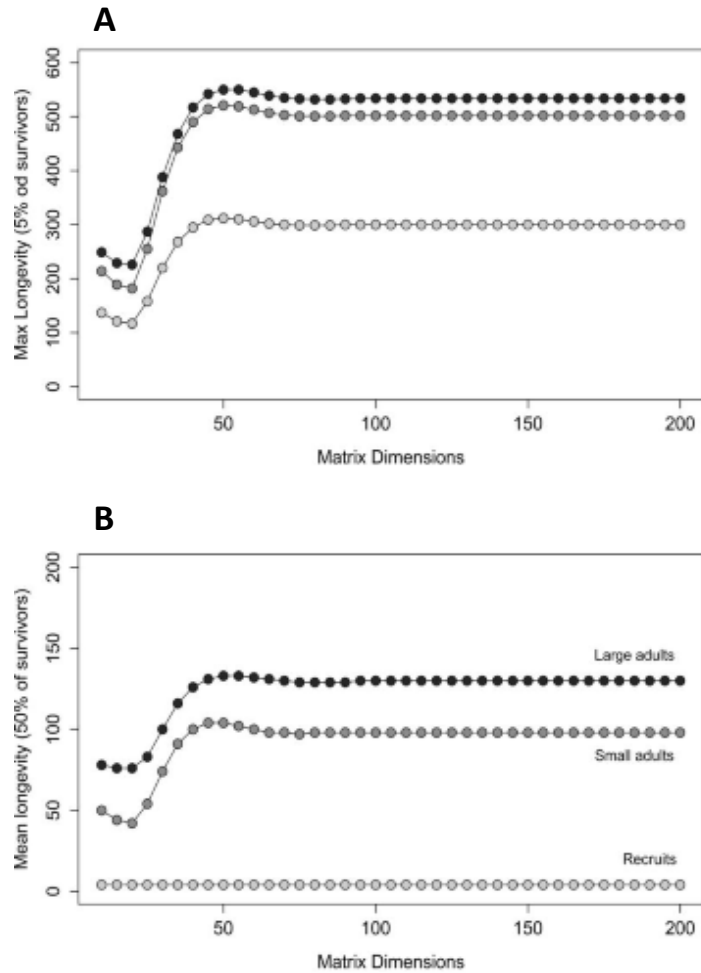
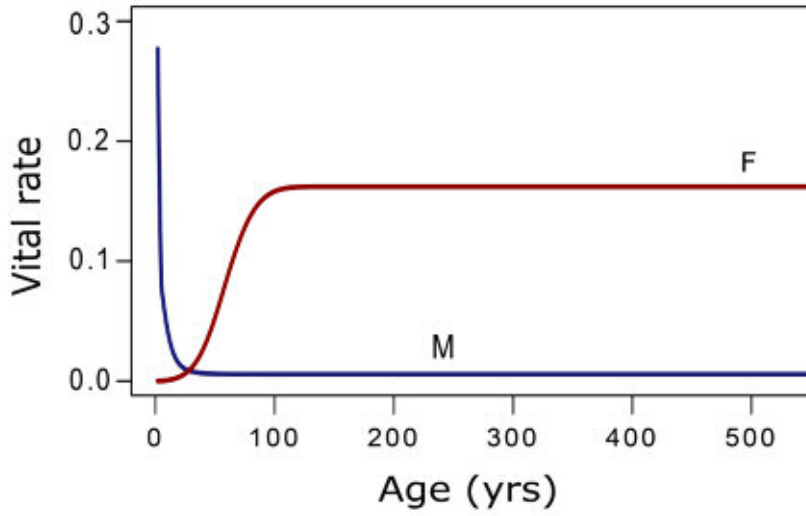


Figure S2. *Corallium rubrum* mean and maximum longevity estimates depending on matrix dimensions. A. Shows maximum potential lifespan, the number of years until 5% of the initial population remains alive. B. Shows the mean lifespan, the number of years until the 50% of the initial population remains alive. Simulations were performed based on 100 individuals starting as recruits (light grey), adults of the smallest size class (grey), and adults of the largest size class (black).



S3. *Corallium rubrum* Normalized age-dependent vital rates: mortality (M) and fecundity (F).

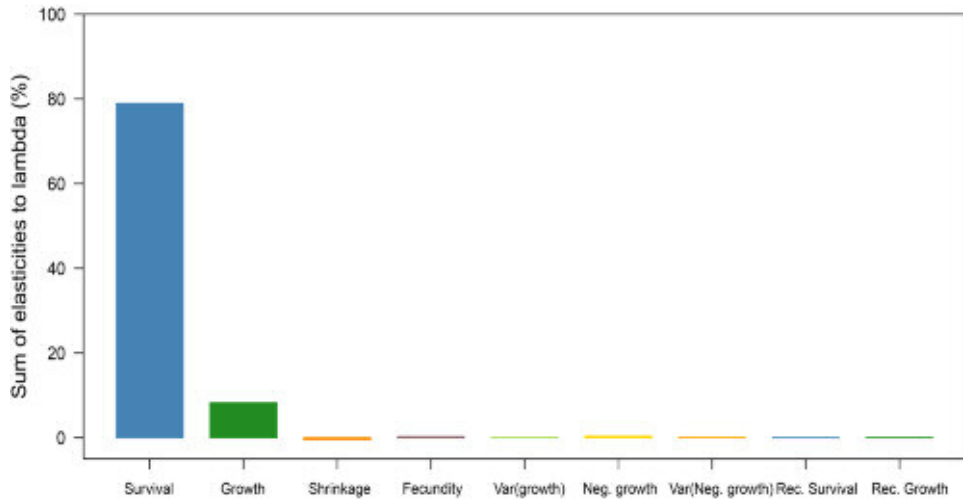


Figure S4. Patterns of sum of elasticity values to the asymptotic population growth rates of different vital rates in the red coral *Corallium rubrum*. From left to right: survival of adult colonies, mean normal growth of adult colonies, probability of extreme shrinkage, fecundity, variance in normal growth of adult colonies, mean negative growth, variance in mean negative growth, recruits survival, and recruits growth.

Table S1. Red coral (*Corallium rubrum*) populations in the NW Mediterranean Sea used in this study.

Name	Longitude	Latitude	Depth (m)	Area	Protection Level	N	Years Surveyed
Dofi	3° 13' 44" E	42° 2' 29" N	17 - 22	Catalan Sea	Partial Protection	148	2006 to 2011
Reina	3° 13' 34" E	42° 2' 49" N	17 - 24	Catalan Sea	Partial Protection	140	2007 to 2011
Banyuls	3°7'50" E	42°28' 60" N	23 - 25	Catalan Sea	Integral Protection	108	2005 to 2007
Maire	5°20'14" E	43°12'32" N	15 - 18	Marseille	Unprotected	140	2002 to 2009
Riou	5° 14' 3" E	43° 6' 13" N	17 - 24	Marseille	Unprotected	204	2005 to 2009
Plane	5° 14' 3" E	43° 11' 54" N	17 - 24	Marseille	Unprotected	154	2003 to 2010
Passe	8° 33' 27" E	42°21' 35" N	24 - 27	Corsica	Integral Protection	145	2006 to 2011
Palazzu	8° 33' 27" E	42° 21' 35" N	19 - 22	Corsica	Integral Protection	137	2003 to 2011

Table S2. Correlation coefficients between maximum depth and maximum lifespan in marine sessile species. The category “All Marine Sessile” includes data on species in five most-represented taxa plus those from other taxa, including bryozoan, hydrozoan, polychaeta, cirripeds and one tunicate species. Statistics shown are: n = number of species included in the analysis, rho = Spearman’s correlation coefficient; P = Pearson’s correlation coefficient.

Spearman’s Rank correlation

Taxa	Maximum Lifespan vs. Maximum Depth Occurrence			Maximum Lifespan vs. Depth at the Study Site		
	n	rho	P value	n	rho	P value
Octocorals	35	0.673	P < 0.001	34	0.546	P = 0.008
Hexacorals	40	0.592	P < 0.001	34	0.704	P < 0.001
Sponges	17	0.647	P < 0.001	16	0.729	P = 0.001
Bivalves	93	0.358	P < 0.001	34	0.700	P < 0.001
Macroalgae	14	0.600	P = 0.023	9	0.597	P = 0.089
All Marine Sessile	223	0.503	P < 0.001	138	0.695	P < 0.001

Pearson’s R correlation

Taxa	Maximum Lifespan vs. Maximum Depth Occurrence			Maximum Lifespan vs. Depth at the Study Site		
	n	R	P value	n	R	P value
Octocorals	35	0.714	P < 0.001	34	0.594	P = 0.0002
Hexacorals	40	0.650	P < 0.001	34	0.790	P < 0.001
Sponges	17	0.739	P < 0.001	16	0.727	P = 0.0014
Bivalves	93	0.456	P < 0.001	34	0.679	P < 0.001
Macroalgae	14	0.620	P = 0.018	9	0.674	P = 0.046
All Marine Sessile	223	0.547	P < 0.001	138	0.674	P < 0.001

Table S3. Summary of the four best supported models predicting longevity estimates based on matrix models of octocorals, hexacorals and sponges.

Model	Predictors / Levels	Estimate	Std. Error	P value	AICc	delta AIC
Maximum Longevity Adults ~ Depth + Dimensions	Intercept	0.849207	0.278340	0.0072	67.219	0
	Maximum Depth	0.020635	0.008199	0.0222		
	Dimensions	-0.157308	0.082079	0.0723		
	Overall Model: $R^2 = 0.4298$; $F = 8.161$; $df = 17$; $P = 0.0033$					
Maximum Longevity Adults ~ Depth	Intercept	0.956596	0.292187	0.0042	67.965	0.746
	Maximum Depth	0.005170	0.001558	0.0038		
	Overall Model: $R^2 = 0.3796$; $F = 11.01$; $df = 18$; $P = 0.0038$					
Mean Longevity Recruits ~ Depth + Dimensions + Depth*Dimensions	Intercept	2.0324517	1.0345084	0.067	68.833	1.613
	Maximum Depth	0.0143110	0.0088083	0.124		
	Dimensions	-0.6189497	0.2300075	0.016		
	Maximum Depth*Dimensions	0.0006347	0.0002998	2.117		
	Overall Model: $R^2 = 0.5463$; $F = 8.625$; $df = 18$; $P = 0.001232$					
Mean Longevity Adults ~ Depth + Dimensions + Depth*Dimensions	Intercept	1.9999077	1.0460125	0.074	69.275	2.056
	Maximum Depth	0.0164804	0.0089062	0.083		
	Dimensions	-0.4057816	0.2325653	0.100		
	Maximum Depth*Dimensions	0.0003458	0.0003032	0.271		
	Overall Model: $R^2 = 0.4397$; $F = 5.97$; $df = 16$; $P = 0.00624$					

Table S4. Summary statistics of best supported multiple linear regression model between maximum depth occurrence and maximum lifespan for all marine sessile species (n = 223). Model support was based on Akaike Information Criteria corrected for small sample sizes (AICc). The second-best model had deltaAICc of 3.568. Data were log-transformed.

Best Fit Model	Predictors / Levels	Estimate	Std. Error	t value	P value
Longevity ~ Depth + Taxa + Method + Depth*Taxa	Intercept	0.67424	0.14385	4.687	< 0.001
	Maximum Depth	0.29642	0.07414	3.998	< 0.001
	Taxa (Hexacoral)	-0.01922	0.34657	-0.055	0.9558
	Taxa (Macroalgae)	-0.43510	0.33371	-1.304	0.1939
	Taxa (Octocoral)	0.62504	0.36226	1.725	0.0861
	Taxa (Sponge)	-0.71140	0.56550	-1.258	0.2100
	Method (Matrix/Survival)	-0.17426	0.09751	-1.787	0.0756
	Method (Oxygen)	0.33719	0.09751	0.883	0.3785
	Method (Radiometric Age)	0.39427	0.38196	2.492	0.0136
	Maximum Depth * Taxa (Hexacoral)	0.21700	0.15820	1.223	0.2228
	Maximum Depth * Taxa (Macroalgae)	0.59011	0.17740	2.176	0.0308
	Maximum Depth * Taxa (Octocoral)	-0.05002	0.27123	-0.294	0.7689
	Maximum Depth * Taxa (Sponge)	0.74739	0.29825	2.506	0.0131
	Overall model statistics: $R^2 = 0.5105$; $F = 18.12$; $df = 185$; $P < 0.0001$				

Table S5. Summary statistics of best supported multiple linear regression models between maximum depth occurrence and maximum lifespan in marine sessile species, including the potential effects of different ageing methods (annual ring counts, radiocarbon dating, demographic simulations and growth curves) and taxonomic levels (Phylum, Class, Order). Model support was based on Akaike Information Criteria corrected for small sample sizes (AICc).

Formula	Intercept	Depth slope (P value)	df	AICc	Delta AICc
Longevity ~ Depth + Class + Method	1.0129	0.3633 (>0.0001)	15	315.0721	0
Longevity ~ Depth + Class + Method + Depth*Class	0.8859	0.4299 (0.00101)	21	316.2796	1.2075
Longevity ~ Depth + Phylum + Depth*Phylum	-2.0621	1.4262 (0.2853)	15	316.8752	1.803146
Longevity ~ Depth + Method + Phylum	-2.8295	1.8155 (0.175)	12	319.1589	4.086796
Longevity ~ Depth + Class + Depth*Class	0.4153	0.6715 (>0.0001)	18	322.0485	4.0868

Table S6. Summary statistics of quantile linear regression models between maximum depth occurrence and maximum lifespan across all marine sessile species (n = 223)

Quantile	Intercept (IC 95%)	Slope (IC 95%)
10%	-0.0682 (-0.516 – 0.083)	0.441 (0.361 – 0.677)
20%	0.0691 (-0.174, 0.173)	0.486 (0.427 – 0.593)
80%	0.811 (0.547 – 0.972)	0.628 (0.570 – 0.791)
90%	0.9031 (0.691 – 1.071)	0.715 (0.550 – 0.918)

Table S7. Multiple linear regression models used to assess the effects of maximum longevity on temporal variance in population growth (realized log-lambda values) in terrestrial plant species (n=25) and marine sessile invertebrates (n=9).

Model	R²	AICc	Delta	Weight
log(Variance) ~ log(Longevity) + Habitat	0.543	125.4652	0	0.6222
log(Variance) ~ log(Longevity) + Habitat + log(Longevity)*Habitat	0.528	128.1723	2.707	0.1607
log(Variance) ~ log(Longevity)	0.466	130.2689	4.804	0.0671
log(Variance) ~ Habitat	0.224	142.9632	17.498	0.0001

Table S8. Mean fecundity values of marine sessile species (n = 28). Fecundity values correspond to the mean annual recruitment rate per adult and were computed based on the F values of each size-class of the population matrix (first row), weighed by the proportion of each class at the Stable Stage Distribution. We only included species in which demographic data in the form of standar matrix population models, including fecundity were available. To allow standard comparisons, we did not include matrices in which the first stage corresponded to larvae or eggs.

Species Name	Taxa	Fecundity	Source
<i>A. islandica</i>	Mollusca	0.001	Ripley & Caswell 2008
<i>P. abrupta</i>	Mollusca	0.004	Ripley & Caswell 2008
<i>X. muta</i>	Mollusca	0.006	McMurray <i>et al.</i> 2010
<i>T. gigas</i>	Mollusca	0.013	Ripley & Caswell 2008
<i>A. colbecki</i>	Mollusca	0.040	Ripley & Caswell 2008
<i>A. compressa</i>	Mollusca	0.071	Mercado-Molina 2011
<i>C. rubrum</i>	Cnidaria	0.077	Present study
<i>A. palmata</i>	Cnidaria	0.077	Vardi <i>et al.</i> 2012
<i>P. astreoides</i>	Cnidaria	0.083	Edmunds 2010
<i>L. digitata</i>	Macroalgae	0.126	Chapman 1990
<i>S. graminea</i>	Porifera	0.135	Cropper 1999
<i>G. demissa</i>	Mollusca	0.254	Ripley & Caswell 2008
<i>N. obscurata</i>	Mollusca	0.261	Dudas. <i>et al.</i> 2007
<i>H. iris</i>	Mollusca	0.547	Somerville <i>et al.</i> 2014
<i>H. sorenseni</i>	Mollusca	0.727	Roggers-Bennett 2006
<i>H. rufenses</i>	Mollusca	0.794	Roggers-Bennett 2006
<i>P. caesia</i>	Cnidaria	0.847	Tanner 1997
<i>B. annulata</i>	Cnidaria	0.875	Nelsen 2008
<i>A. nana</i>	Macroalgae	0.966	Pfister 2005
<i>L. notorcadensis</i>	Mollusca	1.176	Ripley & Caswell 2008
<i>L. miliaris</i>	Mollusca	1.639	Ripley & Caswell 2008
<i>H. laevigata</i>	Mollusca	1.654	Fordham <i>et al.</i> 2013
<i>M. mactroides</i>	Mollusca	1.791	Brazeiro & Defeo 1999
<i>M. californianus</i>	Mollusca	2.314	Carson <i>et al.</i> 2011
<i>M. galloprovincialis</i>	Mollusca	2.768	Carson <i>et al.</i> 2011
<i>Alcyonium sp.</i>	Cnidaria	3.257	McFadden 1991
<i>F. distichus</i>	Macroalgae	8.001	Ang & De Wreede 1993
<i>L. rubra</i>	Mollusca	115.314	Ripley & Caswell 2008

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Table S9. Source studies of longevity estimates and maximum occurrence depth for marine sessile species included in the comparative analysis (n = 241) (See completed references list below).

Name	Family	Taxa	Lifespan Method	Source Lifespan	Source Depth
<i>Acropora cervicornis</i>	Acroporidae	Hexacoral	Matrix/Survival	1	37, 44
<i>Acropora hyacinthus</i>	Acroporidae	Hexacoral	Matrix/Survival	2	38, 44
<i>Acropora palmata</i>	Acroporidae	Hexacoral	Matrix/Survival	3	37, 44
<i>Agaricia agaricites</i>	Agariciidae	Hexacoral	Matrix/Survival	4	37, 44
<i>Leptoseria cucullata</i>	Agariciidae	Hexacoral	Matrix/Survival	9	49, 50, 44
<i>Bartholomea annulata</i>	Aiptasiidae	Hexacoral	Matrix/Survival	19	42
<i>Antipathes dendrochristos</i>	Antipathidae	Hexacoral	Radiometric Age	69, 70	69
<i>Antipathes dichotoma</i>	Antipathidae	Hexacoral	Radiometric Age	58	NA
<i>Antipathes griggi</i>	Antipathidae	Hexacoral	Radiometric Age	70	NA
<i>Caryophyllia inornata</i>	Caryophylliidae	Hexacoral	Growth	76, 86	87
<i>Desmophyllum cristagalli</i>	Caryophylliidae	Hexacoral	Radiometric Age	60	NA
<i>Lophelia pertusa</i>	Caryophylliidae	Hexacoral	Radiometric Age	131	130
<i>Solenosmilia variabilis</i>	Caryophylliidae	Hexacoral	Radiometric Age	132	133
<i>Leptosammia pruvoti</i>	Dendrophylliidae	Hexacoral	Growth	84, 76	81
<i>Balanophyllia elegans</i>	Dendrophylliidae	Hexacoral	Matrix/Survival	79	78
<i>Balanophyllia europea</i>	Dendrophylliidae	Hexacoral	Growth	80	81, 82
<i>Enallopsammia rostrata</i>	Dendrophylliidae	Hexacoral	Radiometric Age	134	135, 136
<i>Ctenactis echinata</i>	Fungiidae	Hexacoral	Growth	52	44
<i>Diaseris distorta</i>	Fungiidae	Hexacoral	Growth	52	37, 44
<i>Fungia actiniformis</i>	Fungiidae	Hexacoral	Growth	52	44
<i>Fungia fungites</i>	Fungiidae	Hexacoral	Growth	52	44
<i>Fungia granulosa</i>	Fungiidae	Hexacoral	Growth	52	49, 44
<i>Fungia paumotensis</i>	Fungiidae	Hexacoral	Growth	52	49, 44
<i>Leiopathes glaberime</i>	Leiopathidae	Hexacoral	Radiometric Age	58, 70	70
<i>Leiopathes sp.</i>	Leiopathidae	Hexacoral	Radiometric Age	57	133
<i>Goniastrea aspera</i>	Merulinidae	Hexacoral	Matrix/Survival	7	44
<i>Goniastrea favulus</i>	Merulinidae	Hexacoral	Matrix/Survival	7	258
<i>Orbicella annularis</i>	Merulinidae	Hexacoral	Matrix/Survival	9	37, 44
<i>Platygyra sinensis</i>	Merulinidae	Hexacoral	Matrix/Survival	7	237
<i>Diploria strigosa</i>	Mussidae	Hexacoral	Matrix/Survival	6	45
<i>Antipathella fiordensis</i>	Myriopathidae	Hexacoral	Growth	71, 70	41
<i>Plumapathes pennacea</i>	Myriopathidae	Hexacoral	Growth	72, 70	NA

<i>Kulamanamana haumeae</i>	Parazoanthidae	Hexacoral	Radiometric Age	57, 59	59
<i>Parazoanthus axinellae</i>	Parazoanthidae	Hexacoral	Matrix/Survival	75	74
<i>Pocillopora damicornis</i>	Pocilloporidae	Hexacoral	Matrix/Survival	2	48, 44
<i>Alveopora japonica</i>	Poritidae	Hexacoral	Growth	181	181
<i>Porites astreoides</i>	Poritidae	Hexacoral	Matrix/Survival	6	49, 50, 44
<i>Fannyella rossii</i>	Primnoidae	Hexacoral	Radiometric Age	64	NA
<i>Stauropathes artica</i>	Schizopathidae	Hexacoral	Radiometric Age	62	NA
<i>Palythoa caesia</i>	Sphenopiidae	Hexacoral	Matrix/Survival	11	41
<i>Alcyonium acaule</i>	Alcyoniidae	Octocoral	Matrix/Survival	75, 76	77
<i>Anthomastus Ritteri</i>	Alcyoniidae	Octocoral	Growth	179	179
<i>Gorgonia ventalina</i>	Anthothelidae	Octocoral	Matrix/Survival	88	91
<i>Chrysogorgia sp.</i>	Chrysogorgiidae	Octocoral	Radiometric Age	177	177
<i>Corallium niobe</i>	Corallidae	Octocoral	Radiometric Age	67	NA
<i>Corallium secundum</i>	Corallidae	Octocoral	Radiometric Age	58	NA
<i>Corallium rubrum</i>	Coralliidae	Octocoral	Matrix/Survival	Present Study	55
<i>Eunicella singularis</i>	Gorgoniidae	Octocoral	Growth	66	73
<i>Leptogorgia sarmentosa</i>	Gorgoniidae	Octocoral	Growth	85	123
<i>Leptogorgia virgulata</i>	Gorgoniidae	Octocoral	Matrix/Survival	8	259
<i>Pseudopterogorgia</i>	Gorgoniidae	Octocoral	Matrix/Survival	14	NA
<i>Pseudopterogorgia acerosa</i>	Gorgoniidae	Octocoral	Matrix/Survival	88	41
<i>Pseudopterogorgia americana</i>	Gorgoniidae	Octocoral	Matrix/Survival	88	91
<i>Halipteris willemoesi</i>	Halipteridae	Octocoral	Radiometric Age	63	NA
<i>Halipteris willemoesi</i>	Halipteridae	Octocoral	Growth	176	176
<i>Isidella tentaculum</i>	Isididae	Octocoral	Radiometric Age	129	NA
<i>Keratosia ornata</i>	Isididae	Octocoral	Radiometric Age	62	NA
<i>Keratosia sp.</i>	Isididae	Octocoral	Radiometric Age	128	NA
<i>Acanella arbuscula</i>	Isididae	Octocoral	Radiometric Age	62	NA
<i>Eunicea laxispica</i>	Plexauridae	Octocoral	Matrix/Survival	88	NA
<i>Eunicea succinea</i>	Plexauridae	Octocoral	Matrix/Survival	88	92
<i>Eunicea tourneforti</i>	Plexauridae	Octocoral	Matrix/Survival	88	NA
<i>Muricea californica</i>	Plexauridae	Octocoral	Growth	83	NA
<i>Muricea fruticosa</i>	Plexauridae	Octocoral	Growth	83	NA
<i>Muriceopsis flavida</i>	Plexauridae	Octocoral	Matrix/Survival	88	92
<i>Paramuricea biscaya</i>	Plexauridae	Octocoral	Growth	177	177
<i>Paramuricea clavata</i>	Plexauridae	Octocoral	Matrix/Survival	12	56
<i>Paramuricea spp.</i>	Plexauridae	Octocoral	Radiometric Age	62	NA

