

ECOLOGICAL NICHE MODELLING OF FIVE GORGONIAN SPECIES WITHIN THE SHALLOW ROCKY HABITAT OF THE FRENCH MEDITERRANEAN COAST

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GORGONIANS
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MARINE PROTECTED AREA
DISTRIBUTION
HABITAT SUITABILITY
MEDITERRANEAN

Abstract.- Engineer species, such as gorgonians, provide several ecosystem services and play a significant role in the maintenance of biodiversity. We tested an ecological niche modelling approach using a unique dataset of spatialized inventories on a regular grid (< 800 m) replicated in 2013 and 2020 along 450 km of coastline for five gorgonian species commonly found in the Mediterranean shallow habitats (10-50 m deep). Ten non-correlated environmental predictors derived from the most advanced geomorphological and hydrological data were used to assess the ecological niche of the five species using maximum entropy (Maxent). The model confirmed that different combinations of depth, rugosity, sea surface and bottom temperature, bottom flow speed and turbidity explain the different gorgonian's distribution. The reduction of the number of presence observations did not alter the quality of the prediction as long as the observation points were spread over the entire variability range of predictors. The latter can be achieved by only including presence observations from highly protected zones of the marine protected areas of the region. Our results provide a greater understanding of the factors shaping the distribution of five gorgonian species commonly found in Mediterranean shallow areas.

INTRODUCTION

The maintenance of the ecosystem services provided by the ocean and that contribute to human welfare is one of the 17 sustainable development goals identified by the General Assembly of the United Nations in 2015 (<https://sdgs.un.org/fr/goals>). Such an objective is encompassed in the European Marine Spatial Planning framework which aims to take into account the interactions between