<i>Plexaura A</i>	Plexauridae	Octocoral	Matrix/Survival	13	NA
<i>Plexaura flexuosa</i>	Plexauridae	Octocoral	Matrix/Survival	88	91
<i>Plexaurella dichotoma</i>	Plexauridae	Octocoral	Matrix/Survival	88	91
<i>Pseudoplexaura porosa</i>	Plexauridae	Octocoral	Matrix/Survival	88	91
<i>Pseudoplexaura wagnaari</i>	Plexauridae	Octocoral	Matrix/Survival	88	91
<i>Primnoa resedaeformis</i>	Primnoidae	Octocoral	Radiometric Age	60, 65	62
<i>Thouarella variabilis</i>	Primnoidae	Octocoral	Radiometric Age	64	NA
<i>Fannyella abies</i>	Primnoidae	Octocoral	Radiometric Age	64	NA
<i>Aplysina cavernicola</i>	Aplysinidae	Sponge	Growth	76	104
<i>Amphimedon compressa</i>	Chalinidae	Sponge	Matrix/Survival	5	102
<i>Haliclona fulva</i>	Chalinidae	Sponge	Growth	76	103
<i>Chondrosia reniformis</i>	Chondrosiidae	Sponge	Growth	76	99
<i>Crambe crambe</i>	Crambeidae	Sponge	Matrix/Survival	76	101
<i>Desmapsamma anchorata</i>	Desmacididae	Sponge	Matrix/Survival	105	106
<i>Monorhaphis chuni</i>	Monorhaphididae	Sponge	Radiometric Age	94	NA
<i>Mycale fistulifera</i>	Mycalidae	Sponge	Matrix/Survival	170	NA
<i>Petrosia ficiformis</i>	Petrosiidae	Sponge	Growth	76	97, 98, 100
<i>Xestospongia muta</i>	Petrosiidae	Sponge	Growth	89	124
<i>Rhabdocalyptus dawsoni</i>	Rossellidae	Sponge	Growth	96	NA
<i>Rosella spp.</i>	Rossellidae	Sponge	Oxygen_Size	93	NA
<i>Rosella racovitzae</i>	Rossellidae	Sponge	Radiometric Age	95	NA
<i>Spirastrella cunctatrix</i>	Spirastrellidae	Sponge	Growth	76	100
<i>Spongia graminea</i>	Spongiidae	Sponge	Matrix/Survival	32	261
<i>Stylocordyla borealis</i>	Stylocordylidae	Sponge	Oxygen_Size	93	NA
<i>Cinachyra antarctica</i>	Tetillidae	Sponge	Oxygen_Size	93	NA
<i>Scalarispongia scalaris</i>	Thorectidae	Sponge	Growth	76	100
<i>Glossus humanus</i>	Glossidae	Bivalve	Growth	212	213
<i>Panopea generosa</i>	Hiatellidae	Bivalve	Growth	216	190, 217
<i>Perna perna</i>	Mytilidae	Bivalve	Growth	236	2237
<i>Pedum spondyloideum</i>	Pectinidae	Bivalve	Growth	180	248
<i>Anadara tuberculosa</i>	Anadarinae	Bivalve	Growth	207	211
<i>Arca noae</i>	Arcidae	Bivalve	Growth	208	195, 209
<i>Arctica islandica</i>	Arcticidae	Bivalve	Growth	127	121
<i>Astarte arctica</i>	Astartidae	Bivalve	Growth	240	41
<i>Astarte borealis</i>	Astartidae	Bivalve	Growth	240	215
<i>Astarte elliptica</i>	Astartidae	Bivalve	Growth	240	241

<i>Astarte montagui</i>	Astartidae	Bivalve	Growth	240	215
<i>Astarte sulcata</i>	Astartidae	Bivalve	Growth	234	234
<i>Clinocardium californiense</i>	Cardiidae	Bivalve	Growth	240	241
<i>Clinocardium ciliatum</i>	Cardiidae	Bivalve	Growth	240	241
<i>Serripes gloenlandicus</i>	Cardiidae	Bivalve	Growth	240	215
<i>Tridacna gigas</i>	Cardiidae	Bivalve	Matrix/Survival	16	205
<i>Cardita affinis</i>	Carditidae	Bivalve	Growth	226	227
<i>Cyclocardia ventricosa</i>	Carditidae	Bivalve	Growth	240	41
<i>Donax denticulatus</i>	Donacidae	Bivalve	Growth	173	215
<i>Donax denticulatus</i>	Donacidae	Bivalve	Growth	233	215
<i>Donax trunculus</i>	Donacidae	Bivalve	Growth	235	225
<i>Mytilopsis leucophaeata</i>	Dreissenidae	Bivalve	Growth	228	215
<i>Glycymeris bimaculata</i>	Glycymerididae	Bivalve	Growth	210	195
<i>Glycymeris pilosa</i>	Glycymerididae	Bivalve	Growth	194	195
<i>Neopycnodonte zibrowii</i>	Gryphaeidae	Bivalve	Radiometric Age	246	246
<i>Haliotis iris</i>	Haliotidae	Bivalve	Matrix/Survival	24	157
<i>Haliotis laevigata</i>	Haliotidae	Bivalve	Matrix/Survival	33, 34	158
<i>Haliotis rufescens</i>	Haliotidae	Bivalve	Matrix/Survival	25	159
<i>Haliotis sorenseni</i>	Haliotidae	Bivalve	Matrix/Survival	171	160
<i>Cyrtodaria kurriana</i>	Hiatellidae	Bivalve	Growth	240	272, 41
<i>Hiatella arctica</i>	Hiatellidae	Bivalve	Growth	240	273
<i>Panomya ampla</i>	Hiatellidae	Bivalve	Growth	240	41
<i>Panopea abrupta</i>	Hiatellidae	Bivalve	Growth	16, 151	152, 168
<i>Hiatella arctica</i>	Hiatellidae	Bivalve	Growth	119	167
<i>Lasaea rubra</i>	Lasaeidae	Bivalve	Matrix/Survival	16	278
<i>Mysella kurilensis</i>	Lasaeidae	Bivalve	Growth	240	240
<i>Laternula elliptica</i>	Laternulidae	Bivalve	Growth	148	150
<i>Keletistes rhizoecus</i>	Lucinidae	Bivalve	Growth	245	245
<i>Lyonsia arenosa</i>	Lyonsiidae	Bivalve	Growth	240	274
<i>Lyonsia vniroi</i>	Lyonsiidae	Bivalve	Growth	240	NA
<i>Mactromeris polynyma</i>	Mactridae	Bivalve	Growth	240	243
<i>Spisula sachalinensis</i>	Mactridae	Bivalve	Growth	240	244
<i>Spisula solidissima</i>	Mactridae	Bivalve	Growth	172	NA
<i>Spisula subtruncata</i>	Mactridae	Bivalve	Matrix/Survival	224	225
<i>Mesodesma mactroides</i>	Mesodesmatidae	Bivalve	Growth	175	174
<i>Mya arenaria</i>	Myidae	Bivalve	Growth	148	149, 196
<i>Mya baxteri</i>	Myidae	Bivalve	Growth	240	NA

<i>Mya pseudoarenaria</i>	Myidae	Bivalve	Growth	240	41
<i>Mya truncata</i>	Myidae	Bivalve	Growth	240	215
<i>Mytilus trossulus</i>	Mytilidae	Bivalve	Growth	240	242
<i>Arvella manshurica</i>	Mytilidae	Bivalve	Growth	240	240
<i>Crenella decussata</i>	Mytilidae	Bivalve	Growth	240	215
<i>Crenomytilus grayanus</i>	Mytilidae	Bivalve	Growth	240	192
<i>Geukensia demissa</i>	Mytilidae	Bivalve	Matrix/Survival	16	156
<i>Limnoperna fortunei</i>	Mytilidae	Bivalve	Growth	218	219
<i>Lithophaga lithophaga</i>	Mytilidae	Bivalve	Growth	206	195
<i>Lithophaga patagonica</i>	Mytilidae	Bivalve	Growth	214	215
<i>Musculus discors</i>	Mytilidae	Bivalve	Growth	240	215
<i>Musculus niger</i>	Mytilidae	Bivalve	Growth	240	215
<i>Mytilus californianus</i>	Mytilidae	Bivalve	Matrix/Survival	35	161
<i>Mytilus galloprovincialis</i>	Mytilidae	Bivalve	Matrix/Survival	35	162
<i>Perna viridis</i>	Mytilidae	Bivalve	Growth	231	232
<i>Adamussium colbecki</i>	Pectinidae	Bivalve	Matrix/Survival	16	278
<i>Aequipecten opercularis</i>	Pectinidae	Bivalve	Radiometric Age	147	146
<i>Argopecten irradians irradians</i>	Pectinidae	Bivalve	Growth	137	143144
<i>Argopecten purpuratus</i>	Pectinidae	Bivalve	Growth	139140	145
<i>Placopecten megellanicus</i>	Pectinidae	Bivalve	Growth	141, 142	153
<i>Siliqua alta</i>	Pharidae	Bivalve	Growth	240	41
<i>Lissarca miliaris</i>	Philobryidae	Bivalve	Matrix/Survival	16	155
<i>Lissarca notorcadensis</i>	Philobryidae	Bivalve	Matrix/Survival	16	154
<i>Zirfaea pilsbryi</i>	Pholadidae	Bivalve	Growth	240	NA
<i>Pinna nobilis</i>	Pinnidae	Bivalve	Matrix/Survival	122	120
<i>Nuttallia obscurata</i>	Psammobiidae	Bivalve	Matrix/Survival	31, 164	241
<i>Pinctada imbricata</i>	Pteriidae	Bivalve	Growth	229	230
<i>Abra alba</i>	Semelidae	Bivalve	Growth	247	41
<i>Macoma balthica</i>	Tellinidae	Bivalve	Growth	240	241
<i>Macoma calcarea</i>	Tellinidae	Bivalve	Growth	240	241
<i>Macoma lama</i>	Tellinidae	Bivalve	Growth	240	241
<i>Macoma loveni</i>	Tellinidae	Bivalve	Growth	240	215
<i>Macoma middendorffi</i>	Tellinidae	Bivalve	Growth	240	41
<i>Tellina lutea</i>	Tellinidae	Bivalve	Growth	240	41
<i>Tellina tenuis</i>	Tellinidae	Bivalve	Growth	238	239
<i>Thracia septentrionalis</i>	Thraciidae	Bivalve	Growth	240	215
<i>Tindaria callistifomis</i>	Tindariidae	Bivalve	Radiometric Age	138	NA

<i>Tindaria callistiformis</i>	Tindariidae	Bivalve	Radiometric Age	169	NA
<i>Diplodonta aleutica</i>	Ungulinidae	Bivalve	Growth	240	240
<i>Callista brevisiphonata</i>	Veneridae	Bivalve	Growth	197, 200	17, 18
<i>Callista chione</i>	Veneridae	Bivalve	Growth	201	203, 204, 222
<i>Gemma gemma</i>	Veneridae	Bivalve	Matrix/Survival	16	215
<i>Liocyma fluctuosum</i>	Veneridae	Bivalve	Growth	240	272
<i>Mercenaria mercenaria</i>	Veneridae	Bivalve	Growth	101	215
<i>Protothaca euglypta</i>	Veneridae	Bivalve	Growth	221	220, 221
<i>Ruditapes philippinarum</i>	Veneridae	Bivalve	Growth	201, 223	202
<i>Yoldia eightsi</i>	Yoldiidae	Bivalve	Growth	191	193
<i>Yoldia myalis</i>	Yoldiidae	Bivalve	Growth	240	215
<i>Yoldia seminuda</i>	Yoldiidae	Bivalve	Growth	240	241
<i>Alaria nana</i>	Alariaceae	Macroalgae	Matrix/Survival	17	267
<i>Leathesia difformis</i>	Chordariaceae	Macroalgae	Matrix/Survival	263	270
<i>Lithothamnium crassiusculum</i>	Corallinaceae	Macroalgae	Radiometric Age	187	187
<i>Ascophyllum nodosum</i>	Fucaceae	Macroalgae	Matrix/Survival	18	41
<i>Fucus distichus</i>	Fucaceae	Macroalgae	Matrix/Survival	23	268
<i>Pelvetia fastigiata</i>	Fucaceae	Macroalgae	Matrix/Survival	264	116
<i>Iridaea splendens</i>	Gigartinaceae	Macroalgae	Matrix/Survival	26	NA
<i>Gracilaria gracilis</i>	Gracilariaceae	Macroalgae	Matrix/Survival	265	
<i>Halimeda incrassata</i>	Halimedaceae	Macroalgae	Matrix/Survival	182	249
<i>Hormosira banksii</i>	Hormosiraceae	Macroalgae	Matrix/Survival	183	NA
<i>Laminaria digitata</i>	Laminariaceae	Macroalgae	Matrix/Survival	27	NA
<i>Laminaria hyperborea</i>	Laminariaceae	Macroalgae	Growth	117	117
<i>Laminaria hyperborea</i>	Laminariaceae	Macroalgae	Matrix/Survival	184	41
<i>Macrocystis pyrifera</i>	Laminariaceae	Macroalgae	Matrix/Survival	30	266
<i>Petrocelis middendorffii</i>	Phylloporaceae	Macroalgae	Growth	113	NA
<i>Pterocladia capillacea</i>	Pterocladaceae	Macroalgae	Growth	112	269
<i>Petrocellis middendorffi</i>	Rhodymeniophycidae	Macroalgae	Growth	186	NA
<i>Cystoseira zosteroides</i>	Sargassaceae	Macroalgae	NA	114	115
<i>Clathromorphum nereostratum</i>	Hapalidiaceae	Macroalgae	Growth	185	188
<i>Adeonellopsis sp</i>	Adeonidae	Bryozoan	NA	189	NA
<i>Cellepora pumicosa</i>	Celleporidae	Bryozoan	Matrix/Survival	22	262
<i>Mucropetraliella ellerii</i>	Petraliellidae	Bryozoan	Matrix/Survival	10	10
<i>Watersipora subtorquata</i>	Watersiporidae	Bryozoan	Matrix/Survival	20	107
<i>Semibalanus balanoides</i>	Archaeobalanidae	Cirriped (Crustacea)	Matrix/Survival	21	NA

<i>Onithochiton quercinus</i>	Chitonidae	Cirriped (Crustacea)	Growth	271	41
<i>Chthamalus montagui</i>	Chthamalidae	Cirriped (Crustacea)	Matrix/Survival	21	NA
<i>Plaxiphora albida</i>	Mopaliidae	Cirriped (Crustacea)	Growth	271	41
<i>Cantellius sp.</i>	Pyrgomatidae	Cirriped (Crustacea)	NA	180	41
<i>Botrylloides violaceous</i>	Styelidae	Tunicate	Matrix/Survival	20	110
<i>Botryllus schlosseri</i>	Styelidae	Tunicate	Matrix/Survival	20	111
<i>Campanularia everta</i>	Campanulariidae	Hydrozoan	NA	125	NA
<i>Campanularia flexuosa</i>	Campanulariidae	Hydrozoan	NA	126	NA
<i>Corymorpha nutans</i>	Corymorphidae	Hydrozoan	NA	250	250
<i>Corymorpha palma</i>	Corymorphidae	Hydrozoan	NA	250	41
<i>Eutima japonica</i>	Eirenidae	Hydrozoan	NA	253	NA
<i>Helecium petrosum</i>	Haleciidae	Hydrozoan	Growth	108	NA
<i>Helecium pusillum</i>	Haleciidae	Hydrozoan	Growth	108	NA
<i>Nemertesia antennina</i>	Plumulariidae	Hydrozoan	NA	251	41,
<i>Errina novaezelandiae</i>	Stylasteridae	Hydrozoan	Growth	178	178, 41
<i>Tubularia indivisa</i>	Tubulariidae	Hydrozoan	NA	125	NA
<i>Tubularia indivisa</i>	Tubulariidae	Hydrozoan	NA	252	41
<i>Galeolaria hystrix</i>	Serpulidae	Polychaete	NA	255	276
<i>Circeis armoricana</i>	Serpulidae	Polychaete	NA	256	41
<i>Ficopomatus enigmaticus</i>	Serpulidae	Polychaete	NA	254	41
<i>Hydroides elegans</i>	Serpulidae	Polychaete	NA	256	NA
<i>Janua pagenstecheri</i>	Serpulidae	Polychaete	NA	256	277
<i>Neodexiospira brasiliensis</i>	Serpulidae	Polychaete	NA	256	41
<i>Pomatoceros triqueter</i>	Serpulidae	Polychaete	NA	256	257
<i>Pseudochitinopoma occidentalis</i>	Serpulidae	Polychaete	NA	256	41
<i>Spirobranchus cariniferus</i>	Serpulidae	Polychaete	NA	255	NA
<i>Spirobranchus giganteus</i>	Serpulidae	Polychaete	NA	256	41

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Table S10. Source studies of long-term demographic variability estimates included in the comparative analysis (n = 34) (See completed references list below).

Species Name	Habitat	Study Period	# Populations	Source ID
<i>Actaea spicata</i>	Terrestrial	1999 - 2005	2	1
<i>Alliaria petiolata</i>	Terrestrial	2003-2008	6	2
<i>Arabis fecunda</i>	Terrestrial	1987-1993, 2009	3	3, 4
<i>Astragalus scaphoides</i>	Terrestrial	1986-2009	3	3, 5
<i>Astragalus tyghensis</i>	Terrestrial	1991-1998, 2008	5	3, 6
<i>Cimicifuga elata</i>	Terrestrial	1992–1996, 1992–1997	3	3, 6
<i>Cirsium pitcheri</i>	Terrestrial	1998-2009	4	3, 7
<i>Cypripedium fasciculatum</i>	Terrestrial	1999-2007	3	3
<i>Dicerandra frutescens</i>	Terrestrial	1988-2008	6	3, 8, 9
<i>Eryngium cuneifolium</i>	Terrestrial	1990-2008	6	10
<i>Gentiana pneumonanthe</i>	Terrestrial	1987-1993, 2009	7	3, 11
<i>Haplopappus radiatus</i>	Terrestrial	1992-1999, 2009	4	3, 6
<i>Horkelia congesta</i>	Terrestrial	1994-1999, 2009	1	3, 12
<i>Hypericum cumulicola</i>	Terrestrial	1994-2007	6	3, 13
<i>Lathyrus vernus</i>	Terrestrial	1988-1991, 2009	10	3, 14
<i>Liatris scariosa</i>	Terrestrial	1995-2000, 2008	1	3
<i>Lomatium cookii</i>	Terrestrial	1994-1999, 2009	2	3, 6
<i>Lomatium bradshawii</i>	Terrestrial	1988–1994, 1996–1997	7	3, 6
<i>Neobuxbaumia macrocephala</i>	Terrestrial	1997-2002, 2009	1	3, 15, 16
<i>Opuntia macrorhiza</i>	Terrestrial	1999-2005	5	17
<i>Phyllanthus emblica</i>	Terrestrial	1999-2004, 2009	1	3
<i>Phyllanthus indofischeri</i>	Terrestrial	1999-2004, 2009	1	3
<i>Silene acaulis</i>	Terrestrial	1995-2000, 2009	4	3, 18
<i>Silene spaldingii</i>	Terrestrial	1987-2009	1	3, 19, 20
<i>Trillium grandiflorum</i>	Terrestrial	1998-2002, 2010	12	3, 21, 22
<i>Plexaura A</i>	Marine	1984-1987	1	23
<i>Corallium rubrum</i>	Marine	2003-2011	8	Present Study
<i>Montastrea annularis</i>	Marine	2001-2009	1	24

<i>Porites astreoides</i>	Marine	1999-2007	1	25
<i>Diploria strigosa</i>	Marine	2002-2007	1	25
<i>Acropora palmata</i>	Marine	2004-2010	1	26
<i>Paramuricea clavata</i>	Marine	1999-2003	1	27
<i>Mesodesma mactroides</i>	Marine	1983-1990	1	28
<i>Yoldia notabilis</i>	Marine	1973-1989, 1977-1989	2	29

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SUPPLEMENTARY METHODS

Integral Projection Model: data collection and model construction

*Demographic monitoring of natural *Corallium rubrum* populations*

A total of 1144 coral colonies from nine *C. rubrum* populations along the North-Western Mediterranean region were individually monitored using photographic techniques over periods ranging from 3 to 9 years between 2003 and 2011 (Montero-Serra *et al. in preparation*).

Six populations were located within Marine Protected Areas (MPA) where harvesting is not allowed but other human uses are allowed with some restrictions. Recreational diving is allowed in Parc Natural del Montgrí, Illes Medes i Baix Ter (Spain) and completely forbidden except for scientific studies in the Réserve Naturelle Marine Cerbère-Banyuls (France), in Parc Marine Côte Bleue (France), and the Réserve Naturelle de Scandola (France). The last three populations are located in the archipelago of Riou near Marseilles (France), where harvesting and recreational diving was allowed during the study (Fig. S1).

To individually identify and track each coral colony during the study period, two to four permanent transects were installed at each population. Transects were monitored once a year using a photographic sampling design. During sampling, a cord was deployed between permanently set screws and a 20 x 20 cm quadrat, sequentially positioned and photographed along the length of each transect. Two photographs were taken from each quadrat using different angles. This allowed us to obtain precise measurements of colony heights using photogrammetric techniques (Drap *et al.* 2013).

Estimation of vital rates

Vital rates (survival, growth, and reproduction) were estimated by fitting size-based continuous functions to long-term data on survival, post-recruit survival, growth and variance in growth (Easterling *et al.* 2000, Morris and Doak 2002). Survival was described by a logistic mixed effects model (GLMM) including colony size and year as fixed factors and population as a random factor.

Growth data from two populations over variable length periods (7 and 8 yrs) were used to fit a model characterizing annual growth rate depending on size. Processed height data from all annual photographs were not available, and growth is so slow in this species that even 7 to 8 year time periods only result in mean growth on the order of 10 to 40mm in maximum height ($2.74 \pm 0.19 \text{ mm} \cdot \text{yr}^{-1}$, see Fig. S2). As for many other clonal species, red coral can both grow and shrink over time. Shrinkage or negative growth is a common process in plants and sessile invertebrates that is usually

accounted for by assuming normally distributed variation in size changes around a mean rate (e.g., Easterling *et al.* 2000, Salguero-Gomez & Casper 2010). However, for many species assuming normally distributed growth variance is problematic, because a few individuals experience sporadic extreme shrinkage events and these results in a highly skewed distribution of growth rates (Shriver *et al.* 2012). We accounted for these two types of growth following the procedures proposed by Shriver and colleagues (2012), describing normal growth and extreme shrinkage as a two-part process. First, we estimated the probability of each colony undergoing “normal growth” (p_n) and “extreme shrinkage” ($1 - p_n$). These probabilities are size- and year-dependent according to the data.

For each type of growth (normal and extreme shrinkage) we first calculated mean annual change in size as.

$$G = (\text{Size}_T - \text{Size}_0) / T$$

where T is the time between the first and final census, Size_0 is the initial size measurement, and Size_T is the final size measurement.

We then extracted the residuals from the best-fitting model and divided the residuals by the number of years to predict annual variance in growth:

$$\text{VAR} = \{(\text{Residuals}) / T\}^2$$

We then used a linear regression model to fit the squared residuals to size, which showed a marginally significant positive relationship for normal growth ($P = 0.08$) and shrinkage ($P < 0.001$).

Recruitment rates were annually quantified from 30 fixed quadrats for each population. We considered effective recruits to be the button-like colonies observed for the first time in a quadrat. We calculated post-recruitment survival by monitoring recruits that we observed for the first time within a quadrat and tracked during the study period. Our data allowed for a robust estimation of survival within the range from 1 to 4 years. Further, we estimated the probability of recruits growing to an adult stage by direct observations in three populations where this process could be clearly identified.

Molecular studies have revealed a fine-scale genetic structure within *C. rubrum* populations and almost negligible dispersal among populations (Ledoux *et al.* 2010). Thus, new recruits were assigned to past individual parents assuming a relationship between colony size and the relative rate of reproduction. Past work has shown that there is an exponential relationship between *C. rubrum* size and polyp number (Santangelo *et al.* 2003), proportion of fertile colonies, and frequency of larvae per polyp in *C. rubrum* (Torrents *et al.* 2005). Thus, we used the squared adult colony size (height) as an approximation for polyp number. We then corrected these data by

the proportion of fertile colonies and percentage of fertile polyps, because both parameters have been shown to increase along with colony size (Torrents et al. 2005).

Fecundity was calculated as follows:

(1)

$$\text{Recruits/Size} = \text{New Recruits per quadrat (t1)} / \sum((\text{Colony Size})^2 * \text{Colony Fertility} * \text{Polyp Fertility}) (\text{t0}) \text{ per quadrat}$$

(2)

$$\text{Fecundity values} = [(\text{Adult Colony Size})^2] * (\text{Colony Fertility} * \text{Polyp Fertility}) * (\text{Rec/Size})$$

Model construction

We coupled continuous size-based vital rates estimates for adults to four age-based stages for recruits to build an IPM for red coral. We simulated the model using 60 classes (where the first four classes are age-based categories for recruits and the rest are size-based). The number of size classes was chosen after calculating asymptotic population growth under multiple matrices constructed using a range of size-classes from 3 to 200. The results of these simulations showed that matrices with a number of size-classes larger than 60 showed a consistent value for population growth rate (Fig S2).

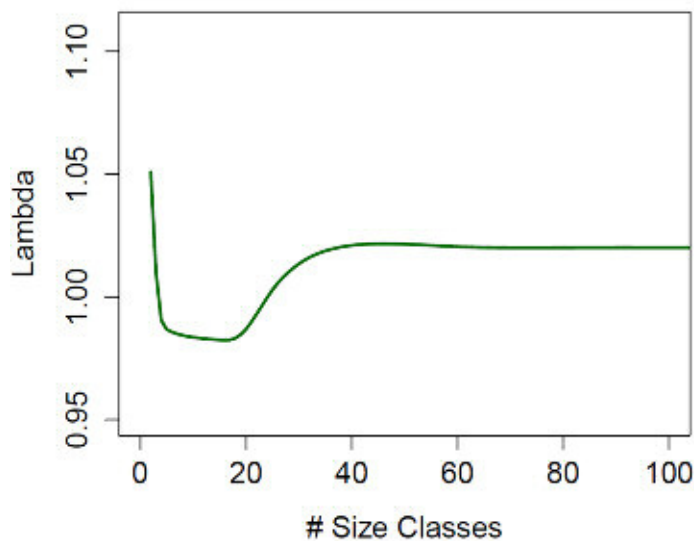


Figure S2. Asymptotic population growth rate (lambda) based on matrix approximations to a continuous IPM model constructed using a range of size-classes from 3 to 200.

We parameterized eight annual matrices that were derived based on the models previously fit; the fecundity and survival functions varied between these years, while growth was not estimated on an annual basis. All the a_{ji} matrix elements below the top row (the reproductive elements) were estimated as:

(3)

$$a_{ji} = s_i (p_{n,i} n_{ji} + (1 - p_{n,i}) m_{ji})$$

where s_i is the probability of survival for the i th size class, $p_{n,i}$ is the probability of “normal growth or shrinkage” for the i th class, n_{ji} is the probability of normally growing or shrinking from the i th class to the j th class, and m_{ji} is the probability of moving from the i th to the j th class with extreme shrinkage. To estimate n_{ji} and m_{ji} values, we used the mid-point size for class i as the starting size, and a normal CDF to calculate the total probability of falling within the range of sizes in the j th class.

References included in the supplementary methods

1. Drap, P., Merad, D., Mahiddine, A., *et al.* (2013) Automating the measurement of red coral in situ using underwater photogrammetry and coded targets. XXIV International CIPA Symposium, Strasbourg, France. Vol. XL-5/W2, pp.231–236
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3. Morris, W.F. & Doak, D.F. (2002) Quantitative Conservation Biology: Theory and Practice of Population Viability Analysis. Sinauer Associates, Inc. Publishers; Sunderland, Massachusetts U.S.A.
4. Torrents, O., Garrabou, J., Marschal, C. & Harmelin, J. G. (2005) Age and size at first reproduction in the commercially exploited red coral *Corallium rubrum* (L.) in the Marseilles area (France, NW Mediterranean). *Biol. Conserv.*, 121(3) 391–397
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6. Santangelo, G., Carletti, E., Maggi, E. & Bramanti, L. (2003) Reproduction and population sexual structure of the overexploited Mediterranean red coral *Corallium rubrum*. *Mar. Eco. Prog. Ser.*, 248: 99–108, 2003
7. Shriver, R. K., Cutler, K., & Doak, D. F. (2012). Comparative demography of an epiphytic lichen: support for general life history patterns and solutions to common problems in demographic parameter estimation. *Oecologia*, 170(1), 137-146.

Supplementary Materials Chapter 2

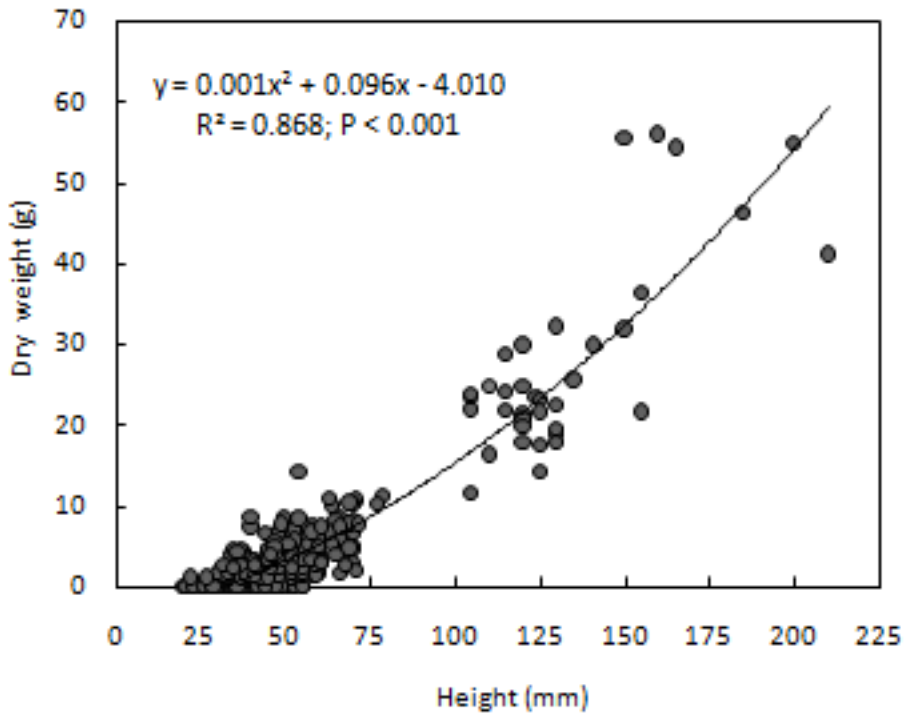


Figure S1. Height – weight relationship obtained from death corals (n = 300).

Supplementary Materials Chapter III

Table S1. Mediterranean red coral populations surveyed annually.

Table S2. Logistic Generalized Mixed Effects Models fitted to long-term demographic data on annual colony survival rates.

Table S3. Logistic Generalized Mixed Effects Models fitted to long-term demographic data on annual colony shrinkage rates.

Figure S1. Local disturbance levels used to simulate the interactive effects of warming and marine protected areas. To account for a continuous gradient of local disturbance intensity in Figure 5, we computed ten levels of local disturbance intensity by multiplying the protection factor in the shrinkage probability GLMM (Table S3) by ten levels. The maximum intensity (black line) corresponds to shrinkage probability outside MPAs, while the minimum probability (light grey) corresponds to shrinkage rates observed in populations within MPAs.

Figure S2. Annual transition probabilities of large red coral colonies (<100mm in height) in protected and unprotected populations. Large red coral colonies can survive and stay in the same size-class (stasis) or survive but transit to smaller size-classes due to partial mortality events (shrinkage). A total of 36 annual and

Table S1. Mediterranean red coral populations surveyed annually.

Name	Depth (m)	Area	Protection Level	N	Years Surveyed
Dofi	17-22	Catalan Sea (Spain)	Partial Protection	148	2006 to 2011
Reina	17-24	Catalan Sea (Spain)	Partial Protection	140	2007 to 2011
Banyuls	23 - 25	Catalan Sea (Spain)	Integral Protection	108	2005 to 2007
Maire	15 - 18	Marseille (France)	Unprotected	140	2002 to 2009
Riou	17-24	Marseille (France)	Unprotected	204	2005 to 2009
Plane	17-24	Marseille (France)	Unprotected	154	2003 to 2010
Passe	24 – 27	Corsica (France)	Integral Protection	145	2006 to 2016
Palazzu	19 – 22	Corsica (France)	Integral Protection	137	2003 to 2016

Table S2. Logistic Generalized Mixed Effects Models fitted to long-term demographic data on annual colony survival rates.

Model	AICc	Δ AICc	Parameter	Estimate	P-value
Survival ~ Colony Size + (1 Population) + (1 Year)	923.5	0	Intercept	4.360851 \pm 0.508109	<0.0001
			Colony Size	0.015677 \pm 0.003904	<0.0001
Survival ~ Colony Size + MPA + (1 MPA/Population) + (1 Year)	925.3	1.8	Intercept	4.177284 \pm 0.675142	<0.0001
			Colony Size	0.015572 \pm 0.003917	<0.0001
			MPA	0.316082 \pm 0.769349	0.6810
Survival ~ Colony Size + MPA + Colony Size*MPA + (1 MPA/Population) + (1 Year)	927.3	3.8	Intercept	4.182451 \pm 0.76020	<0.0001
			Colony Size	0.015398 \pm 0.01244	0.2161
			MPA	0.310012 \pm 0.87232	0.7223
			Colony Size*MPA	0.000193 \pm 0.01307	0.9881

Table S3. Logistic Generalized Mixed Effects Models fitted to long-term demographic data on annual colony shrinkage rates.

Model	AIC	Δ AIC	Parameter	Estimate (\pm SE)	P-value
Shrinkage ~ Colony Size + MPA + Colony Size*MPA + (1 MPA/Population) + (1 Year)	1928.5	0	Intercept	-4.508317 \pm 0.599878	<0.0001
			Colony Size	0.046478 \pm 0.004421	<0.0001
			MPA	-0.580281 \pm 0.763198	0.447
			Colony Size*MPA	-0.040954 \pm 0.005398	<0.0001
Shrinkage ~ Colony Size + MPA + (1 MPA/Population) + (1 Year)	1987.9	59	Intercept	-3.457779 \pm 0.458558	<0.0001
			Colony Size	0.020850 \pm 0.002419	<0.0001
			MPA	-2.542601 \pm 0.576368	<0.0001
Shrinkage ~ Colony Size + (1 Population) + (1 Year)	1994.5	66	Intercept	-5.319992 \pm 0.748843	<0.0001
			Colony Size	0.020597 \pm 0.002444	<0.0001

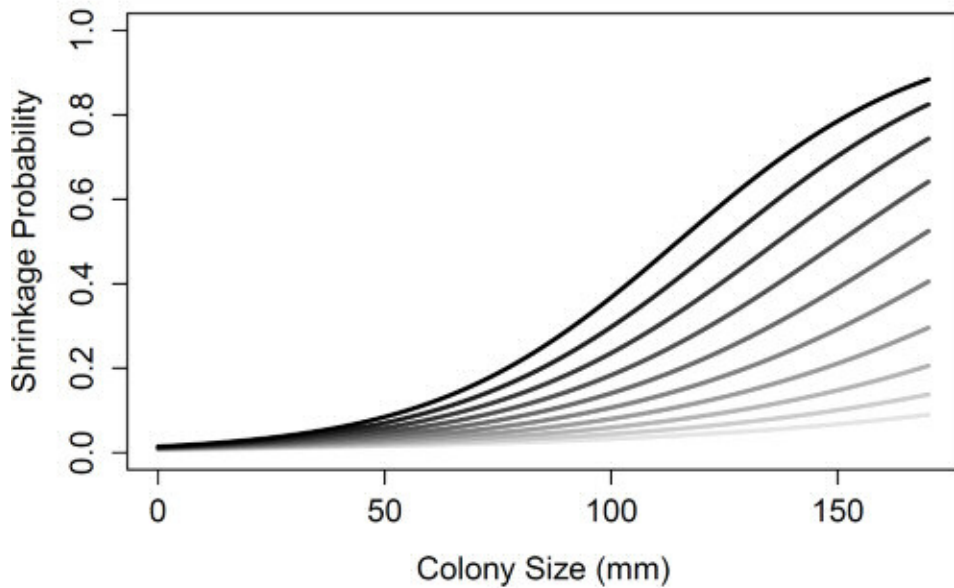


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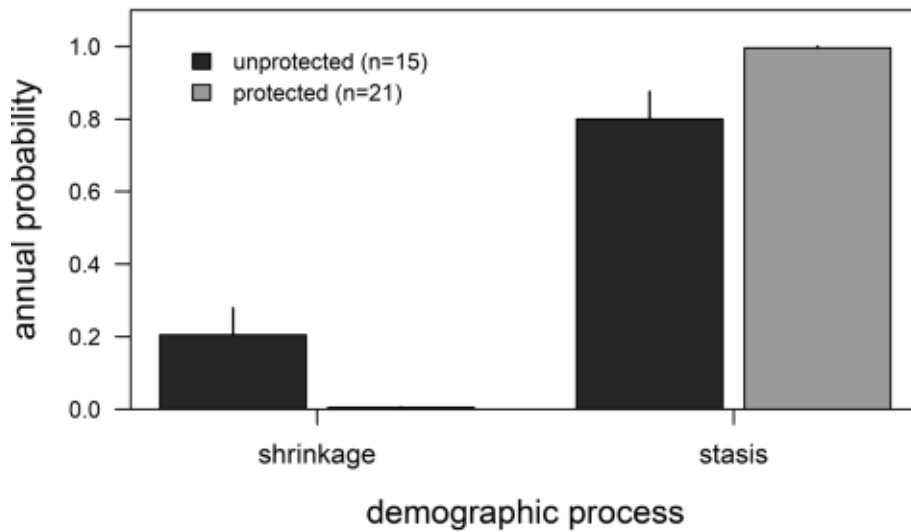


Figure S2. Annual transition probabilities of large red coral colonies (<100mm in height) in protected and unprotected populations. Large red coral colonies can survive and stay in the same size-class (stasis) or survive but transit to smaller size-classes due to partial mortality events (shrinkage). A total of 36 annual and population level matrices were constructed from field demographic data on four protected and three unprotected red coral populations in the NW Mediterranean Sea.

Supplementary Materials Chapter IV

Figure S1. Population size frequency distribution of the transplanted colonies represented by the brown bars and the brown line. Green lines represent the size-structure of three mature and well-preserved populations based on data from Linares *et al.* (2010): Carry, Banyuls, and Palazzu. In order to aid data visualization, we fitted a cubic smoothing spline to the colony-size frequency data of each population using the *Smth.spline()* function in R.

Figure S2. Size-dependent growth of *C. rubrum* colonies based on data from two natural populations.

Figure S3. Growth rates in natural vs. transplanted coral colonies (log-scale) for 17 coral species.

Supplementary Methods. Full description of the demographic models.

Table S1. Summary statistics for the linear model fitted to the expected recovery periods.

Table S2. Source data of survival in transplant experiments and growth references of the marine sessile species included in the comparative analysis.

Table S3. Source data of population matrix models used to project potential recovery periods.

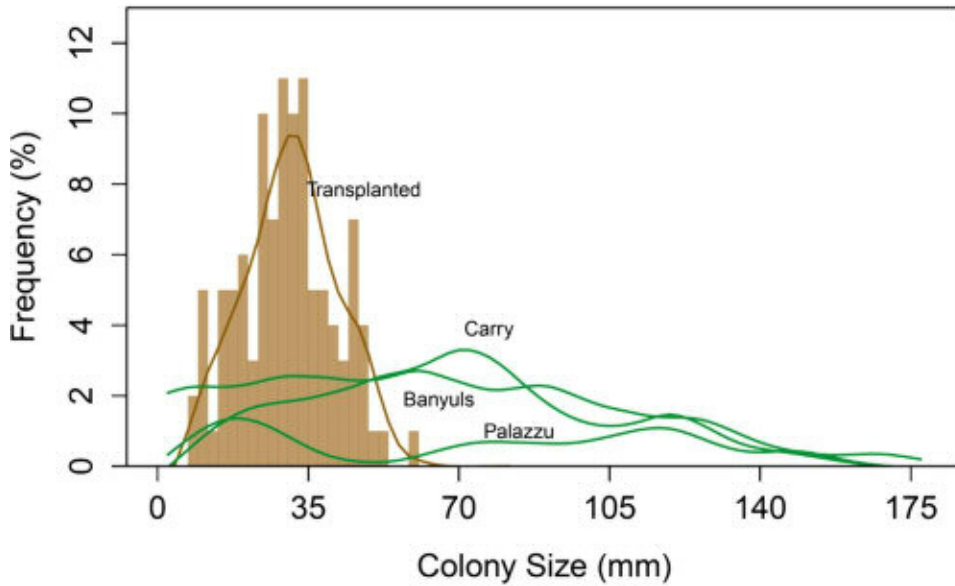


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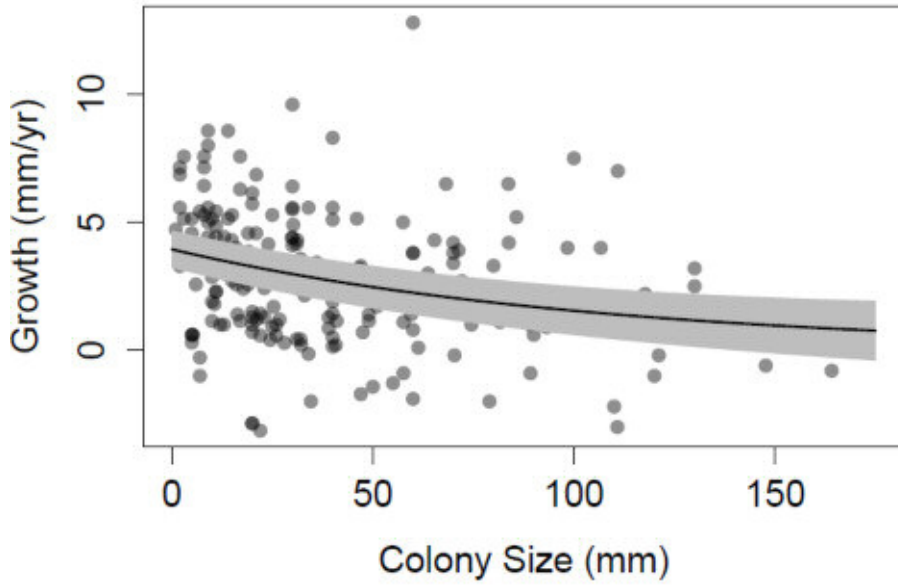


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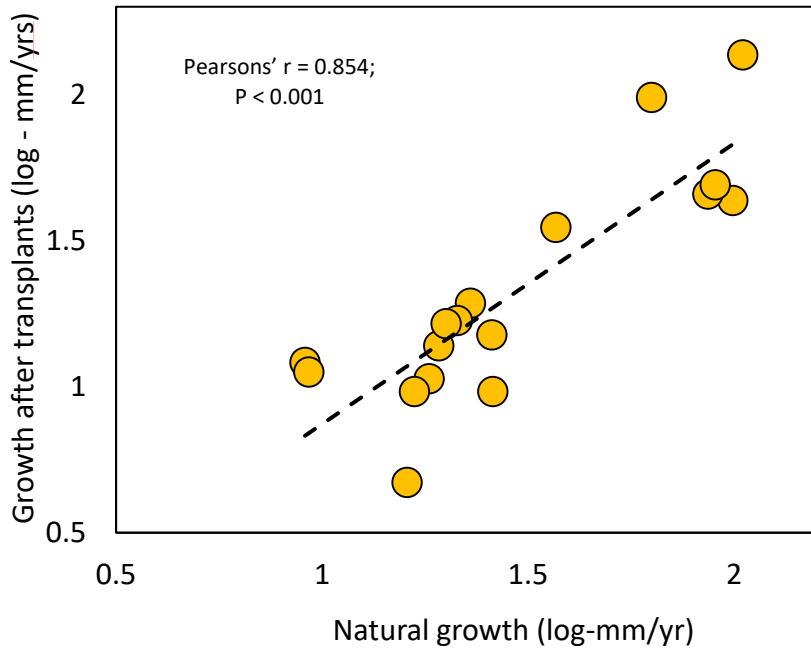


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Supplementary Methods Chapter IV

Integral Projection Model: data collection and model construction

*Demographic monitoring of natural *Corallium rubrum* populations*

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To individually identify and track each coral colony during the study period, two to four permanent transects were installed at each population. Transects were monitored once a year using a photographic sampling design. During sampling, a cord was deployed between permanently set screws and a 20 x 20 cm quadrat, sequentially positioned and photographed along the length of each transect. Two photographs were taken from each quadrat using different angles. This allowed us to obtain precise measurements of colony heights using photogrammetric techniques (Drap *et al.* 2013).

Estimation of vital rates

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For each type of growth (normal and extreme shrinkage) we first calculated mean annual growth as.

$$G = (\text{Size}_T - \text{Size}_0) / T$$

where T is the time between the first and final census, Size_0 is the initial size measurement, and Size_T is the final size measurement.

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$$\text{VAR} = \{(\text{Residuals}) / T\}^2$$

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Fecundity was calculated as follows:

(1)

$$\text{Recruits/Size} = \text{New Recruits per quadrat (t1)} / \sum((\text{Colony Size})^2 * \text{Colony Fertility} * \text{Polyp Fertility}) (\text{t0}) \text{ per quadrat}$$

(2)

$$\text{Fecundity values} = [(\text{Adult Colony Size})^2] * (\text{Colony Fertility} * \text{Polyp Fertility}) * (\text{Rec/Size})$$

Model construction

We coupled continuous size-based vital rates estimates for adults to four age-based stages for recruits to build an IPM for red coral. We simulated the model using 60 classes (where the first four classes are age-based categories for recruits and the rest are size-based). The number of size classes was chosen after calculating asymptotic population growth under multiple matrices constructed using a range of size-classes from 3 to 200. The results of these simulations showed that matrices with a number of size-classes larger than 60 showed a consistent value for population growth rate (Fig S2).

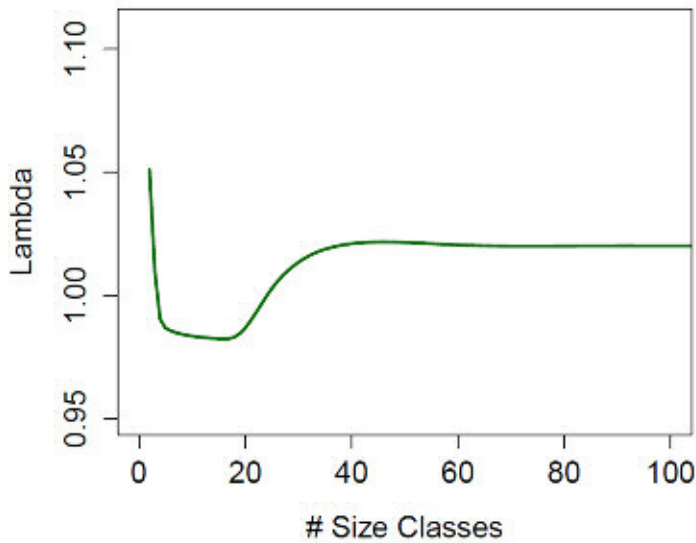


Figure S4. Asymptotic population growth rate (lambda) based on matrix approximations to a continuous IPM model constructed using a range of size-classes from 3 to 200.

We parameterized eight annual matrices that were derived based on the models previously fit; the fecundity and survival functions varied between these years, while growth was not estimated on an annual basis. All the a_{ji} matrix elements below the top row (the reproductive elements) were estimated as:

(3)

$$a_{ji} = s_i (p_{n,i} n_{ji} + (1 - p_{n,i}) m_{ji})$$

where s_i is the probability of survival for the i th size class, $p_{n,i}$ is the probability of “normal growth or shrinkage” for the i th class, n_{ji} is the probability of normally

growing or shrinking from the i th class to the j th class, and m_{ji} is the probability of moving from the i th to the j th class with extreme shrinkage. To estimate n_{ji} and m_{ji} values, we used the mid-point size for class i as the starting size, and a normal CDF to calculate the total probability of falling within the range of sizes in the j th class.

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TABLE S1. Summary statistics for the linear models fitted to the expected recovery periods ($n = 41$). A log transformation was previously applied to the response variable.

Model	Threshold at 20%					Threshold at 80%				
	Estimate	DF	t	P	R ²	Estimate	DF	t	P	R ²
Survival + Dimensions	2.50 ± 0.55	38	4.56	> 0.001	0.42	3.01 ± 0.53	38	5.63	> 0.001	0.50
(Dimensions)	0.03 ± 0.01	38	2.93	0.029	0.42	0.03 ± 0.01	38	1.98	0.055	0.50
Survival*Dimensions	-	-	-	0.274	-	-	-	-	0.243	-
Survival	2.73 ± 0.57	39	4.82	> 0.001	0.36	3.21 ± 0.43	39	5.44	> 0.001	0.46
Dimensions	0.05 ± 0.02	39	2.57	0.014	0.12	0.04 ± 0.02	39	2.30	0.027	0.10

TABLE S2. Source data of survival rates in transplant experiments and growth rates of the marine sessile species included in the comparative analysis. Numbers correspond to references listed below the table.

Species Name	Family	Taxa	Survival Transplants	Growth Natural	Growth Transplants
<i>Acropora cervicornis</i>	<i>Acroporidae</i>	Scleractinia	1, 2, 3, 4, 94	41, 42, 43, 44	92, 93, 94
<i>Acropora digitifera</i>	<i>Acroporidae</i>	Scleractinia	5	61	NA
<i>Acropora divaricata</i>	<i>Acroporidae</i>	Scleractinia	6	NA	6
<i>Acropora echinata</i>	<i>Acroporidae</i>	Scleractinia	88	88	88
<i>Acropora abrolhosensis</i> / <i>A. gomezi</i>	<i>Acroporidae</i>	Scleractinia	87, 90	NA	90
<i>Acropora hemprichii</i>	<i>Acroporidae</i>	Scleractinia	8, 9, 10	49, 50	NA
<i>Acropora humilis</i>	<i>Acroporidae</i>	Scleractinia	6, 9	47, 52, 54	6
<i>Acropora hyacinthus</i>	<i>Acroporidae</i>	Scleractinia	5, 6, 11, 53	51	6
<i>Acropora intermedia</i> / <i>A. nobilis</i>	<i>Acroporidae</i>	Scleractinia	11, 53	52	NA
<i>Acropora micropthalma</i>	<i>Acroporidae</i>	Scleractinia	87	NA	NA
<i>Acropora millepora</i>	<i>Acroporidae</i>	Scleractinia	12,53	NA	NA
<i>Acropora muricata</i> / <i>A. formosa</i>	<i>Acroporidae</i>	Scleractinia	7, 8, 10, 20, 111	46, 47, 48	99
<i>Acropora nasuta</i>	<i>Acroporidae</i>	Scleractinia	8, 10	47, 51	10
<i>Acropora palmata</i>	<i>Acroporidae</i>	Scleractinia	1, 2, 13, 14, 15, 16	47, 55, 56, 57, 58, 97	97
<i>Acropora pulchra</i>	<i>Acroporidae</i>	Scleractinia	NA	114	114
<i>Acropora tenuis</i>	<i>Acroporidae</i>	Scleractinia	17, 18	NA	NA

<i>Acropora variabilis</i>	<i>Acroporidae</i>	Scleractinia	19	NA	NA
<i>Acropora verweyi</i>	<i>Acroporidae</i>	Scleractinia	9	54	NA
<i>Acropora yongei</i>	<i>Acroporidae</i>	Scleractinia	20	98, 99	NA
<i>Corallium rubrum</i>	<i>Coralliidae</i>	Octocorallia	Present Study	Present Study	NA
<i>Echinopora lamellosa</i>	<i>Mussidae</i>	Scleractinia	24, 111	NA	NA
<i>Eunicella cavolini</i>	<i>Gorgoniidae</i>	Octocorallia	115	NA	NA
<i>Eunicella singularis</i>	<i>Gorgoniidae</i>	Octocorallia	115	116, 117, 118	NA
<i>Eunicella verrucosa</i>	<i>Gorgoniidae</i>	Octocorallia	115	119,120	NA
<i>Favia stelligera</i>	<i>Mussidae</i>	Scleractinia	9	62	NA
<i>Isopora brueggemanni</i>	<i>Acroporidae</i>	Scleractinia	20	63	NA
<i>Leptoseris gardineri</i>	<i>Agariciidae</i>	Scleractinia	88	88	88
<i>Madracis mirabilis</i>	<i>Astrocoeniidae</i>	Scleractinia	27	NA	NA
<i>Merulina scabricula</i>	<i>Merulinidae</i>	Scleractinia	24, 111	NA	NA
<i>Montipora aequituberculata</i>	<i>Acroporidae</i>	Scleractinia	111	113	NA
<i>Montipora digitata</i>	<i>Acroporidae</i>	Scleractinia	28, 29, 111, 112	100	28
<i>Montipora foliosa</i>	<i>Acroporidae</i>	Scleractinia	87	63	NA
<i>Montipora pulcherrima</i>	<i>Acroporidae</i>	Scleractinia	88	NA	88
<i>Paramuricea clavata</i>	<i>Plexauridae</i>	Octocorallia	30, 115	67, 121	NA
<i>Pavona varians</i>	<i>Agariciidae</i>	Scleractinia	19	68, 71, 73	NA
<i>Pavona cactus</i>	<i>Agariciidae</i>	Scleractinia	87, 88	88, 89	88
<i>Pocillopora damicornis</i>	<i>Pocilloporidae</i>	Scleractinia	9, 19, 31, 111	51, 68, 69, 70	NA
<i>Pocillopora verrucosa</i>	<i>Pocilloporidae</i>	Scleractinia	6, 8, 10, 20	51, 63, 73	NA

<i>Porites astreoides</i>	<i>Poritidae</i>	Scleractinia	32	73, 74	NA
<i>Porites cylindrica</i>	<i>Poritidae</i>	Scleractinia	8, 10, 33, 87	75,61	76, 91
<i>Porites lobata</i>	<i>Poritidae</i>	Scleractinia	6	77, 78, 79	6
<i>Porites lutea</i>	<i>Poritidae</i>	Scleractinia	6	80, 81	6
<i>Porites nigrescens</i>	<i>Poritidae</i>	Scleractinia	6	NA	6
<i>Porites porites</i>	<i>Poritidae</i>	Scleractinia	1, 2	82, 83, 84, 85	97
<i>Porites rus</i>	<i>Poritidae</i>	Scleractinia	111	76	76
<i>Stylophora pistillata</i>	<i>Pocilloporidae</i>	Scleractinia	9,19,31,40	101, 102, 103	104
<i>Xestospongia muta</i>	<i>Petrosiidae</i>	Sponge	41	86	NA
<i>Cymodocea nodosa</i>	<i>Cymodoceaceae</i>	Seagrass	21, 22	105	NA
<i>Halodule wrightii</i>	<i>Cymodoceaceae</i>	Seagrass	25, 110	105	NA
<i>Posidonia australis</i>	<i>Posidoniaceae</i>	Seagrass	26, 34	105	NA
<i>Posidonia oceanica</i>	<i>Posidoniaceae</i>	Seagrass	26, 35, 36, 37, 108	105	NA
<i>Posidonia sinuosa</i>	<i>Posidoniaceae</i>	Seagrass	26, 106	105	NA
<i>Ruppia maritima</i>	<i>Ruppiceae</i>	Seagrass	38	NA	NA
<i>Syringodium filiforme</i>	<i>Cymodoceaceae</i>	Seagrass	25	105	NA
<i>Thalassia testudinum</i>	<i>Hydrocharitaceae</i>	Seagrass	25	105	NA
<i>Zostera capricorni</i>	<i>Zosteraceae</i>	Seagrass	26	NA	NA
<i>Zostera marina</i>	<i>Zosteraceae</i>	Seagrass	38, 109	105	NA
<i>Zostera noltii</i>	<i>Zosteraceae</i>	Seagrass	107	NA	NA
<i>Cystoseira barbata</i>	<i>Fucaceae</i>	Macroalgae	23	NA	NA
<i>Cystoseira compressa</i>	<i>Fucaceae</i>	Macroalgae	23	NA	NA

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TABLE S3. Source data of matrix population models used to project recovery periods. Numbers correspond to references listed below the table. *Only matrices that included fecundity data (n = 41) were used to project recovery periods.

Name	Family	Taxa	Matrix Dimensions	Source ID
<i>Acropora cervicornis</i>	Acroporidae	Hexacoral	3	1
<i>Acropora hyacinthus</i>	Acroporidae	Hexacoral	4	2
<i>Acropora palmata</i>	Acroporidae	Hexacoral	4	3
<i>Adamussium colbecki</i>	Pectinidae	Bivalve	3	16
<i>Agaricia agaricites</i>	Agariciidae	Hexacoral	3	4
<i>Alaria nana</i>	Alariaceae	Algae	3	17
<i>Amphimedon compressa</i>	Chalinidae	Sponge	3	5
<i>Artica islandica</i>	Arcticidae	Bivalve	3	16
<i>Ascophyllum nodosum</i>	Fucaceae	Algae	5	18
<i>Bartholomea annulata</i>	Aiptasiidae	Hexacoral	5	19
<i>Botrylloides schlosseri</i>	Styelidae	Tunicate	3	20
<i>Botrylloides violaceous</i>	Styelidae	Tunicate	3	20
<i>Celleporella pumicosa</i>	Celleporidae	Bryozoan	3	22
<i>Chthamalus montagui</i>	Chthamalidae	Cirriped	4	21
<i>Corallium rubrum</i>	Coralliidae	Octocoral	60	Present Study
<i>Diploria strigosa</i>	Mussidae	Hexacoral	3	6
<i>Fucus distichus</i>	Fucaceae	Algae	9	23
<i>Gemma gemma</i>	Veneridae	Bivalve	3	16
<i>Geukensia demissa</i>	Mytilidae	Bivalve	3	16
<i>Goniastrea aspera</i>	Merulinidae	Hexacoral	5	7
<i>Goniastrea favulus</i>	Merulinidae	Hexacoral	5	7
<i>Goniastrea favulus</i>	Merulinidae	Hexacoral	5	7
<i>Haliotis iris</i>	Haliotidae	Bivalve	3	24
<i>Haliotis laevigata</i>	Haliotidae	Bivalve	7	33, 34
<i>Haliotis rufescens</i>	Haliotidae	Bivalve	4	25
<i>Haliotis sorenseni</i>	Haliotidae	Bivalve	8	25
<i>Iridaea splendens</i>	Gigartiniaceae	Algae	8	26
<i>Laminaria digitata</i>	Laminariaceae	Algae	5	27
<i>Lasaea rubra</i>	Lasaeidae	Bivalve	3	16

<i>Leptogorgia Virgulata</i>	Gorgoniidae	Octocoral	5	8
<i>Leptoseris cucullata</i>	Agariciidae	Hexacoral	3	9
<i>Lissarca miliaris</i>	Philobryidae	Bivalve	3	16
<i>Lissarca notorcadensis</i>	Philobryidae	Bivalve	3	16
<i>Macrocystis pyrifera</i>	Laminariaceae	Algae	8	30
<i>Mesodesma mactroides</i>	Mesodesmatidae	Bivalve	5	29
<i>Mucropetraliella ellerii</i>	Petraliellidae	Bryozoan	5	10
<i>Mya arenaria</i>	Myidae	Bivalve	8	28
<i>Mytilus californianus</i>	Mytilidae	Bivalve	3	35
<i>Mytilus galloprovincialis</i>	Mytilidae	Bivalve	3	35
<i>Nutallia obscurata</i>	Psammobiidae	Bivalve	5	31
<i>Orbicella annularis</i>	Merulinidae	Hexacoral	3	9
<i>Palythoa caesia</i>	Sphenopidae	Hexacoral	4	11
<i>Panopea abrupta</i>	Hiatellidae	Bivalve	3	16
<i>Paramuricea clavata</i>	Plexaura A	Octocoral	7	12
<i>Plexaura A</i>	Plexaura A	Octocoral	3	13
<i>Pocillopora damicornis</i>	Pocilloporidae	Hexacoral	4	2
<i>Goniastrea sinensis</i>	Merulinidae	Hexacoral	5	8
<i>Porites astreoides</i>	Poritidae	Hexacoral	3	6
<i>Pseudopterogorgia</i>	Gorgoniidae	Octocoral	13	14
<i>Semibalanus balanoides</i>	Archaebalanidae	Cirriped	3	21
<i>Spongia graminea</i>	Spongiidae	Sponge	10	32
<i>Tridacna gigas</i>	Cardiidae	Bivalve	3	16
<i>Watersipora subtorquata</i>	Watersiporidae	Bryozoan	3	20
<i>Xestospongia muta</i>	Petrosiidae	Sponge	6	15

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Supplementary Materials Chapter V

Table S1. Source data of distribution and depths of *Corallium rubrum* and *Paramuricea clavata*.

Table S2. Sample sizes of distribution and depth data for *Corallium rubrum* and *Paramuricea clavata* populations.

Table S3. Source data for environmental predictors included in the models.

Table S4. Collinearity among environmental predictors

Table S5. Logistic generalized additive mixed models fitted to presence/absence data for *Paramuricea clavata* and *Corallium rubrum* at five depth layers (from 5-15, 15-25, 25-35, 35-45, and 45-55).

Table S6. Clusters classification according to projected trajectories of change.

Figure S1. Correlations among environmental predictors

Figure S1. Examples of model validation comparing predicted probability of occurrence with independent presence and absence data.

Figure S2. Model validation results. Adjusted R squared values among 50 permutations comparing predicted probability of occurrence against independent presence and absence data for *C. rubrum* and *P. clavata*.

Figure S3. Projected changes by species.

Table S1. Sources of geographic distribution data of *Paramuricea clavata* and *Corallium rubrum* populations.

Source Type	Project name (website)	Reference
Citizen Science Program	ReefCheck (reefcheckmed.org)	Cerrano et al. 2017
	Cigesemed (cs.cigesmed.eu)	Gerovasileiou et al. 2016
	SeaWatchers (observadoresdelmar.es)	NA
Literature Review	NA	Di Camilo et al. 2018
Literature Review	NA	Boavida et al. 2016

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Table S2. Total geographic and depth distribution data for *Corallium rubrum* and *Paramuricea clavata*. Filtering was done by randomly removing all duplicated data points and leaving only one for each location.

Species	PRESENCES						Filtered absences	TOTAL
	All data	Filtered presences*	No depth data	Single depth observation	Range of depths	Upper limit known		
Paramuricea clavata	1769	1019	199	242	186	392	199	1769
Corallium rubrum	1030	645	92	160	128	265	166	1277

Table S3. Total geographic and depth distribution data for *Corallium rubrum* and *Paramuricea clavata*.

Variable	Spatial Resolution	Source
Mean Annual Nitrate Concentration	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Mean Annual Phosphate Concentration	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Mean Annual Net Primary Productivity	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Annual Mean Temperature	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Mean August Temperature	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Mean September Temperature	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Max August Temperature	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Max September Temperature	0.083° x 0.083°	Ocean model (Macías et al. 2015)
Concavity	0.0083 x 0.0083°	MARSPEC (Sbrocco et al. 2013)
Slope	0.0083 x 0.0083°	MARSPEC (Sbrocco et al. 2013)
NS Exposure	0.0083 x 0.0083°	MARSPEC (Sbrocco et al. 2013)
EW Exposure	0.0083 x 0.0083°	MARSPEC (Sbrocco et al. 2013)

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Table S4. Collinearity among environmental predictors

	August Mean T ^a	August Max T ^a	September Max T ^a	Annual Mean T ^a	Annual Mean PPR	Annual Mean Pho
August Mean T ^a	0.963958					
August Max T ^a	0.937181	0.908165				
September Max T ^a	0.875857	0.803242	0.901352			
Annual Mean T ^a	-0.37764	-0.38413	-0.38215	-0.19237		
Annual Mean PPR	-0.35813	-0.34841	-0.39383	-0.25247	0.856813	
Annual Mean Pho	0.10049	0.06259	0.098412	0.122621	0.27999	0.386776

Table S5. Logistic generalized additive mixed models fitted to presence/absence data for *Paramuricea clavata* and *Corallium rubrum* at five depth layers (from 5-15, 15-25, 25-35, 35-45, and 45-55).

Species	Forumula	Adjusted R ²	Sample Size
<i>P. clavata</i>	Presence ~ Depth	0.344	n = 1198
	Presence ~ PPR	0.0589	
	Presence ~ Annual T ^a	0.268	
	Presence ~ August T ^a	0.343	
	Presence ~ Annual T ^a + Depth	0.413	
	Presence ~ August T ^a + Depth	0.342	
	Presence ~ August T ^a + PPR	0.353	
	Presence ~ Annual T ^a + PPR	0.266	
	Presence ~ August T^a + PPR + Depth	0.402	
	Presence ~ Annual T^a + PPR + Depth	0.430	
<i>C. rubrum</i>	Forumula	Adjusted R²	n = 497
	Presence ~ Depth	0.257	
	Presence ~ PPR	0.349	
	Presence ~ Annual T ^a	0.418	
	Presence ~ August T ^a	0.347	
	Presence ~ Annual T ^a + Depth	0.333	
	Presence ~ August T ^a + Depth	0.226	
	Presence ~ August T ^a + PPR	0.238	
	Presence ~ Annual T ^a + PPR	0.341	
	Presence ~ August T^a + PPR + Depth	0.404	
Presence ~ Annual T^a + PPR + Depth	0.424		

Table S6. Number of sites classified in each cluster (K-means analysis) for each species and emission's scenario.

Species	Scenario	Clusters according trajectories of change							
		A	B	C	D	E	F	G	H
<i>C. rubrum</i>	RCP 4.5	35	68	193	60	19	138	50	30
	RCP 8.5	1	68	1	33	34	210	115	131
<i>P. clavata</i>	RCP 4.5	113	109	246	101	49	335	2	0
	RCP 8.5	4	109	97	15	228	487	15	0

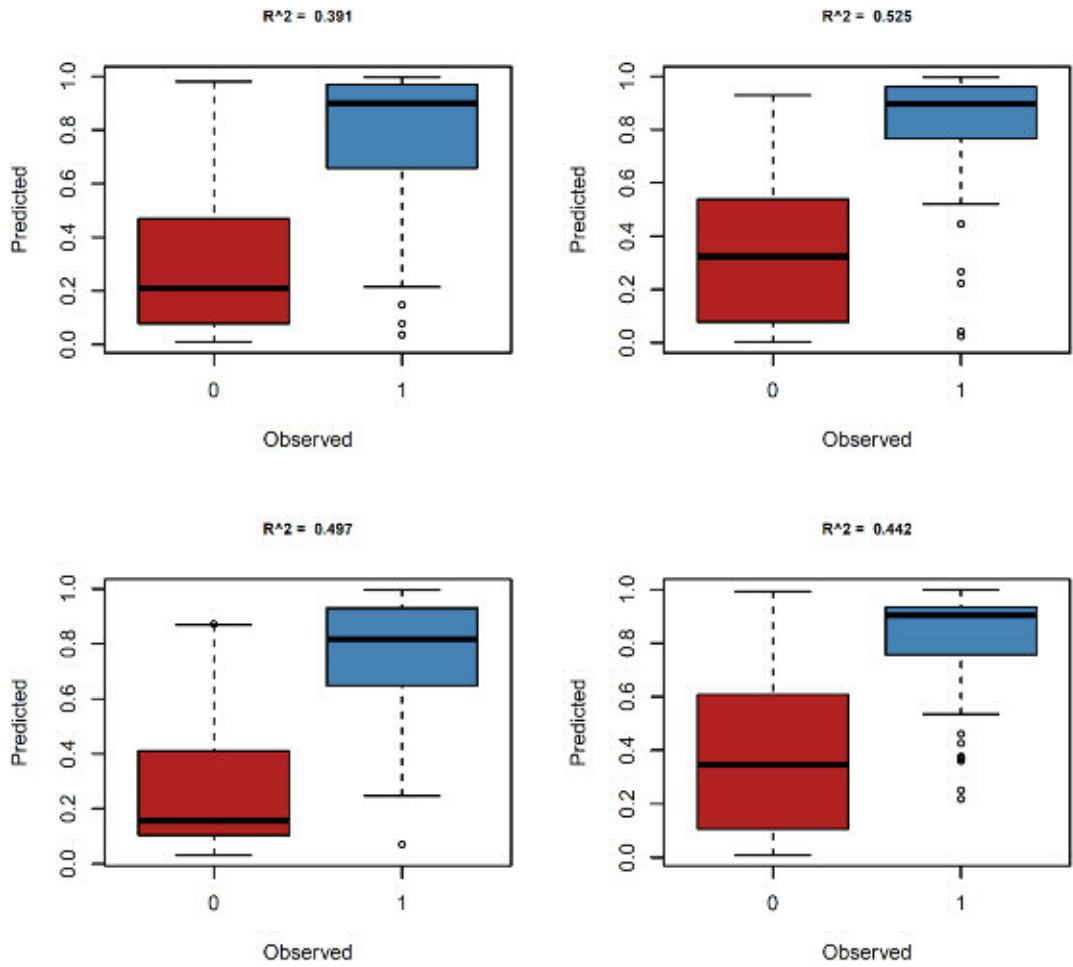


Figure S1. Examples of model validation comparing predicted probability of occurrence with independent presence and absence data.

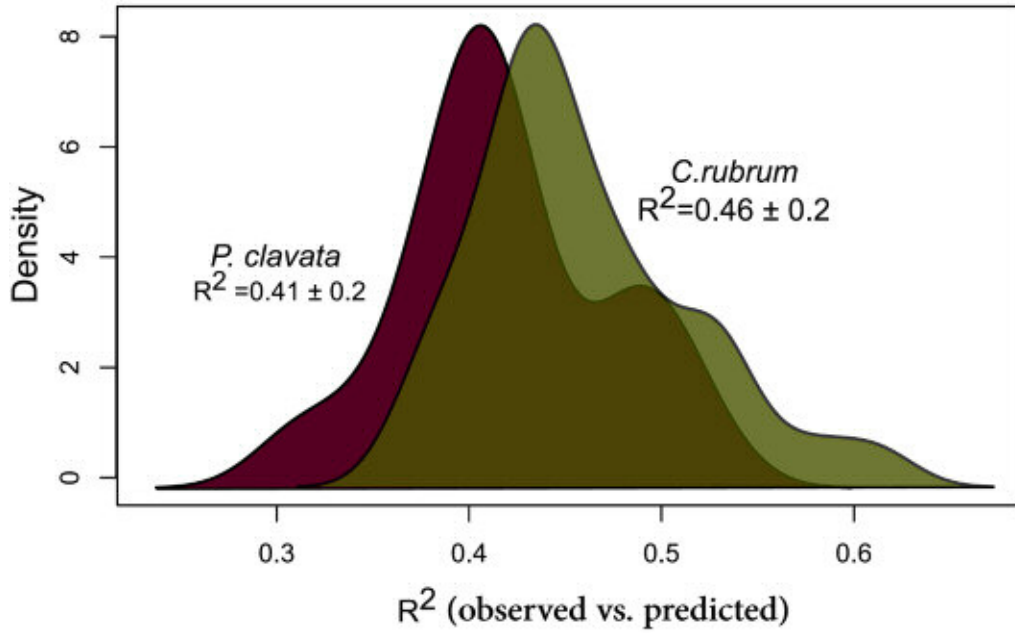


Figure S2. Model validation results. Adjusted R squared values among 50 permutations comparing predicted probability of occurrence against independent presence and absence data for *C. rubrum* and *P. clavata*.

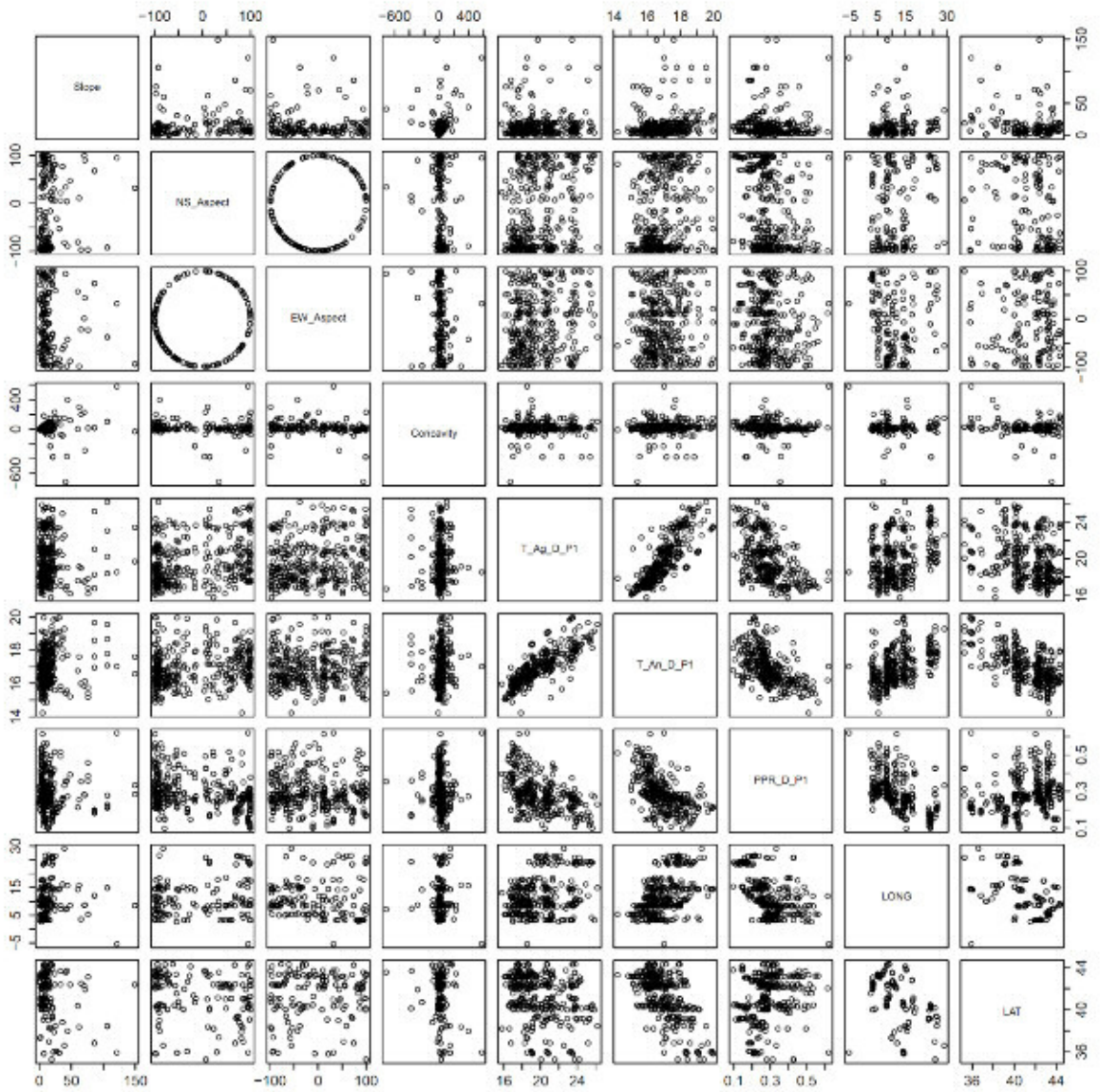


Figure S3. Correlations between environmental predictors. Only annual and August temperature showed a high correlation.

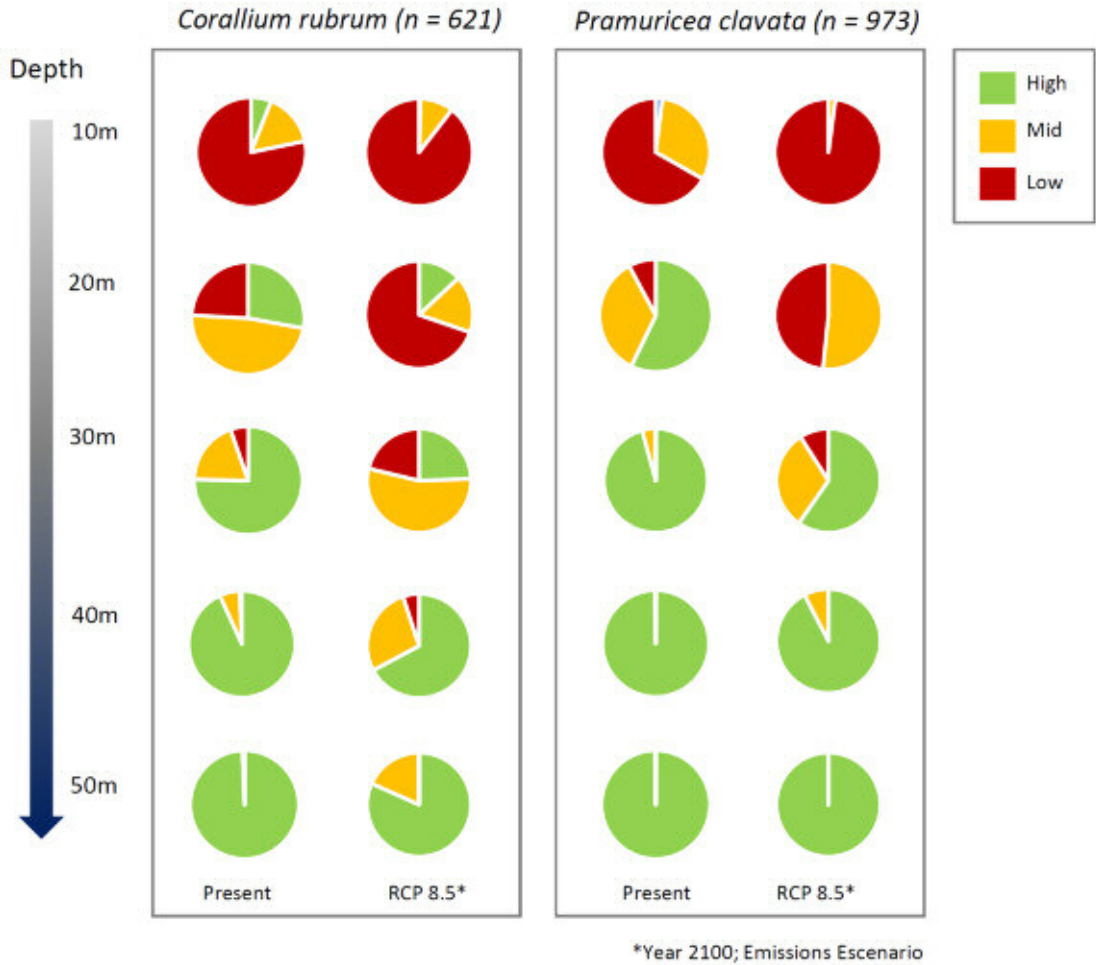


Figure S4. Species-specific predicted habitat suitability for present and future (2100) conditions according to RCP8.5. Suitability values were classified in three levels: high (>0.67), medium (0.34 to 0.67) and low (<0.34).

Research



Cite this article: Montero-Serra I, Linares C, Doak DF, Ledoux JB, Garrabou J. 2018

Strong linkages between depth, longevity and demographic stability across marine sessile species. *Proc. R. Soc. B* **285**: 20172688.

<http://dx.doi.org/10.1098/rspb.2017.2688>

Received: 1 December 2017

Accepted: 1 February 2018

Subject Category:

Ecology

Subject Areas:

ecology, evolution

Keywords:

comparative demography, *Corallium rubrum*, deep sea, integral projection models, life history, longevity

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Electronic supplementary material is available online at <https://dx.doi.org/10.6084/m9.figshare.c.4002595>.

Strong linkages between depth, longevity and demographic stability across marine sessile species

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Understanding the role of the environment in shaping the evolution of life histories remains a major challenge in ecology and evolution. We synthesize longevity patterns of marine sessile species and find strong positive relationships between depth and maximum lifespan across multiple sessile marine taxa, including corals, bivalves, sponges and macroalgae. Using long-term demographic data on marine sessile and terrestrial plant species, we show that extreme longevity leads to strongly dampened population dynamics. We also used detailed analyses of Mediterranean red coral, with a maximum lifespan of 532 years, to explore the life-history patterns of long-lived taxa and the vulnerability to external mortality sources that these characteristics can create. Depth-related environmental gradients—including light, food availability, temperature and disturbance intensity—drive highly predictable distributions of life histories that, in turn, have predictable ecological consequences for the dynamics of natural populations.

1. Introduction

From sequoias to small desert shrubs to naked mole rats, the study of species with unusually long lifespans has long fascinated evolutionary ecologists [1–4]. Understanding the conditions where extreme life histories dominate and the potential consequences of these singular strategies for the dynamics of natural populations is central to ecology and evolution, and may also have important implications for conservation biology [5]. However, to date ecologists have only found weak correlations between extrinsic factors and longevity, and we still have relatively few full demographic descriptions for extremely long-lived species.

In terrestrial landscapes, high longevities are common in relatively inhospitable environments characterized by low availability of resources, such as desert and alpine ecosystems [6–8]. However, some species of extreme longevity inhabit highly productive habitats (e.g. the coastal redwood), and climate or habitat factors generally provide only weak predictions of life-history traits [9–12]. The marine realm is perhaps a more promising setting to examine the external factors driving life-history variation than are terrestrial habitats, owing to the set of physical–chemical properties that lead to strong and predictable depth-related gradients at smaller spatial scales than are observed in other systems [13]. The main abiotic factors controlling bottom-up processes in marine communities (light, food availability, temperature and disturbance intensity) all generally decrease with depth (e.g. [14,15]); these factors are also often invoked as likely selective agents favouring short or long lifespans [16]. If environment plays a role in shaping the evolution of longevity, we, therefore, expect an increase of lifespan along depth gradients. Yet, quantitative analyses addressing this hypothesis are scarce.

In this study, we first explore longevity patterns of marine habitat-forming species, asking how predictable longevity is across depths and whether there are consistent consequences of individual longevity for population dynamics. We discuss these results in the light of similar analyses for terrestrial plant species. Next, we conduct a focused analysis of the iconic Mediterranean red coral, examining aspects of population biology of an extremely long-lived species that may be general for high-longevity species. Our findings represent an important step towards a better understanding on the role of environmental conditions as drivers of the evolution and maintenance of longevity and have implications for the conservation of dominant structural species found at deep habitats, where demographic data are scarce.

2. Material and methods

(a) Review of demographic studies

Life-history strategies are complex combinations of multiple traits that describe the timing and magnitude of the species' reproduction, growth and survival, which in turn generate life-history metrics such as longevity, generation time and net reproductive rate. For two reasons, we have focused here on longevity as a proxy for this broader diversity in life-history strategies. First, several comparative demography studies have shown that longevity is highly correlated to many other demographic traits; in particular, longevity is positively correlated to age at maturity and generation time and negatively correlated to reproductive output and growth [17,18]. Second, unlike many complicated life-history traits that require detailed demographic study to estimate, longevity can be estimated in quite simple ways and is widely reported, making it especially suited to broad-scale comparisons.

To construct our dataset, we searched the literature for studies reporting longevity estimates and demographic data by using several combinations of keywords. We restricted our analysis to marine sessile species and used the terms 'marine sessile', 'coral', 'hexacoral', 'octocoral', 'macroalgae', 'seaweed', 'bivalve' and 'bryozoan'. These were combined with either 'demographic model' and 'matrix model' or 'longevity' and 'lifespan' in the Web of Knowledge. We found 144 suitable studies reporting data for 241 species (electronic supplementary material, table S9). We include studies of polychaetes that form cemented tubes, and thus are effectively sessile, as well as studies of bivalves that are and are not strictly sessile.

Maximum longevity is estimated through different methodologies. Some studies directly report estimates of maximum observed lifespan using growth rings or geochemical analysis or both. Others report standard demographic data in the form of a matrix model or annual survival rates of adult individuals or colonies. The latter were used to derive potential maximum lifespan based on demographic simulations using the same methodology as for the red coral (see §2d(iii), Demographic analysis below). When demographic data were based on non-annual transitions, annual lifespan was calculated as $L_{\text{ann}} = L \times 12/T$, where L_{ann} is the annual lifespan, L is the non-corrected lifespan value and T is the period described by each transition in months.

As a proxy for habitat, we used depth of occurrence, quantified by the maximum depth reported for the species. While mean or median depth would be more informative, accurate information on these central depths is not available for a wide range of species. Maximum depth is more commonly reported and likely to be correlated with a species' characteristic depth range. Maximum depths were obtained by searching in the literature including the keywords 'depth', 'depth range' and 'maximum depth' combined to the species scientific name (electronic supplementary material, table S9).

(b) Demographic variability using matrix population models

To quantify temporal variance in population growth, we used stochastic demographic models based on at least 5-year study periods (four annual transition matrices) for marine sessile species ($n = 9$) and terrestrial plants ($n = 25$) (electronic supplementary material, table S10). For each species, we simulated the fate of 1000 initial individuals for 1000 years, assuming equal probabilities for each reported annual transition, and computed changes in population size by dividing the total number of individuals at each time (t) by the previous number ($t - 1$). We then log-transformed these realized annual lambda values and computed their variance. Two long-term demographic studies on bivalve species reported only observed annual lambdas, thus we computed variance in population growth directly from these observed lambdas after a log-transformation.

(c) Statistical analysis

We used Pearson's and Spearman's rank correlations and linear models to assess the relationships between depth and longevity as well as longevity and population growth variability. We also explored multiple combinations of potential predictors and their interactions by fitting a set of multiple linear models and using Akaike information criteria corrected for small sample sizes to select the best models. First, a set of linear models was fitted to the whole longevity dataset to test for effects of depth while accounting for different methodological differences in longevity calculations and also to explore potential significant interactions between depth and taxonomic class. We also ran a separate set of models to test if other taxonomic levels (Phylum or Order) had more explanatory power or shifted the results. A second set of models was fitted using only the species for which estimates were based on demographic models. For these species, we explored which longevity definition (see §2d(iii), Demographic analysis below) had a better fit with maximum depth. Finally, to assess the effect of longevity on demographic variability, a third set of models was fitted to sessile species for which four or more annual transition models were available, setting variance in population growth as a response variable and maximum longevity, habitat and their interaction as potential predictors. In these and all other statistical tests, longevity was log-transformed for analysis to achieve more linear results.

(d) Study case: the Mediterranean red coral *Corallium rubrum*

(i) Natural history

Precious corals are long-lived species found in several seas across the world. The red coral *Corallium rubrum* is a long-lived octocoral of the Mediterranean Sea. It is a depth-generalist species commonly observed in the northwestern Mediterranean at shallower depths in overhangs and caves but its bathymetrical distribution can reach up to 1000 m deep [19]. Owing to the high value of its carbonate skeleton for the ornamental jewellery industry, *C. rubrum* has been intensively harvested for millennia. Exploitation—both legal and illegal—as well as ongoing climate change are current threats to this species [20–25].

(ii) Study area and demographic monitoring

A total of 1144 coral colonies from eight *C. rubrum* populations along the northwestern Mediterranean region were individually monitored using photographic techniques ($n = 30$ photoquadrats of 20×20 cm at each population) over periods ranging from 3 to 10 years between 2003 and 2011 (electronic supplementary material, table S1; see [26] for detailed information).

(iii) Demographic analysis

Based on the photographic time-series data, we could estimate size-dependent annual survival, fecundity and growth rates. We used a set of generalized mixed effects models (GLMMs) to describe annual survival and fecundity rates. Because the goal was to describe the general life history of the red coral, population and year were included in the models as random factors. Growth rates are extremely low for this species [27,28], thus a nonlinear growth/shrinkage model was fitted to data of initial and final colony sizes measured for 247 colonies in two populations over periods of 7–8 years. Owing to relatively rare ‘extreme shrinkage’ of some colonies, we divided the growth/shrinkage process into normal growth and extreme retrogression, each with their own mean change of size (see the electronic supplementary material). The growth/shrinkage rates model and the fixed effects from the survival and reproduction GLMMs, which included colony size as a fixed effect and population and year as random factors, were then used to construct a deterministic integral projection model (hereafter IPM). We analysed this model as a large matrix based on 80 colony size-classes (see [26]; electronic supplementary material for a full description on model construction).

Maximum potential lifespan was estimated by deterministic simulations using the demographic IPMs. Specifically, we simulated the fate of 100 individuals starting at the same stage and recording the number of years until 50% (mean lifespan) and 5% (maximum potential lifespan) of individuals were still alive. To explore the expected mean and maximum lifespan conditional on starting at different life-stages, we ran three set of simulations: starting with 100 recruits, starting with 100 of the smallest size-class adults (first reproductive individuals [29]), and starting with 100 largest size-class adults. In all results reported in the main text, we use adult maximum longevity. To explore how robust these longevity metrics are to changes in the dimension of the transition matrix derived from the IPMs, we computed all these metrics using matrices constructed with 10 to 200 size-classes.

To assess the elasticity of deterministic population growth to each vital rate, we performed a perturbation analysis of size-dependent survival, fecundity, and mean normal growth and mean extreme shrinkage, using an 80 size-class model. Each size-dependent vital rate was separately increased or decreased (by 1% and 5%). A mean demographic model was derived that included each perturbation and its deterministic population growth rate (dominant eigenvalue) was computed. Finally, the size-dependent elasticity of each vital rate to λ was calculated using:

$$e_{a_j} = \frac{a_j}{\lambda} \cdot \frac{\partial \lambda}{\partial a_j},$$

where a is the vital rate, j the size-specific interval, λ the deterministic population growth and e_{a_j} is the elasticity for vital rate a_j . Since underlying vital rates were perturbed separately (e.g. mean growth, mean extreme shrinkage and survival), no compensation to control for shifting effects on other vital rates was required (see approach 4 in [30]).

We calculated mean age-specific survival and fecundity rates by simulating a cohort of 1000 new recruits and computing at each time step the survival rate (N_t/N_{t-1}) and the fecundity (R_t/N_{t-1}), where N is the total adult abundance and R is the total number of recruits.

3. Results

(a) Depth, longevity and population stability

The comparison of maximum lifespan revealed a great diversity of life histories across marine sessile species (figure 1). Hydrozoans, bryozoans, cirripeds and polychaetes displayed the shortest lifespans, ranging from weeks to several decades. These generally short-lived taxa have received significantly

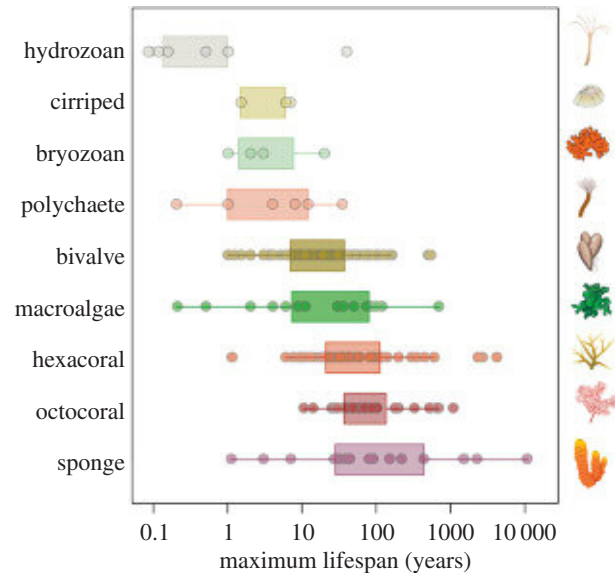


Figure 1. Longevity patterns across marine sessile species ($n = 241$). Data were obtained from the literature and correspond to the maximum potential lifespan reported for each species for the main marine sessile taxa. See the electronic supplementary material, table S9 for specific species data. (Online version in colour.) Images: Integration and Application Network, University of Maryland Center for Environmental Science (ian.umces.edu/symbols/).

less attention in the literature compared with other major taxa such as bivalves and corals. The other groups had a great range of lifespans, up to hundreds of years for some calcareous macroalgae and bivalves and to thousands of years for cnidarians and sponges. Most macroalgae species ranged from less than 1 year to several decades, but an outlier with exceptional longevity is the Antarctic calcareous algae *Clathromorphum nereostratum*. Marine bivalve’s lifespans ranged from less than 1 year to more than 500 years for the Antarctic scallop *Arctica islandica* and the giant deep-sea oyster *Neopycnodonte zibrowii*, which is the longest-lived known non-clonal animal. Modular and structural taxa such octocorals, hexacorals and sponges were dominated by longer-lived species. Extreme cases were the deep-sea coral *Gerardia* sp., found in Hawaii, with an estimated age of 4700 years and the longest-lived known marine species, the sponge *Monorhaphis chuni*, observed at 1000 m depth with an estimated age of 11000 years [31,32].

We correlated $\log(\text{maximum lifespan})$ with $\log(\text{maximum depth})$ and found strong positive relationships in the five main taxa for which enough data were available to perform meaningful analyses: octocorals (Spearman’s $\rho = 0.673$; $p < 0.001$; $n = 35$), hexacorals (Spearman’s $\rho = 0.592$; $p < 0.001$; $n = 40$), sponges (Spearman’s $\rho = 0.647$; $p < 0.001$; $n = 17$), macroalgae (Spearman’s $\rho = 0.600$; $p = 0.023$; $n = 14$) and bivalves (Spearman’s $\rho = 0.491$; $p < 0.001$; $n = 93$) (figure 2; electronic supplementary material, table S2). The positive relationship between depth and lifespan was consistent across different measures of longevity, although adult maximum longevity showed the strongest relationships with depth (electronic supplementary material, table S3). This relationship is also consistent when accounting for taxon or matrix dimension; the best supported model includes an interaction between depth and taxon, with significantly larger slopes for sponges ($p = 0.0131$) and macroalgae ($p = 0.0308$) than for the other taxa (electronic supplementary material, table S4; overall model statistics: $R^2 = 0.511$; $F_{185} = 18.12$; $p < 0.001$). Use of other taxonomic levels does not change this qualitative result,

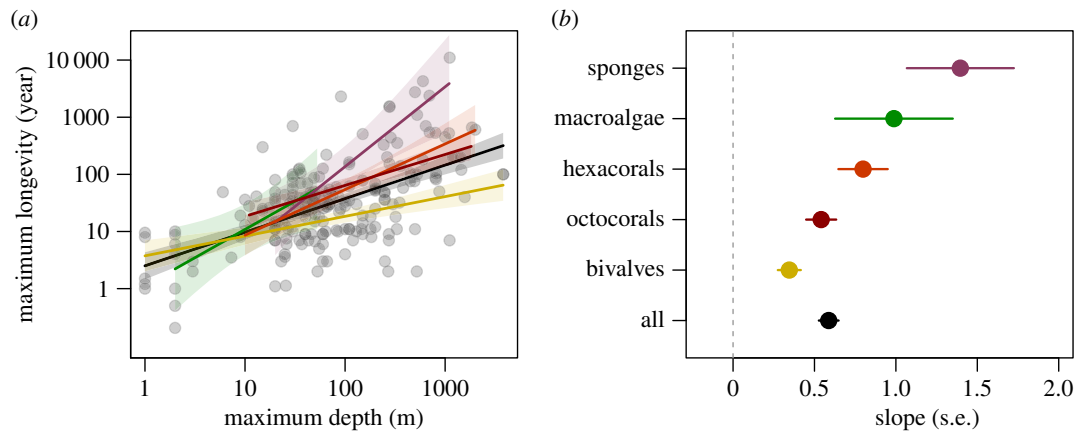


Figure 2. (a) Relationship between maximum depth of occurrence and maximum longevity across marine sessile species. Colours correspond to the labels in (b), and lines show the best fit linear model for each taxon. Black line shows the linear relationship for all taxa. Shaded areas represent standard errors. (b) Slope corresponds to the estimated slope of maximum depth on maximum longevity for each taxon, with error bars showing ± 1 s.e. (Online version in colour.)

and model fit is best for Class (electronic supplementary material, table S5). We also ran quantile regressions between $\log(\text{depth})$ and $\log(\text{longevity})$ to see if depth is a more powerful regulator of extreme life-history values, but found no significant differences in the slope of the depth–longevity relationship for 10% up to 90% quantiles (electronic supplementary material, table S6).

To assess population consequences of longevity, we regressed variance in realized log-lambda on longevity across marine sessile invertebrates and terrestrial plants and found a similar, strongly negative relationship for both. Short-lived species displayed higher variability in population growth over time than did longer-lived species ($n = 34$; Spearman's $\rho = -0.73$; $p < 0.001$). The pattern was consistent in terrestrial plants ($n = 25$, Spearman's $\rho = -0.68$; $p = 0.0002$) and marine sessile invertebrates ($n = 9$; Spearman's $\rho = -0.67$, $p = 0.0214$) (figure 3; electronic supplementary material, table S7).

(b) The extreme life-history of *Corallium rubrum* and its population dynamics

The red coral *C. rubrum* displayed slow dynamics over the eight populations studied along the northwestern Mediterranean (electronic supplementary material, figure S1), a pattern driven by extremely high adult survival, ranging between 98.05% and 99.76% from the smallest to the largest reproductive *C. rubrum* colonies (figure 4a). Monitoring of post-recruitment survival revealed that relative to later survival rates, annual survival is low during the first years after recruitment with a mean of 69.5% (figure 4c). However, by the age of four, young colonies already displayed a high survival probability of 92.2%. Overall, mortality rates decreased rapidly during the early stages of the red coral lifetime, following a type III survivorship curve, and then decreased steadily, more similar to type II survivorship (figure 5a). Recruitment was very limited; from 246 quadrats (about 30 per population) individually monitored for recruitment at the eight studied populations, 83.7% showed no annual recruitment. The remaining 16.3% of quadrats were divided into 10.7% with only a single recruit present, 5.1% with two to four recruits and only 0.5% with five or more recruits (figure 4b,d). Mean annual *per capita* recruitment, weighted by the relative abundance of different sized individuals, was 0.077. This is considerably lower than the average *per capita* recruitment for a range of marine sessile

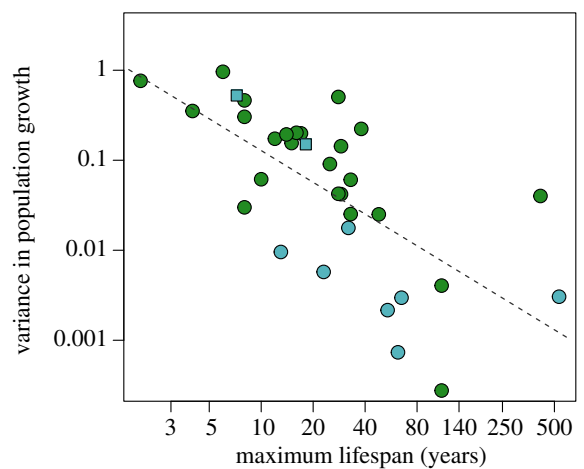


Figure 3. Relationship between temporal variability in population growth and maximum longevity in terrestrial plants (green, $n = 25$, Spearman's $\rho = -0.68$; $p = 0.0002$) and marine sessile (light blue, $n = 9$; Spearman's $\rho = -0.67$, $p = 0.0214$) species, and overall ($n = 34$; Spearman's $\rho = -0.73$; $p < 0.0001$). Squared light blue dots correspond to two bivalve species whose variance was computed directly from observed annual lambdas. (Online version in colour.)

invertebrates for which there is standard demographic data available ($n = 28$, median = 0.547–0.727, mean = 5.14 ± 4.02 (s.d.); electronic supplementary material, table S8), although higher than the same estimate for six other species (electronic supplementary material, table S8).

Demographic simulations revealed high maximum potential future longevity for *C. rubrum* colonies, ranging from 501 to 532 years for small and large adult, respectively (figure 5a; electronic supplementary material, figure S2). The mean expected lifespan was also relatively high for adult red coral colonies and ranged from 98 years for small adult colonies to 129 years for the larger red coral colonies. Our simulations also revealed potential artefacts owing to matrix dimensionality. When using relatively small transition matrices (less than 60 size-classes) simulations underestimated values of maximum and mean longevity, the values stabilized around 60–80 size-classes and were consistent thereafter (electronic supplementary material, figure S2).

Perturbation analysis revealed a disproportionately large influence of survival rates on asymptotic population growth

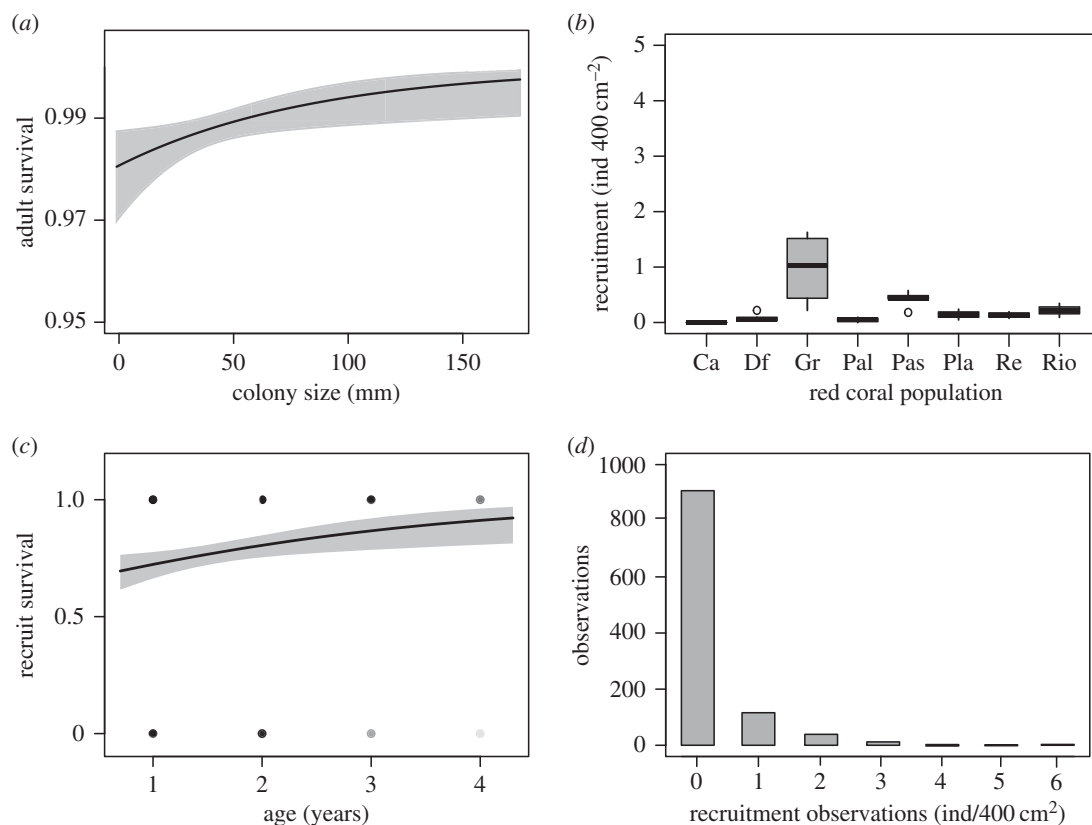


Figure 4. Long-term demographic traits of the red coral *Corallium rubrum*. (a) Adult survival probability depending on colony size. The line represents a logistic generalized linear mixed effects model including population and year as random effects and colony size as a fixed effect. (b) Mean recruitment rates in eight *C. rubrum* populations. (c) Age-dependent post-recruitment survival probability. (d) Frequency of annual recruitment observations in eight *C. rubrum* populations. In (a,c) grey areas show ± 1 s.e.

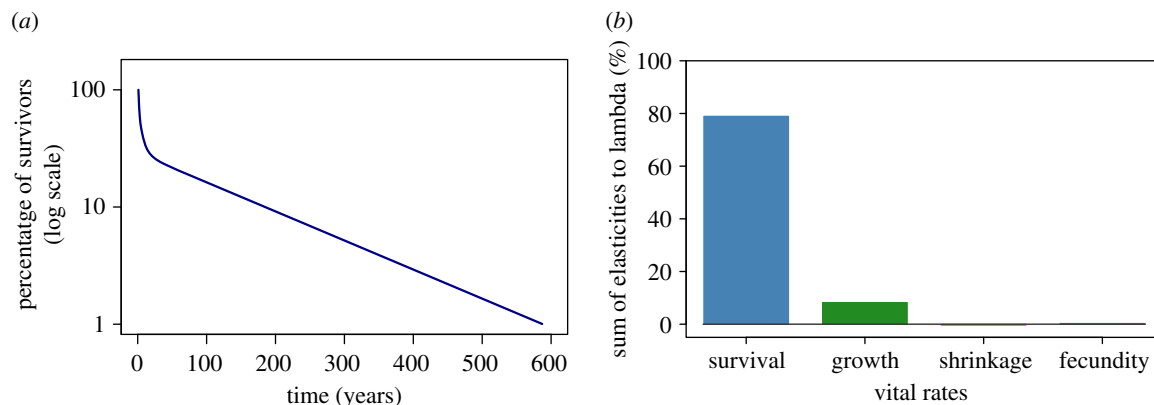


Figure 5. Demographic analyses for the red coral *Corallium rubrum*. (a) Survivorship curve based on a deterministic simulation of 100 individuals. (b) Patterns of elasticity to the asymptotic population growth rates of different vital rates, with each bar indicating elasticity summed across all size-classes. Other vital rates have even lower elasticities (electronic supplementary material, figure S4). (Online version in colour.)

(figure 5b). Elasticity of lambda to fecundity, mean growth and mean extreme shrinkage, as well as all other vital rates, was markedly lower (electronic supplementary material, figure S4). Model fitting of age- or size-dependent fecundity and survival showed no evidence of senescence (electronic supplementary material, figure S3).

4. Discussion

Demography is a powerful tool to understand the evolution of life histories and has provided important insights into plant, mammal and bird's strategies [5,33]. However, a lack

of long-term and broad spatial-scale data on the dominant species of marine ecosystems, particularly those living at deeper depths, hinders our understanding of the processes shaping the dynamics of these threatened ecosystems [34]. In this study, we analysed the long-term dynamics of the long-lived red coral *C. rubrum* and published data for a broad range of marine sessile invertebrates and macroalgae. The results revealed strong positive relationships between depth, longevity and demographic stability, showing strong habitat-determined predictability of key demographic processes in marine ecosystems.

Marine sessile species have a wide range of maximum life-spans across taxa and habitats, ranging from weeks in some

intertidal species to hundreds of years for the red coral *C. rubrum*, and several thousands of years for some deep-sea corals and sponges (figure 1). Our most striking finding was that maximum lifespan was strongly and positively correlated with a species' maximum depth occurrence in hexacorals, octocorals, sponges, and although weaker, also in bivalves (figure 2). These results demonstrate a strong role of habitat features at shaping the distribution of longevity patterns.

In the shallowest waters, short-lived species dominate (figure 2), perhaps owing to high disturbance rates and competition in high-energy environments where fast-growing species have a distinct advantage. By contrast, species displaying diverse life-history strategies coexist at intermediate depths, including short-lived invertebrates that also occur intertidally and longer-lived massive corals, macroalgae and seagrasses [35,36]. The high energy availability owing to constant solar irradiation may allow the presence of a large spectrum of life-history strategies that have diversified to occupy multiple niches, following analogous processes than those shaping tropical rainforest plant communities [37]. Finally, in deeper waters, environmental constraints seem to favour the dominance of slower life histories and longer lifespans. Depth gradient determines changes in environmental factors such as light and food availability, temperature and disturbance intensities. In turn, these factors are likely to control the evolution of lifespans through several mechanisms. First, deeper habitats are generally less productive, limiting food supply for consumers and favouring slow growth and longer lifespans [38]. Empirical studies exploring physiological trade-offs have shown that the elevated metabolic costs of fast strategies can be detrimental in the absence of high availability of resources [39]. Also, as light is strongly depth-dependent, organisms dwelling at deeper habitats may experience a 'competition release', mainly by the absence of macroalgae and other fast-growing autotroph organisms that are constrained to the photic zone [40]. Similarly, deeper habitats are less diverse and thus organisms could face less predation pressure, pushing selective forces to favour somatic maintenance and, therefore, life extension [41]. Finally, the relative predictability of deep environments may enable species to survive with lower reproductive and mortality rates. Shallow habitats are more exposed to external sources of mortality owing to extreme physical disturbances (i.e. hurricanes and storms) and temperature fluctuations [13,42], making them less suitable for organisms whose long-term success rely on extremely high survival of adult stocks [43].

Previous anecdotal data on longevity in sessile and mobile marine organisms support our findings. For instance, the highest longevity reported to date corresponds to the sponge *M. chuni*, found at 1000 m depth with an estimated age of 11000 years [32]. A similar pattern has been also observed in marine fishes [44]. Indeed, a recent study based on radiocarbon dating on eye tissue revealed that the deep-resident Greenland shark *Somniosus microcephalus*, which can live up to 400 years, may be the longest-lived vertebrate on the Earth [45]. Overall, these results demonstrate a strong role of depth-related environmental gradients in shaping life-history strategies across marine taxa. The strength of this pattern is striking and contrasts with the weak environmental correlates of longevity or other life-history traits that seem typical in terrestrial systems [9–12]. Nonetheless, our findings open new questions regarding the existence of potentially analogous processes driving the evolution of life-history strategies along elevation and/or productivity gradients in terrestrial landscapes.

Further analyses exploring the relationship between depth and different environmental parameters could provide a more mechanistic understanding of this general pattern of increasing species maximum longevity towards deeper marine waters.

At least three demographic mechanisms may contribute to the strong relationship observed between depth and longevity in marine sessile organisms. First, species may show relatively indiscriminate settlement. Under this scenario, the longevity pattern may arise because faster-growing, shorter-lived species outcompete slow species in productive, shallow sites, while at deeper sites only slow-growing species with high longevity may persist. Second, species may have evolved specific settlement cues so that larvae of long-lived species preferentially settle in either deep or shallow sites. Finally, to some extent these life-history patterns could result from demographic compensation, reflecting phenotypic plasticity in life histories (i.e. fecundity, growth or survival rates) as a function of local constraints on population growth rates [46]. In reality, all three mechanisms are likely to contribute to this matching of life history to environment.

(a) The extreme life-history features of the red coral *Corallium rubrum*

While our survey shows that mesophotic and deep-sea ecosystems are predictably inhabited by species of extreme longevity, we have few detailed studies of the population ecology of these taxa. We found that *C. rubrum*, a widely distributed Mediterranean red coral, showed a consistent pattern of slow population dynamics driven by extremely high survival, recruitment limitation and apparently negligible senescence (figure 5; electronic supplementary material, figure S3a). The species revealed a maximum potential longevity of up to 532 years, overlapping with lifespan ranges of other deep-sea organisms. Although previous research had shown that *C. rubrum* settlement can be highly heterogeneous [47,48], in this study, we observed a much more homogeneous pattern of low recruitment in the long term (figure 4b,d; electronic supplementary material, table S8). Recruitment limitation has been observed in other long-lived temperate invertebrates [49–51] and can seriously hinder the ability of populations to recover after intense perturbations [52].

For species with limited recruitment, adult survival becomes of paramount importance to ensure their persistence [43]. We found extremely high survival for *C. rubrum* (figures 4a and 5a), with estimated mortality even lower than found in previous experimental studies [27,50]. In agreement with previous observations in long-lived gorgonians [43], the perturbation analysis supported that survival is the vital rate demonstrating the highest effect on population growth in *C. rubrum* (figure 5b). The absence of detectable senescence is common in sessile marine species and terrestrial plants [2,53], and may be related to high investments in structural tissues, although some authors have suggested that modularity may also play an important role [53]. It should also be noted that our data, like that for most other studies of extremely long-lived species, lack the precision necessary to carefully test for senescence. Precious corals build an energetically demanding hard carbonate skeleton; while preventing a fast rate of colony growth, it may provide the structural basis that allows for an extremely high survival.

The shallow occurrences of the monitored populations of this long-lived species could seem to make it an outlier in the depth-longevity pattern. However, this coral is distributed from shallow to deep waters up to 1000 m depth [19] and its shallow occurrences are exclusively found in dim-light and dark habitats where it cannot be outcompeted by fast-growing organisms [40]. Given the overall pattern observed for marine species, the dynamics of red coral populations dwelling at deeper habitats are likely to be even slower and more dependent on adult survival. However, we must be cautious with these extrapolations and note that demographic data on deeper locations is not available owing to logistical constraints. Exceptional shallow presence has been also observed in other important cold-water corals, such as *Lophelia pertusa*, which can occur at 25 m in sediment laden fjords where light attenuation reduces the abundances of autotroph organisms, and thus competition; while at the other end of the scale this species may be found as deep as 3300 m [54].

(b) Population-level consequences of extreme life histories

As important as unravelling the drivers of extreme longevity are the implications of these strategies for dynamics of natural populations. We show that maximum longevity strongly predicts temporal stability in population growth across terrestrial and marine sessile species (figure 3). This finding agrees with previous studies suggesting that long-lived species can strongly buffer environmental stochasticity [55,56]. As observed for the red coral *C. rubrum*, this buffering capacity relies on high adult survival and low reproductive success. While conferring stability under natural conditions, this extreme life history will also hinder the ability of long-lived species to overcome increased mortality rates, but whether these species will be able to cope with rapidly changing perturbation regimes driven by the ongoing global change is still unclear. For instance, while the red coral has

persisted after millennia of historical overharvesting owing to the combination of consistent survival of partially harvested colonies [25] and small size at maturity [29], new human-related stressors such as global warming have increased mortality rates and put some shallow populations at risk [20,21,57]. Worryingly, although declines may already be driving populations of some long-lived species towards collapse, in some cases, these trajectories may be too subtle to be noticed by ecologists and managers [58]. The consistent depth relationship for longevity also implies greater sensitivity of species and communities occurring at greater depths to human perturbations, amplifying recent calls to better monitor and protect these vulnerable ecosystems [34].

Data accessibility. All data are deposited in Dryad: <http://dx.doi.org/10.5061/dryad.p0b6b> [59].

Authors' contributions. I.M.-S., C.L., D.F.D. and J.G. designed the study. C.L., J.B.L. and J.G. collected the demographic data in the field. I.M.-S. extracted data from the literature. I.M.-S. and D.F.D. performed analyses. I.M.-S. wrote the first draft of the manuscript, and all authors contributed substantially to revisions.

Competing interests. We declare we have no competing interests.

Funding. This study was partially funded by the Spanish Ministry of Economy and Innovation Biorock project (CTM2009-08045), the Smart project (CGL2012-32194), the TOTAL Foundation Perfect project and the European Union's Horizon 2020 research and innovation programme under grant agreement no 689518 (MERCES). This output reflects only the authors' view and the European Union cannot be held responsible for any use that may be made of the information contained therein. I.M.-S. was supported by a FPI grant (BES-2013-066150), C.L. by a Ramon y Cajal (RyC-2011-08134) and J.B.L. by a postdoctoral grant (SFRH/BPD/74400/2010) from the Portuguese Foundation for Science and Technology. Support for D.F.D. came from National Science Foundation awards 1340024 and 1242355. I.M.-S., C.L., J.B.L. and J.G. are part of the Marine Conservation research group (2014 SGR 1297) from the Generalitat de Catalunya.

Acknowledgements. We thank K. Kaplan, M. Pages, A. Griffith and one anonymous reviewer for their valuable comments on early versions of this manuscript.

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RESEARCH ARTICLE

Harvesting Effects, Recovery Mechanisms, and Management Strategies for a Long-Lived and Structural Precious Coral

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Citation: Montero-Serra I, Linares C, García M, Pancaldi F, Frleta-Valić M, Ledoux J-B, et al. (2015) Harvesting Effects, Recovery Mechanisms, and Management Strategies for a Long-Lived and Structural Precious Coral. PLoS ONE 10(2): e0117250. doi:10.1371/journal.pone.0117250

Academic Editor: Chaolun Allen Chen, Biodiversity Research Center, Academia Sinica, TAIWAN

Received: August 12, 2014

Accepted: December 19, 2014

Published: February 23, 2015

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Data Availability Statement: All relevant data are within the paper and its Supporting Information files.

Funding: Support for this work was provided by a FPI grant (BES-2013-066150) to I. Montero-Serra and by a Ramon y Cajal research contract (RyC-2011-08134) to C. Linares. This study was partially funded by the Spanish Ministry of Economy and Innovation Biorock project (CTM2009-08045) and Smart project (CGL2012-32194). The authors are part of the Marine Conservation research group (2009 SGR 1174) from the Generalitat de Catalunya. The funders had no role in study design, data

Abstract

Overexploitation is a major threat for the integrity of marine ecosystems. Understanding the ecological consequences of different extractive practices and the mechanisms underlying the recovery of populations is essential to ensure sustainable management plans. Precious corals are long-lived structural invertebrates, historically overfished, and their conservation is currently a worldwide concern. However, the processes underlying their recovery are poorly known. Here, we examined harvesting effects and recovery mechanisms of red coral *Corallium rubrum* by analyzing long-term photographic series taken on two populations that were harvested. We compared the relative importance of reproduction and re-growth as drivers of resilience. Harvesting heavily impacted coral populations causing large decreases in biomass and strong size-class distribution shifts towards populations dominated by small colonies. At the end of the study (after 4 and 7 years) only partial recovery was observed. The observed general pattern of low recruitment and high mortality of new recruits demonstrated limited effects of reproduction on population recovery. Adversely, low mortality of partially harvested adults and a large proportion of colonies showing new branches highlighted the importance of re-growth in the recovery process. The demographic projections obtained through stochastic models confirmed that the recovery rates of *C. rubrum* can be strongly modulated depending on harvesting procedures. Thus, leaving the basal section of the colonies when harvesting to avoid total mortality largely enhances the resilience of *C. rubrum* populations and quickens their recovery. On the other hand, the high survival of harvested colonies and the significant biomass reduction indicated that abundance may not be an adequate metric to assess the conservation status of clonal organisms because it can underestimate harvesting effects. This study highlights the unsustainability of current harvesting practices of *C. rubrum* and provides urgently needed data to improve management practices that are still largely based on untested assumptions.

collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

Introduction

Overfishing is a major threat to the integrity of world's marine ecosystems. Historical exploitation of marine resources has resulted in drastic population declines, species extinctions and the general simplification of marine food webs [1], [2]. This structural impoverishment has also hindered the resilience of marine populations, lowering their ability to recover after recurrent natural and human disturbances [3], [4]. Understanding the ecological consequences of different extractive practices on the structure and function of populations, especially of those organisms exhibiting a long life span and slow dynamics, is therefore one of the greatest challenges in conservation biology [5].

Precious corals have been harvested and traded worldwide since ancient times due to the high economic value of their carbonate axial skeleton [6], [7], [8]. These sessile invertebrates are considered habitat forming species with a keystone role on coastal systems because they provide structural complexity and host high levels of biodiversity [9]. Additionally, they can enhance fisheries by providing shelter during early stages for some commercially important species of rock fishes, shrimps and crabs [10]. The red coral *Corallium rubrum* is a precious octocoral endemic to the Mediterranean rocky bottoms and adjacent Atlantic waters. Available data showed recent large declines in the Mediterranean yields, suggesting that this fishery is unsustainably managed following boom and bust cycles [11], [7]. Intensive harvesting has resulted in significant shifts in the size structure of current *C. rubrum* populations [12], [7], causing a decrease in biomass and mean and maximum colony size [13], [14]. Moreover, current warming and acidification trends associated to global change are major threats for shallow red coral populations [15], [16], [17], [18]. Consequently, there is a growing concern for the conservation of *C. rubrum* and the rest of the precious corals worldwide as evidenced by the recent struggle to include the Genus *Corallium* in the Appendix II of Cites [7], [19], [20], [21].

The need for a responsible fishery for *C. rubrum* and the negative effects of current practices are widely accepted [11], [21], [22]. However, given the fragility and vulnerability to different threats of this species, all studies addressing harvesting effects and recovery have been carried out by comparing size-class distribution of different populations and periods. [11], [23], [24], [25]. To date, biomass reduction has been only quantified once after a poaching event [26]. Beyond the reported shifts in size structures of *C. rubrum* populations [11], [24], sound field and experimental data on the effects of different extractive practices and the recovery process are essentially lacking in the literature.

Due to their low population dynamics and limited dispersal capacities [27], [28], precious corals are considered to have low resilience to disturbances that cause high adult mortality such as harvesting. Paradoxically, *C. rubrum* has not undergone extinction despite being exposed to intensive harvesting since ancient times. In fact, the biological processes underlying coral populations' persistence are largely unknown. The early maturity of *C. rubrum* (2.4 cm in height) corresponding to an age of 6–10 yr [29, 30], could partially explain this persistence. It allows the colonies to contribute to reproductive output before they reach the size of interest for harvesters. On the other hand, colonial organisms have the capacity to recover after events of partial mortality (e.g. breakage of a branch). Indeed, red coral colonies with signs of breakage and recent re-growth of new branches have been observed [29], [31]. The resilience to harvesting will therefore depend on two main mechanisms: (1) re-growth or clonal growth of colonies suffering partial mortality (i. e. when leaving the basal section) and (2) recruitment by sexual or asexual reproduction (i. e. when eradicating the whole colony). Yet, there is no assessment on the relative contribution to recovery of these two mechanisms from a long-term perspective. This assessment would provide a scientific basis for new management measures that enhance the sustainability of this natural resource.

In this study, we examined harvesting effects and the recovery processes of *C. rubrum* using 5–7 years photographic series on two populations located on the coast of Marseilles (France, NW Mediterranean). Fishermen unexpectedly harvested two *C. rubrum* populations that were already being studied by our research team and this represented a unique opportunity to establish before / after comparisons as well as to explore directly the recovery mechanisms by following individually affected as well as unaffected *C. rubrum* colonies. Our overall goal was to elucidate the demographic process underlying the recovery of *C. rubrum* after harvesting events with a particular focus on the relative contribution of re-growth of harvested colonies and reproduction. More precisely we quantified (1) survival and re-growth rates of harvested colonies, (2) survival of non-affected colonies and (3) recruitment rates and survival of recruits. Based on these data, we developed biomass projections to compare population trajectories under different harvesting practices. We contend that the results provide meaningful insights to inform new legislation measures to promote resilience and ensure the conservation of this threatened species and other precious corals also at deeper depths, where most of fishing efforts are concentrated and this type of data is more difficult to obtain.

Methods

1 Study Area

We monitored two Mediterranean red coral populations located along the rocky coast of Massif des Calanques in the SE of Marseilles, France. Due to the specific habitat characteristics of the area, such as submerged cavities and overhangs, along with instability of the water column during summer [32], the development of red coral populations at shallow depths (15–22m) is favored. The studied populations are located in a vertical wall at Riou Island (43° 10' 23.47" N i 5° 23' 13.24" E) (hereafter Riou) and in a cave-like tunnel at Maire Island (43°12'32.34"N; 5°20' 14.01"E) (hereafter Maire).

2 Coral Population Monitoring

In each population we monitored 30 quadrants (20 x 20 cm) by setting up 2 permanent plots using PVC screws fixed to holes in the rocky substratum. Each plot was variable in length, depending on the complexity of the substratum, and 40 cm wide. In each sampling, a cord was deployed between the screws and quadrants assembled with a scale were sequentially positioned and photographed above and below the cord throughout the length of each transect. Two photographs from each quadrat (using 2 slightly different angles, ~30°) were used for analysis with photogrammetric techniques which allowed measuring the height of colonies [33]. Transects were photographed using a NIKON D70 with a housing and 2 electronic strobes.

At Maire, permanent plots were installed at 15 m after a harvesting event was observed during June 2002 and monitored till 2009 but due to logistic constraints, pictures were not taken during 2007 and 2008. Colonies that suffered partial mortality were evident because fishermen remove the branches leaving a wide colony basis, short in height with injuries (naked tissue), that unequivocally belongs to an older red coral colony that was recently pruned. In contrast, at Riou the permanent plots were installed in 2005 just before the harvesting event. Between April and June 2006 fishermen were observed in the area and subsequent surveys clearly detected their impact in the permanent plots (Fig. 1).

Our study did not involve any sampling of or damage to red coral colonies. The authors had all the permits provided by the French authority Affaire Maritimes to perform scientific surveys in the study area.

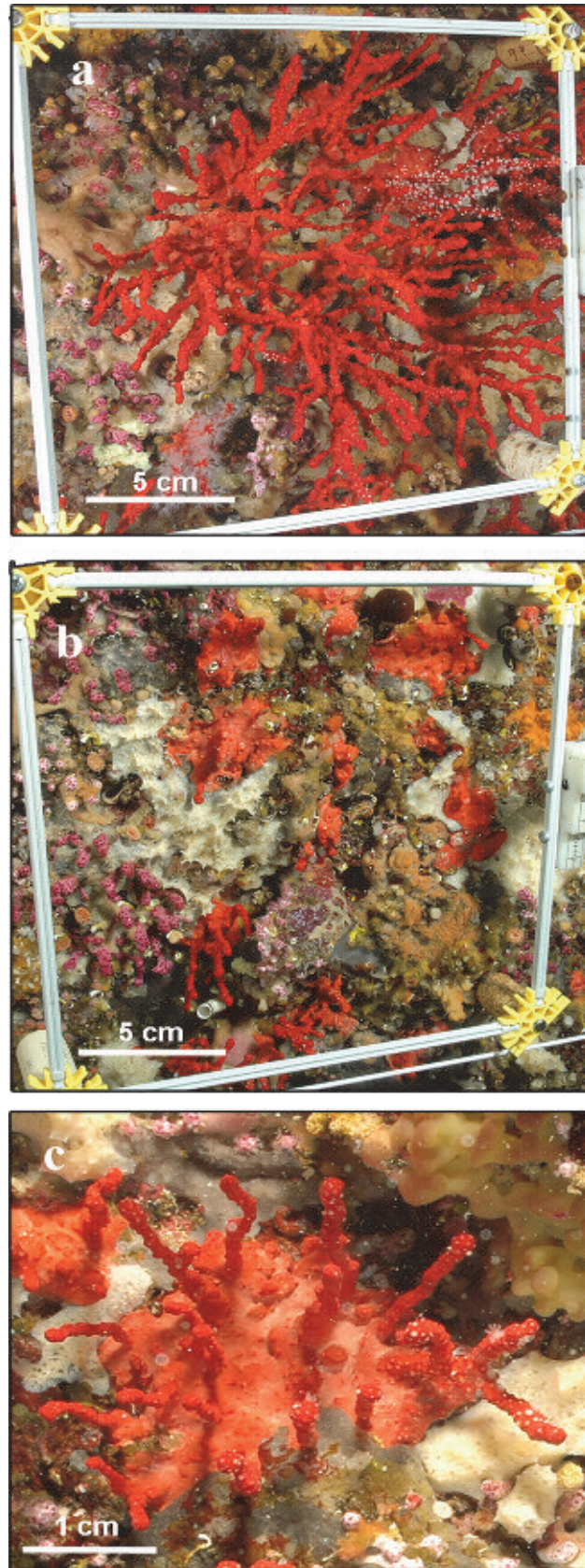


Fig 1. Harvesting effects on red coral populations. Partial mortality at Riou is shown a) before and b) after a harvesting event. c) Detail of re-growth of new branches on a partially harvested colony. Photo credit: Medrecover (www.medrecover.org).

doi:10.1371/journal.pone.0117250.g001

3 Demographic parameters

From the photographic series, the harvested and non-harvested colonies were individually identified and compared year to year to estimate mortality rates, number of branches and colony height. We also estimated adult population density (colonies number per 400 cm²) and recruitment rates. We considered a recruit the new button like colonies observed ranging from a minimum detection size of 2 mm to 10 mm in height. Annual recruitment rates were estimated as the number of recruits appearing for the first time in a photoquadrat and post-recruitment mortality were estimated by following these recruits through the study period.

With this dataset we were able to assess:

3.1 Changes in size-structure

To assess population size-structure changes and the recovery process, we measured maximum height of all colonies and classified them into five size classes (0–30mm; 30–60mm; 60–90mm; 90–120mm, >120mm). Riou population was measured before harvesting (2005), right after (2006) and four years after (2009). Maire population was measured right after harvesting (2002) and seven years after (2009). Biomass variations were estimated by applying a height-weight polynomial equation to height data. The equation was previously calculated from height and weight data measured on 300 dead red coral colonies collected in previous studies in the same region [35], and from different poaching events [26]. The resulting curve was: Weight (g) = 0.001(Height, mm)² + 0.096(Height, mm) – 4.010 (R² = 0.868, P < 0.001) (S1 Fig).

3.2 Recovery mechanisms

We quantified the cumulative survival probability of non-harvested adults, harvested adults, and new recruits based on annual mortality rates, calculated as follows:

$$m = \frac{\left(1 - \frac{N_t}{N_0}\right)}{\Delta_t} \quad (1)$$

where m is the annual mortality rate of *C. rubrum* colonies, N is the total number of colonies, and Δt is the duration of the study period. In addition we assessed the degree of recovery of affected colonies by quantifying the frequency of colonies showing new branches and the number of branches per colony during the study period.

4 Simulations of recovery process under different harvesting practices

To analyze whether recovery rates depend on harvesting practices in the long-term, we developed a set of basic stochastic demographic models under two different scenarios with contrasting levels of total (TM) and partial mortality (PM): a) removal of almost all colonies causing 90% TM and 10% PM; b) leaving the basal section in almost all colonies causing 10% TM and 90% PM. The second scenario was based on the observed mortality values at Riou (see [results](#)). The two scenarios were calculated by setting initial population densities according to the corresponding levels of total mortality. To run the model we used demographic parameters (adult density, annual recruitment rates, adult and post-recruitment mortality, and biomass increase rates per colony) obtained from the two monitored populations during the study period. The two harvesting scenarios were projected by running 100 stochastic simulations during 30 years according to the following steps:

First we estimated the changes in the number of colonies following the next formulation:

$$\begin{aligned} N_{T(t)} &= N_{R(t)} + N_{A(t)} \\ N_{R(t)} &= N_{R(t-1)}(1 - g)(1 - m_g) + r \\ N_{A(t)} &= N_{A(t-1)}(1 - m_A) + N_{R(t-1)}g(1 - m_R) \end{aligned} \tag{2}$$

where N_T is total population, N_R is total number of recruits, N_A is the total number of adults, m_R and m_A are annual mortality rate of recruits and adults respectively, r is the annual recruitment rate and g is the proportion of recruits growing to the adult stage:

$$g = \frac{G}{h_{\max} - h_{\min}} \tag{3}$$

where G is the recruit growth rate ($2.5 \text{ mm} \cdot \text{yr}^{-1}$ in height, from reference [34]), h_{\min} and h_{\max} are the lower and upper height intervals for recruits.

Secondly, we estimated the evolution of biomass recovery using the biomass increases (ΔB) per colony and abundance values from the demographic models previously described. Mean (SD) biomass increases were estimated according to the next equations applied to colonies suffering partial mortality at Riou ($n = 57$) and Maire ($n = 67$) right after harvesting and 4 and 7 years after respectively:

$$\Delta B = \frac{1}{N} \sum_{i=1}^N \left(\frac{B_f - B_0}{\Delta t} \right)_i \tag{4}$$

$$B_{(t)} = NT_{(t)} \cdot \Delta B \tag{5}$$

where B_0 is the colony biomass right after harvesting, B_f is the total biomass at the end of the study period, Δt is the number of years, and NT is the total number of colonies in each population.

Results

1 Harvesting effects

Harvesting affected 30.1% of the colonies at Riou ($n = 209$) and 49.6% of the colonies at Maire ($n = 139$). At Riou, total mortality was around 9.5% and partial mortality was 90.5%, causing a large biomass loss with a 98.5% of reduction in total dry weight (Fig. 2). Harvesting targeted mainly medium and large colonies, triggering a dramatic size-distribution shift toward populations characterized by a large proportion of small colonies (Figs. 1 and 3A). At Maire, because the monitoring started after the harvesting event, we could not determine the loss of biomass nor the size-class distribution shift. Nonetheless, the same size-distribution pattern was observed after the harvesting event with no large colonies present and a great proportion of small-size colonies (Fig. 3B).

2 Temporal trends

Demographic parameters did not show large differences among sites. Mean annual density was 12.26 ± 0.80 colonies $\cdot 400 \text{ cm}^{-2}$ at Riou ($n = 30$) and 11.36 ± 0.52 colonies $\cdot 400 \text{ cm}^{-2}$ at Maire ($n = 30$). Densities were quite stable during the study period (S2 Fig.), and recruitment rates were low at both sites: 0.98 ± 1.40 (SD) colonies $\cdot 400 \text{ cm}^{-2}$ at Maire and 0.21 ± 0.53 (SD) colonies $\cdot 400 \text{ cm}^{-2} \cdot \text{yr}^{-1}$ at Riou. Harvested colonies suffering from severe partial mortality did not show a differential "post affection" survival as they showed similar high values than unaffected

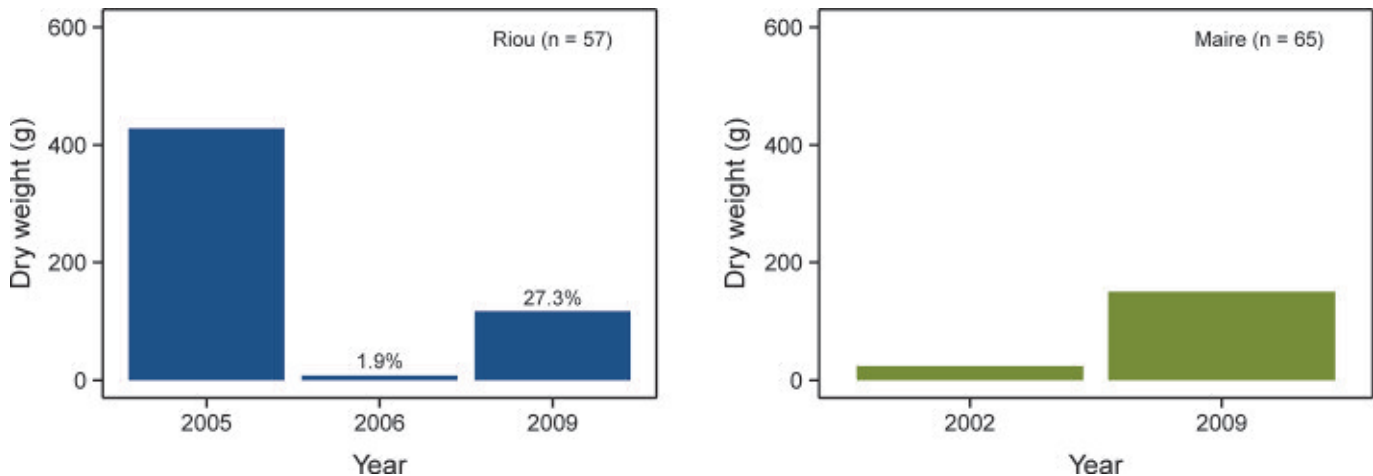


Fig 2. Biomass changes during the study period at (a) Riou and (b) Maire.

doi:10.1371/journal.pone.0117250.g002

colonies (Fig. 4A, 4B). This could explain the observed stability in adult density in both populations throughout the study period (S2 Fig.). Conversely, survival of recruits was very low and decreased rapidly at both populations (Fig. 4A and 4B).

3 Rate of recovery

The two studied red coral populations demonstrated very low rate of recovery. A small increase of biomass was observed at Riou after four years, representing 22% of the initial biomass recorded prior to the harvesting event (Fig. 2). A slight recovery was also observed by comparing size-class distribution from Riou and Maire populations after four and seven years respectively (Fig. 3). Despite a general increase in the proportion of medium size-classes, there was no recovery of the largest sizes (>100mm) and very little of colonies >80mm at Riou. At the end of the study, small colonies (0–60 mm) still represented a large proportion of the whole population (92% at Riou and 89% at Maire). Re-growth was observed in more than 75% of the harvested colonies after two years at Riou and four years at Maire (Fig. 4C). Mean number of branches in harvested colonies also increased quickly in both populations (Fig. 4D).

4 Recovery process simulations

Biomass simulations based on demographic projections showed that harvesting practices strongly influence the recovery process of *C. rubrum* populations. Harvesting practices in which the colony basis remains attached showed recovery times from 15 to 25 yr, whereas events causing total mortality (eradicating the whole colony) slowed down the recovery process, preventing the populations from recovering to initial conditions even 30 years after being harvested (Fig. 5). Riou population showed a slightly higher recovery rate than Maire population. However the influence of harvesting practices at slowing down the recovery rate when increasing total mortality was consistent for both populations (Fig. 5).

Discussion

This study provides new insights into the recovery mechanisms of a heavily exploited precious coral and illustrates how different harvesting practices can strongly modulate the recovery of affected populations. Our results also highlighted the detrimental effects of harvesting and the low rate of recovery displayed by these long-lived invertebrates.

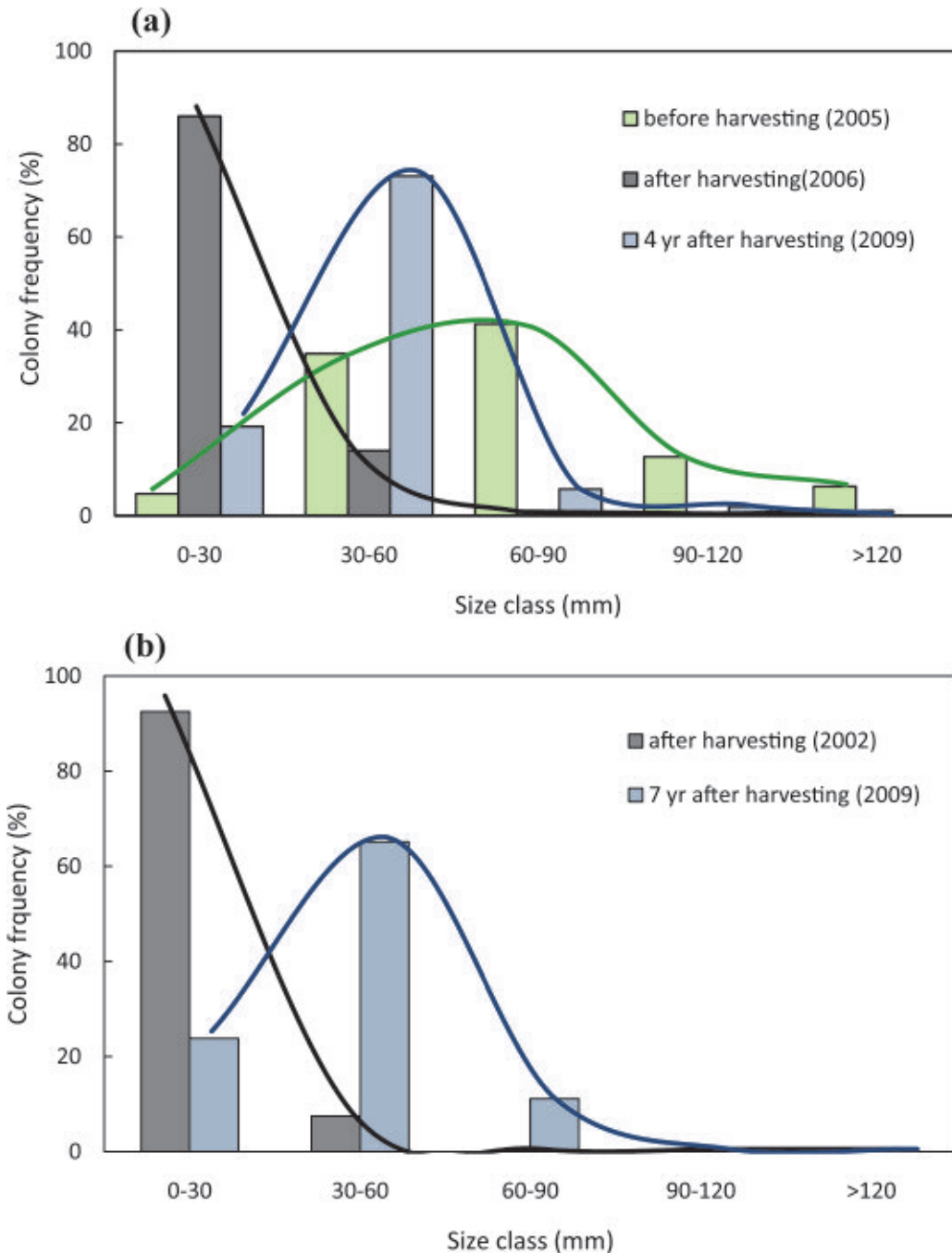


Fig 3. Size-distribution changes on harvested red coral populations. a) Size-distribution at Riou before harvesting, after harvesting, and four years after harvesting. b) Size-distribution at Maire after harvesting and seven years after harvesting.

doi:10.1371/journal.pone.0117250.g003

Harvesting heavily impacted red coral populations has led to drastic reductions of biomass and size-structure shifts towards populations dominated by small colonies. Although both populations still persisted at the end of the study period (after 4 and 7 years), they were far from initial conditions in terms of biomass and sizes. The slow recovery processes reported here confirms the long-lasting negative consequences of harvesting on slow-growing sessile organisms

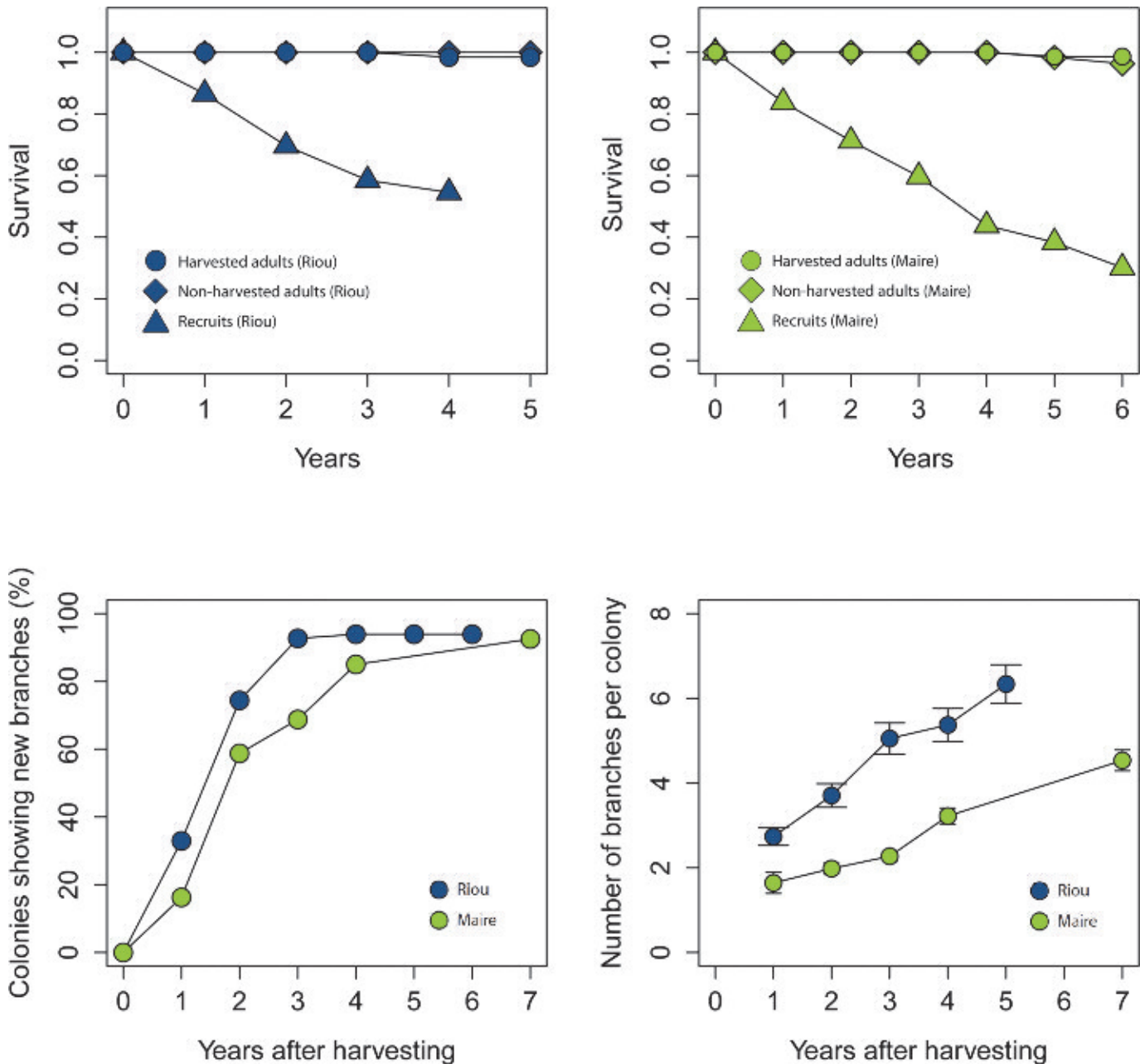


Fig 4. Demographic parameters estimated during the study period at Maire and Riou populations. Cumulative survival probability estimated at Riou (a) and Maire (b) for non-harvested colonies, harvested colonies, and recruits. (c) Temporal variation on the proportion of colonies showing new branches after fishing. (d) Temporal variation of mean (\pm SD) number of branches after harvesting.

doi:10.1371/journal.pone.0117250.g004

and indicates that extractive practices under the current legal framework are not sustainable from a long-term perspective [5].

We investigated the relative contribution of reproduction and re-growth of harvested colonies as drivers of recovery of red coral populations. The high survival of harvested colonies suffering partial mortality and fast growth of new branches demonstrated that re-growth plays a key role in recovery and persistence of *C. rubrum* (Fig 4C, 4D). Similar to previous long-term studies conducted in French and Spanish localities [13], [26], the observed low recruitment

rates and very low survival probability of new recruits suggest that recovery through reproduction is very limited (Fig. 4A, 4B). This demographic strategy is shared by other temperate gorgonian [36], and is concordant with life-history theory, which states that adult survival is inversely proportional to reproductive success [37]. Indeed, biomass simulations developed to compare different harvesting practices showed that when the whole colony is removed and thus reproduction is the main recovery mechanism, the process is much slower and it can take much more than 30 years to return to initial conditions previous to harvesting. Conversely, when fishermen leave the basal section of the colonies recovery is enhanced through re-growth of new branches (Fig. 5). The biological reference point used in this study (size-class distributions at Riou prior to harvesting) does not represent an undisturbed population; therefore, recovery leading to pristine conditions would take longer periods than those reported here. Further, we must note that the models used here do not completely account for the complexity of demographic processes in long lived species. *C. rubrum* is characterized by a low dispersal that leads its populations to be generally self-seeding [27], [38], and breeding units seem also to be restricted in space, suggesting that density may play an important role in the reproduction of this species [28]. Thus, events causing total mortality on adult colonies could lower the potential for recovery even more through reproduction [40]. Alternatively, negative density-dependent processes affecting recruitment and post-recruitment survival were also reported in corals due to intra-specific competition and resources limitation (i. e. availability of suitable substrate to settle) [39]. Further research should therefore address these questions by applying more complex modeling techniques to long-term experimental data. However, the observed large and consistent differences on population trajectories depending on harvesting procedures revealed by our study represent an important first step in the characterization of the ecological consequences of different harvesting practices on *C. rubrum* populations.

Reported recent declines on the Mediterranean yields sparked a wide debate about the conservation status of *C. rubrum*. This discussion raised questions about the sustainability of this fishery and if this species should be categorized as threatened [7], [19], [20], [21], [41]. Our results show that populations demonstrate a high degree of persistence in terms of density due to high survival of affected colonies. This may explain the existence of red coral populations despite intensive harvesting during the past few centuries. Nevertheless, we emphasize the large reduction of biomass and the size-distribution shifts toward populations dominated by small colonies likely hindered their structural function. A general simplification of benthic communities may also have negative consequences for multiple trophic levels that use these habitats as shelter during early life-stages [10], [42]. This finding should be especially relevant for policy-makers. Indeed, abundance is a parameter commonly used by international organizations such as IUCN or CITES to categorize vulnerability levels in commercial species [7], [43]. However, our results challenge the suitability of abundance-based metrics to assess the conservation status of clonal (modular) organisms by showing that large declines in biomass due to partial mortality of colonies can remain masked. Furthermore, abundance measures do not reflect the reduction in reproductive potential caused by harvesting colonial organisms, which exhibit a size-based exponential increase in reproductive output [44]. The development of a new metric focused on size and biomass parameters is crucial to assess the conservation status of precious corals and to develop sustainable fisheries management plans.

On the other hand, benthic communities are currently facing multiple perturbations derived from the ongoing global change such as warming and ocean acidification [16], [17], [18], [45]. The precautionary principle accounting for potential synergistic effects of fishing and climate warming should thus be considered in these new regulations [15], [16], [46]. While climatic perturbations are generally diffuse and difficult to minimize, fishing restrictions on no-take areas can be useful tools at a local scale to enhance the resilience of sessile invertebrates through

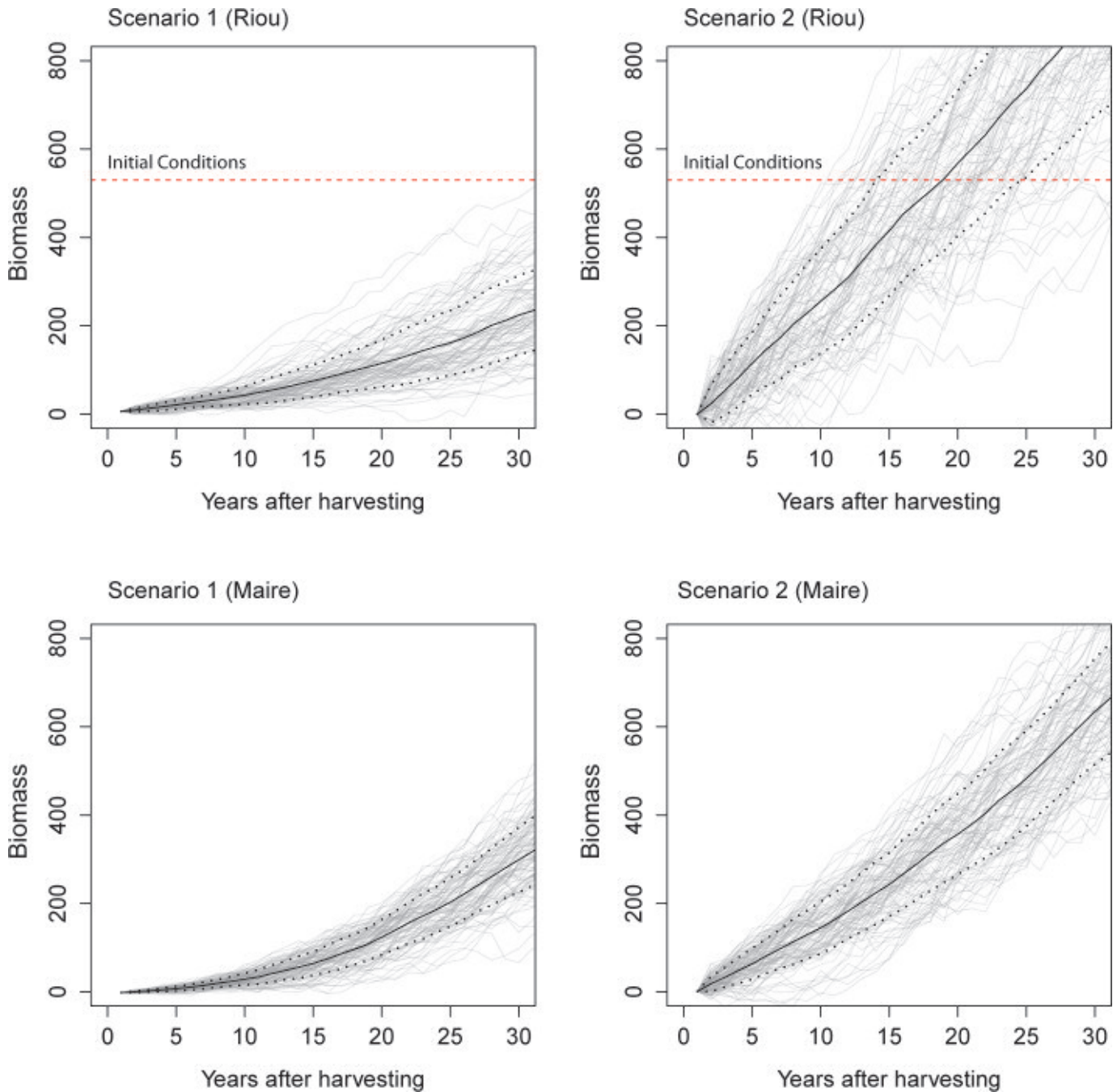


Fig 5. Biomass projections of red coral populations to compare different harvesting scenarios. Scenario 1) 90% total mortality and 10% partial mortality. Scenario 2) 10% total mortality and 90% partial mortality. Horizontal broken lines show initial conditions at Riou. Black lines represent the mean trend and dot lines the standard deviation ($n = 100$).

doi:10.1371/journal.pone.0117250.g005

increased larval production of large individuals [47]. MPAs also enhance the recovery of exploited populations in terms of biomass and size-structure [14], [26]. However given the low recruitment rates and limited dispersal of gorgonian larvae [27], it is unlikely that protected populations act as a source to exploited populations ensuring their long-term persistence [28]. Further, poaching and diving activities hinder the enhancement of coral populations' recovery

within MPAs when regulations are not enforced [27]. Therefore, we should not just consider improving the actual network of MPAs but also moving towards a more restrictive legal framework outside of the protected areas. Several new regulations have been proposed during the last decades to improve *C. rubrum* fisheries, including the establishment of annual quotas and minimum harvesting sizes and the ban on using dredges. However, these current guidelines are still largely based on untested assumptions and have failed to improve the sustainability of coral fisheries [20]. For instance, a significant step was the recent recommendation on total ban on harvesting of *C. rubrum* populations under 50m depth [22], although poaching in shallow waters is still widespread and individual countries may avoid this prohibition by developing specific management plans. Rotating systems were also proposed for precious corals fisheries in the Pacific and the Mediterranean though there still is a large uncertainty around the recovery periods of affected populations [20]. Here, we show how red coral populations may take much more than three decades to recover and that total removal of colonies can significantly slow down this process further. Our approximations are consistent with previous studies conducted within French and Spanish MPAs where populations showed only partial recovery after 20 to 30 yrs of protection [11], [14], [26]. Additionally, there has been a recent development of new jewelry manufacturing processes that allow using small pieces of coral that can be ground to powder and mixed with epoxy or other substances [22]. This development worsens the effects of harvesting by targeting whole size ranges and including entire colonies. Recovery of populations affected by these new practices might be even more jeopardized since re-growth mechanisms are dramatically limited. Finally, this study highlights the importance of long-term ecological monitoring programs because they provide reliable data on key ecological processes such as the recovery patterns of natural populations after human-induced disturbances and enable the improvement of conservation strategies.

Supporting Information

S1 Fig. Height – weight relationship obtained from death corals (n = 300).
(TIF)

S2 Fig. Annual adult densities at the study sites.
(TIF)

Acknowledgments

We acknowledge the support of all the people that helped us with the photographic surveys during all the years. The authors are part of the Marine Conservation Research Group (2009 SGR 1174) from the Generalitat de Catalunya.

Author Contributions

Conceived and designed the experiments: CL JG. Performed the experiments: CL JG JBL FZ. Analyzed the data: IMS MG FP MFV DM PD CL JG. Contributed reagents/materials/analysis tools: IMS CL PD JG. Wrote the paper: IMS CL JBL JG.

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LETTER

Accounting for Life-History Strategies and Timescales in Marine Restoration

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Keywords

Comparative demography; coral reefs; *Corallium rubrum*; integral projection models; life-history tradeoffs; octocorals; restoration; Mediterranean sea; transplants.

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Received

29 March 2016

Revised

20 December 2016

Accepted

28 December 2016

doi: 10.1111/conl.12341

Abstract

Understanding the drivers of restoration success is a central issue for marine conservation. Here, we explore the role of life-history strategies of sessile marine species in shaping restoration outcomes and their associated timescales. A transplantation experiment for the extremely slow-growing and threatened octocoral *Corallium rubrum* was highly successful over a relatively short term due to high survival and reproductive potential of the transplanted colonies. However, demographic projections predict that from 30 to 40 years may be required for fully functional *C. rubrum* populations to develop. More broadly, a comprehensive meta-analysis revealed a negative correlation between survival after transplanting and growth rates among sessile species. As a result, simulated dynamics for a range of marine sessile invertebrates predict that longer recovery times are positively associated with survival rates. These results demonstrate a tradeoff between initial transplantation efforts and the speed of recovery. Transplantation of slow-growing species will tend to require lower initial effort due to higher survival after transplanting, but the period required to fully recover habitat complexity will tend to be far longer. This study highlights the important role of life history as a driver of marine restoration outcomes and shows how demographic knowledge and modeling tools can help managers to anticipate the dynamics and timescales of restored populations.

Introduction

Marine coastal ecosystems host high levels of biodiversity and provide goods and services to a large proportion of the world's human population (Palumbi *et al.* 2008). The cumulative effects of multiple stressors such as overfishing, habitat destruction, and pollution together with new global threats (i.e., climate change and biological invasions) have driven compositional changes, local extinctions, and wholesale destruction of many benthic communities (Jackson *et al.* 2001; Airoldi & Beck 2007). To face this challenge, actions at both global (i.e., reduction of greenhouse gas emissions) and local levels are urgently needed (Kennedy *et al.* 2013). At local

scales, fishery regulations and marine protected areas can help to reduce or remove threats (Edgar *et al.* 2014). Even so, when the resilience of natural systems has been seriously diminished, active restoration may be necessary as a complementary tool to restore damaged populations and communities (Possingham *et al.* 2015).

Over the last few decades, the success of ecological restoration efforts in terrestrial landscapes has improved dramatically, with successful examples of enhancing ecosystem structure and function, and the provision of ecosystem services (Benayas *et al.* 2009; Perrings *et al.* 2015). In the marine realm, however, restoration approaches have generally been successful only at very small spatial scales and continue to present many

challenges (Edwards & Gomez 2007; Rinkevich 2015). Strikingly, the degree of success in marine restoration actions is not generally related to the underlying costs of the project (Bayraktarov *et al.* 2016). This is partly due to the high methodological constraints, but also due to relatively poor understanding of the drivers underlying successful actions.

Habitat-forming species such as corals and seagrasses have been the primary targets of marine restoration activities, and transplanting asexually produced units (i.e., coral fragments or seagrass shoots) has been proposed as the tool of choice for recovering habitats by bypassing sensitive early life stages (Edwards & Clark 1998). Recently, research has shown that corals display high survival rates after transplanting when compared to the dominant organisms found in seagrass beds, oyster reefs, or saltmarsh ecosystems (Bayraktarov *et al.* 2016). However, this generalization ignores the high diversity in life-history strategies of the dominant species in these benthic communities (Darling *et al.* 2012; Madin *et al.* 2014). Indeed, life-history tradeoffs between demographic rates have been observed in hard coral species, suggesting potential effects on short-term and long-term restoration success (Edwards & Clark 1998; Dizon & Yap 2006; Glassom & Chadwick 2006). Yet, quantitative evidence of how the life history of target species shapes restoration outcomes is lacking in the scientific literature. To advance the theoretical framework of marine restoration and provide tools to enhance the effectiveness of transplantation efforts, we need to go beyond habitat type toward a fuller quantitative analysis of how life-history patterns determine the best strategies or allow better prediction of the speed and eventual success of restoration efforts.

To date, most studies of transplant success in marine systems have focused on survival rates of transplanted individuals over relatively short monitoring periods (usually less than 2 years) (Bayraktarov *et al.* 2016). However, a broad goal of restoration efforts is to recover structural complexity that can provide ecosystem services at rates similar to natural ones. Thus, when planning restoration actions, managers should consider the factors affecting the time required from any transplantation action to reach the restoration goals for the target species and habitat. Long-term monitoring programs can provide suitable data to inform this issue, but funding often constrains the duration of monitoring after restoration actions and experiments (Precht & Robbart 2006; Lindenmayer & Likens 2009). Demographic modeling methods such as matrix and integral projection models (IPMs) (Morris & Doak 2002; Ellner & Rees 2006) can be used to synthesize individual data into predictions of the longer term development of transplanted populations (Linares *et al.* 2008).

In the present study, we combined demographic monitoring of transplanted and natural colonies of a temperate coral species, a comprehensive literature review of tradeoffs in the life histories of sessile marine species, and the use of population projection models to explore the dynamics of transplant efforts targeting species with different life histories. Our results support the utility of explicitly linking life-history theory to marine restoration and provide an illustrative example of anticipating the expected dynamics and timescales of restored ecosystems.

Methods

Study system

The precious red coral *Corallium rubrum* is a structural octocoral of a highly diverse coralligenous assemblage of the Mediterranean Sea and also possesses important cultural and economic value. Due to historical overexploitation, most shallow populations of *C. rubrum* can be considered functionally impaired and many are ecologically extinct (Bruckner 2009; Tsounis *et al.* 2010). To reverse this situation, an international agreement urged Mediterranean countries to strengthen their *C. rubrum* fishery regulations during the last decade (Cau *et al.* 2013). Unfortunately, the lack of enforcement of regulations on coral harvesting along with poaching is widespread across the Mediterranean basin and represents a major problem for the management of the species, hindering the effectiveness of its conservation (Linares *et al.* 2012).

Study area and transplant experiment

In 2011, the Catalan authorities intercepted 14.5 kg of illegally harvested *C. rubrum* along the Montgrí Coast (Catalonia, Spain). About 300 red coral colonies, a small portion of the intercepted colonies, were selected for a transplant experiment. These colonies were initially kept at 16 °C and fed in aquarium facilities at the Institute of Marine Sciences in Barcelona (Spain). After 1 week, the colonies were transported in coolers to the Parc Natural del Montgrí, Illes Medes i Baix Ter in the NW Mediterranean and transplanted onto a rocky wall ranging from 15 to 17 m depth using a two-component epoxy putty as glue. The site was chosen because some sparse red coral colonies were found in the vicinity, indicating its suitability for the species (Figure 1).

Demographic traits

Four transects were established within the transplanted population and surveyed through photographic sampling after transplantation, in May 2011, and again in

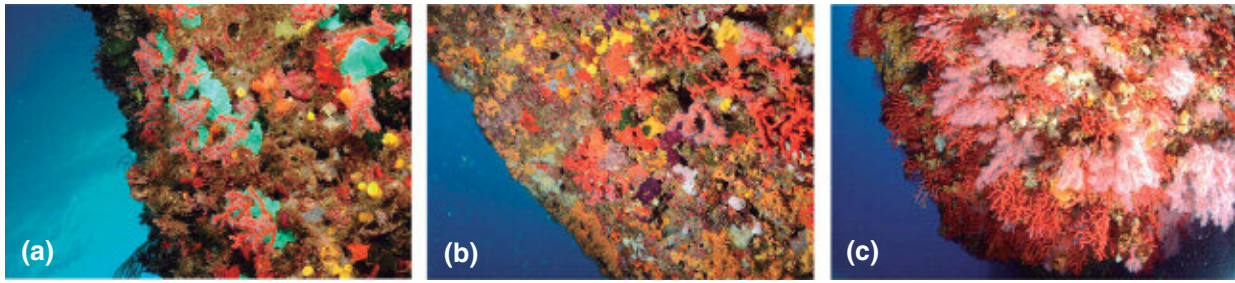


Figure 1 Restoration of *Corallium rubrum* populations. (a) A new population was transplanted in 2011; (b) most transplanted colonies survived in 2015, after 4 years of transplantation; and (c) natural well-protected *C. rubrum* populations with large colonies were used as a baseline to assess the time periods required for restoration actions. Images: J. Garrabou.

May 2015 (Figure 1). Survival rates of the transplanted colonies were quantified by individually identifying coral colonies from the photographic series from 2011 and 2015. Natural survival rates of *C. rubrum* colonies were calculated from long-term data on eight natural populations. (See Supplementary Methods for a complete description of surveys.) Reproductive potential of colonies was estimated for a sample of transplanted colonies ($n = 35$) outside the monitored transects and from a natural adjacent population ($n = 35$) in late June of 2015 by counting *C. rubrum* larvae found inside the polyps of the fertile female colonies (Tsounis *et al.* 2006). Samples were collected by SCUBA diving and fixed in 4% formaldehyde. At the laboratory, 15 polyps per sample were dissected and larvae found inside the polyps were counted.

Literature review

We explored life-history tradeoffs in marine restoration experiments following two steps. First, we systematically reviewed all transplantation experiments of marine sessile species that we could identify in a search of the literature up to November 2015. Using *Google Scholar*, we searched for a combination of the terms “restoration,” “transplantation,” or “rehabilitation” with a second term related to marine sessile taxa: “coral,” “gorgonian,” “sponge,” “macroalga,” or “seagrass.” We then selected those studies that conducted experimental transplants as a restoration technique and reported survival rates at least 1 year after transplanting.

We also compiled data on growth rates of sessile marine species since this vital, or demographic, rate is highly correlated to overall life history (Darling *et al.* 2012). We searched available studies reporting standard data on linear extension rates to approximate average species-specific growth rates for corals (Madin *et al.* 2016). In seagrasses, mean horizontal rhizome elongation rates were used as an indicator of the species growth rate (see Marbà

& Duarte 1998); thus, seagrass and sessile invertebrates were analyzed separately.

Demographic projections

Red coral

To synthesize data on growth, survival, and reproduction into predictions of population growth and increasing sizes within populations, we used IPMs parameterized with long-term demographic data from several natural red coral populations. Full description of data analysis and model construction are given in the Supplementary Material S1. Based on annual IPMs, we computed 1,000 stochastic projections assuming that all annual models can occur with equal probability at each time-step. Maximum height of transplanted colonies was measured in 2015 using photogrammetric techniques (Drap *et al.* 2013) and the distribution of heights was used to establish the starting population vector for the projections.

Linares *et al.* (2010) argue that the structural complexity of *C. rubrum* populations can be assessed by quantifying the proportion of large colonies (>100 mm), since these larger colonies provide structural complexity. Based on this parameter, we compared the outputs from our population projections to the proportion of large colonies in three relatively unimpacted *C. rubrum* populations that are located within old and well-enforced Mediterranean marine protected areas (Figure 1 & S1, Linares *et al.* 2010).

Comparative analyses

We also searched the literature for published matrix population models of other marine sessile species. We then used these models to perform deterministic population projections of 100 individuals starting at the smallest size class and computing time periods until the population reached a proportion of large individuals (largest size

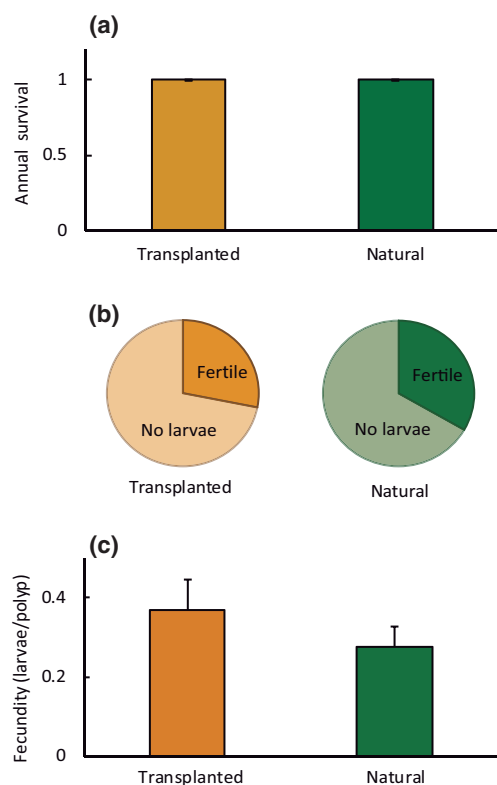


Figure 2 Demographic traits in transplanted and natural *C. rubrum* populations. (a) Mean annual survival rates; (b) proportion of fertile colonies; and (c) mean polyp fecundity, calculated as the frequency of larvae found per polyp within fertile colonies.

class) equivalent to the 20% and 80% of the expected proportion when reaching the stable stage distribution.

Results

Demographic traits of *C. rubrum* transplanted colonies

After 4 years, 99.1% of transplanted *C. rubrum* colonies were still alive. Annual survival rates of transplanted colonies did not show significant differences from control populations (Figure 2a). Transplanted colonies also had similar reproductive potential to colonies in natural populations, considering both the proportion of fertile colonies and the frequency of larvae per polyp (Figure 2b & c).

Comparative survival and growth in transplant experiments/actions

We found 50 studies that allow calculation of mean annual survival rates after at least 1 year following transplanting for a total of 59 marine structural species

(Figure 3a). These included 40 species of hexacorals, which have a mean annual survival of 60.8% (range of 6.8–98.6%); five species of gorgonians, including the present study, with mean annual survival of 48.1 (range of 30.0–99.1%); one species of sponge, with mean annual survival of 85.7%; 11 species of seagrasses, with mean annual survival of 42.5% (range of 28.9–69.2%), and two seaweeds, with mean annual survival of 43.1% (range of 25.1–80.0%). We observed a significant negative correlation between survival after transplantation and the species mean growth rates measured under natural conditions in marine sessile invertebrates (Figure 3b; $n = 35$; Pearson's $r = 0.47$, $P = 0.005$; Spearman rho = 0.37, $P = 0.046$). Seagrass species revealed a parallel pattern (Figure 3c; $n = 8$; Spearman rho = 0.81, $P = 0.022$), although the relationship was only marginally significant according to Pearson's correlation (Pearson's $r = -0.69$; $P = 0.059$). Growth data measured in natural and transplanted colonies for coral species were also highly correlated (Figure S3, $n = 17$; Pearson's $r = 0.85$; $P < 0.001$).

Demographic projections and recovery periods

Red coral

The transplanted population in 2015 was dominated by small individuals (most red coral colonies were < 35 mm in height, Figure 1 & S1), while natural red coral colonies had extremely low growth rates (Figure S2). The stochastic IPMs incorporating these traits showed that a period ranging from 30 to 40 years after transplanting is needed for populations to have a proportion of large colonies comparable to that seen in the well-preserved *C. rubrum* populations used as an ecological reference (Figure 4).

Comparative analyses

The simulated recovery periods for 41 marine sessile species were highly variable in length, ranging from years to several decades (Figure 5 & Table S3). The expected recovery length was strongly and positive associated with the species' mean survival rate regardless of the conservation goal ($n = 41$; 20% threshold: $R^2 = 0.419$; $P < 0.001$; 80% threshold: $R^2 = 0.495$; $P < 0.001$). After accounting for potential artifacts due to different matrix dimensions, mean survival rates were still a strong predictor of the expected recovery periods (Table S1).

Discussion

Marine restoration is a relatively young discipline with most efforts only operating at very small spatial scales (Bayraktarov *et al.* 2016). Filling knowledge gaps on the processes underlying restoration success is therefore

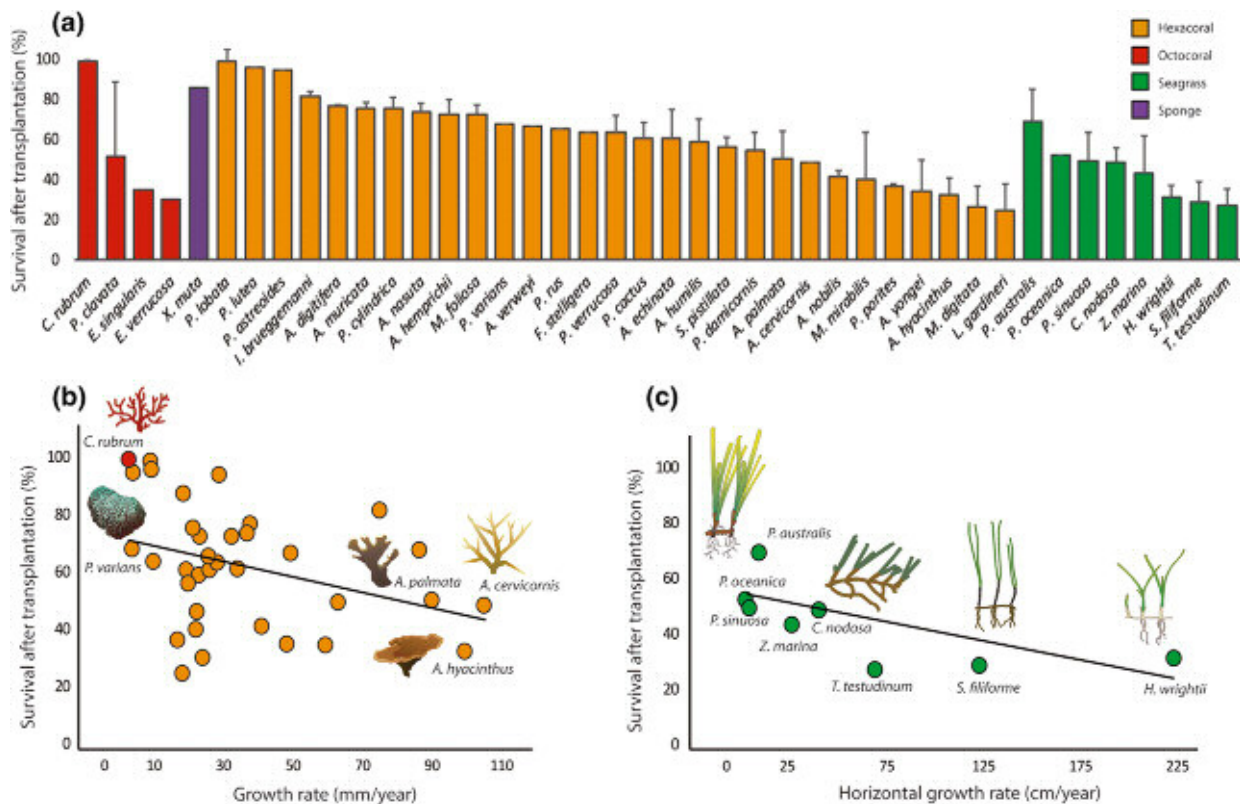


Figure 3 Survival rates of marine sessile species in transplant experiments. (a) Mean annual survival rates (Mean \pm SE). (b) Life-history tradeoff between survival after transplantation and growth rates in 35 marine sessile invertebrate species and (c) life-history tradeoff between survival after transplantation and growth rates in eight seagrass species. Each dot represents a species for which mean annual survival after transplantation and mean growth rate could be calculated from a range of published studies (see Table S1). In seagrass species, growth represents mean horizontal rhizome elongation rate (see Marbà & Duarte 1998). Images: Integration and Application Network, University of Maryland Center for Environmental Science (ian.umces.edu/symbols/).

crucial to help further develop this field and ensure meaningful planning and success over larger spatial and temporal scales. In this study, we quantify the role of life history in shaping restoration outcomes and demonstrate a consistent tradeoff between survival and growth across different taxa with contrasting life history and functional traits, which in turn drives a tradeoff between required minimal transplantation effort at the start of a project and the minimum possible speed of ecosystem recovery.

Anticipating mortality patterns after transplantation is central to the design of any restoration action since it may determine the initial attaching effort required to achieve specific conservation goals. Here, a systematic review of transplantation experiments from tropical and temperate habitat-forming species revealed a negative tradeoff between growth and survival after transplantation that was supported in spite of differences in experimental techniques and physical properties of the environment that were not explored (Figure 3b). Slow growing massive hard corals such as *Porites*

astreoides and *P. lutea* and the sponge *Xestospongia muta* showed the highest survival after transplantation, with rates ranging from 86% to 98%. On the contrary, fast-growing corals such as *Acropora cervicornis*, *A. yongei*, and *A. palmata* had survival rates that ranged from 35% to 44%. These results were consistent with previous transplant experiments in tropical coral species with contrasting life histories (Edwards & Clark 1989; Dizon & Yap 2006). Similarly, among seagrasses, the three slow-growing *Posidonia* species showed higher shoot survival after transplantation (from 49% to 69%) compared to faster growing *Syringodium filiforme* (29%) and *Halodule wrightii* (27%) (Figure 3c). Our findings are in agreement with allocation theory, which predicts that tradeoffs between vital rates such as growth, reproduction, and survival may arise from energetic constraints acting at physiological levels (Stearns 1989). Further, branching morphologies associated with faster life histories may increase exposure to physical damage and result in higher mortality rates (Madin et al. 2014).

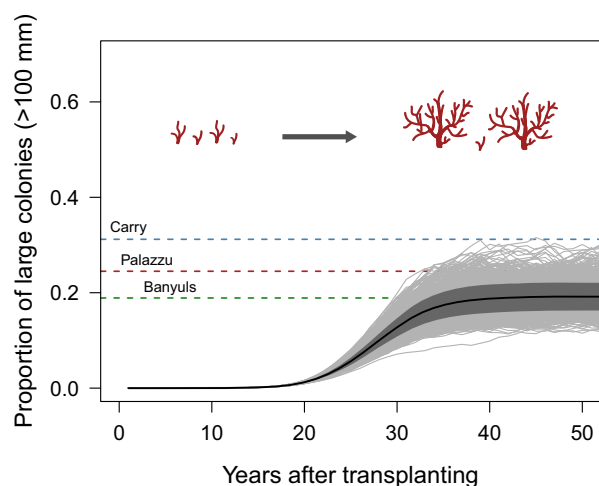


Figure 4 (a) Predicted temporal dynamics of the *C. rubrum* population size frequency distribution. Black line represents the mean and shaded area represents the standard error of 1,000 stochastic projections. The size frequency distribution of three natural and well-protected *C. rubrum* populations (dotted lines) was used as an ecological baseline (Linares et al. 2010).

There are also a number of external drivers that can strongly influence restoration success such as predation and herbivory, density of transplants, and catastrophic events (Shaish et al. 2010; Gomez et al. 2014). In spite of the clear importance of these effects, our results show that species' life histories can still provide strong predictive power concerning the outcome of trans-

plantation projects. Better understanding of both the intrinsic and extrinsic drivers of mortality patterns after transplantation would be ideal and could lead to the implementation of more successful restoration designs, since this combined approach can better define both anticipated time periods for restoration and also the relative benefits of direct transplantation effort.

There have been major international calls to ban the international trade in precious coral and to implement management regulation aimed to ensure the conservation of these species (Bruckner 2014). Yet, the feasibility of restoration actions for these emblematic species has remained uncertain and this may hinder the potential for development of future restoration plans. In the present study, we observed that the colonies of the octocoral *C. rubrum* were extremely resistant to the stress of transplantation, displaying high survival rates similar to those in natural populations (Figure 2a, Garrabou & Harmelin 2002). It is remarkable to observe this high survival rate in transplanted *C. rubrum* colonies that were subject to the stresses of being harvested, kept out of the water in the poachers' nets, transported, maintained in aquaria for 1 week, and then transplanted back into natural habitat. Yet, these transplanted *C. rubrum* colonies had a similar proportion of fertile colonies and even higher frequency of larvae per polyp after 4 years than observed for colonies in natural populations (Figure 2b & c). Assessing reproductive potential is also critical when working with most marine sessile species which, like *C. rubrum*, show limited larval dispersal and

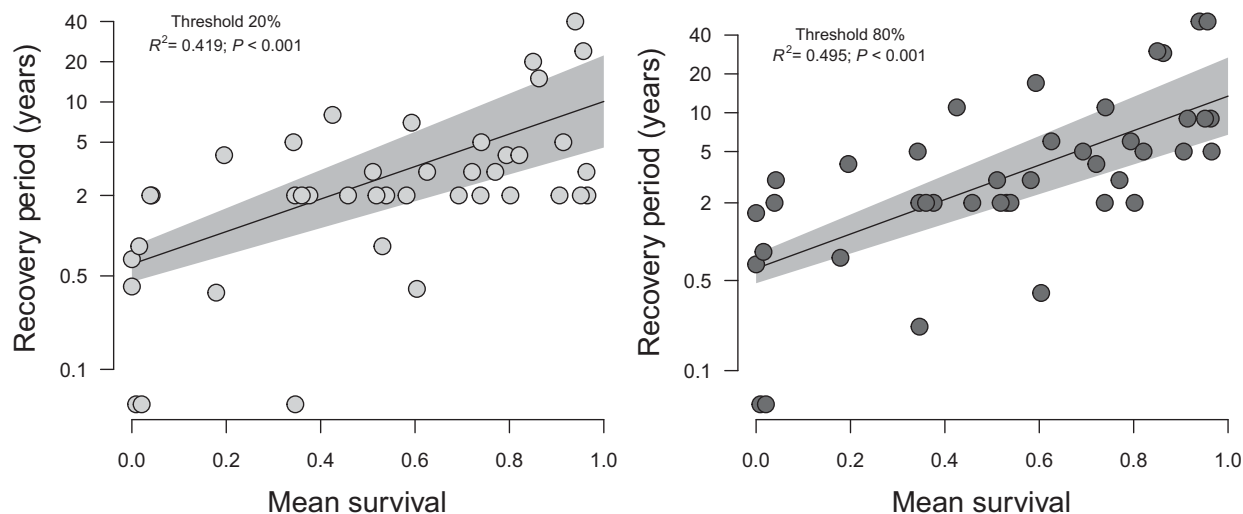


Figure 5 Projected recovery times for 41 marine sessile invertebrate species using published matrix populations models (Table S3) and setting a recovery threshold of (left) 20% and (right) 80% of the number of large colonies expected in a population at the stable stage distribution. The black lines and the shaded areas correspond to the mean and standard errors of the linear models (Table S1).

high self-recruitment rates (Ledoux *et al.* 2010). Indeed, to effectively recuperate populations through a single transplantation effort, newly restored populations must also be viable in the long term, with reproduction reaching natural rates. Here, the high survival and reproductive potential displayed by transplanted *C. rubrum* confirmed the potential success of this restoration action and strongly support the feasibility of these techniques, at least at local spatial scales, with potential applications for other long-lived precious coral species.

As important as choosing a suitable species and restoration method is considering the appropriate time scale and ecological baselines over which to evaluate restoration outcomes or to expect the restoration of ecological functions (Bull *et al.* 2014). Stochastic projections developed here revealed that periods ranging from 30 to 40 years may be necessary for newly established *C. rubrum* populations to show a colony size distribution comparable to those observed in well-preserved natural populations (Linares *et al.* 2010). These results suggest that, similar to relatively fast-growing terrestrial forest systems (Vesk *et al.* 2008), long-lived coral stands can take up to several decades to recover their functionality and to allow the development of associated organisms, such as fish and invertebrates, as may occur in tropical coral reefs (e.g., Cabaitan *et al.* 2008). More interestingly, we found that potential recovery periods can be accurately predicted by the specific mean survival, demonstrating the strong influence of the species' life histories on the temporal scales associated with restoration actions (Figure 5).

Overall, this study demonstrates a tradeoff between initial transplantation effort needed to achieve a target density of individuals and the speed of recover that may be achieved in a restoration action. For instance, targeting fast-growing species such as *A. cervicornis* or *A. hyacinthus* (with survival rates ranging from 40% to 50%) will require a twofold to threefold initial amount of attached colonies to obtain the same density of survivors compared to actions targeting slow-growing-resistant species such as the red coral *C. rubrum* or the massive coral *P. lutea*. On the other hand, life histories of the target species will also have a strong effect on the expected recovery periods that may vary as much as 20–30 years (Figure 5). Finally, because life history and functional traits are highly correlated (Adler *et al.* 2014), favoring specific strategies can have long-term consequences for habitat complexity and ecosystem responses to global change (Ortiz *et al.* 2014).

Acknowledgments

The authors thank the Agents Rurals de Catalunya for their invaluable work against red coral poaching.

We are indebted to A Lorente for his support. We thank E Aspillaga and L Navarro for field assistance. P Capdevila, A Gori, K Kaplan, A Medrano, and two anonymous reviewers provided valuable comments on the manuscript. Funding was provided by the Spanish MINECO (CTM2009-08045 and CGL2012-32194), the Oak Foundation, the TOTAL Foundation Perfect Project, and the European Union's Horizon 2020 research and innovation programme under grant agreement No 689518 (MERCES). This output reflects only the author's view and the European Union cannot be held responsible for any use that may be made of the information contained therein. IMS was supported by a FPI grant (BES-2013-066150), CL by a Ramon y Cajal (RyC-2011-08134), and JBL by a Postdoctoral grant (SFRH/BPD/74400/2010). Authors are part of the Medrecovery group (2014SGR1297).

Supporting Information

Additional Supporting Information may be found in the online version of this article at the publisher's web site:

Figure S1. Population size frequency distribution of the transplanted colonies represented by the brown bars and the brown line.

Figure S2. Size-dependent growth of *C. rubrum* colonies based on data from two natural populations.

Figure S3. Growth rates in natural versus transplanted coral colonies (log-scale) for 17 coral species.

Table S1. Summary statistics for the linear model fitted to the expected recovery periods.

Table S2. Source data of survival in transplant experiments and growth references of the marine sessile species included in the comparative analysis.

Table S3. Source data of population matrix models used to project potential recovery periods.

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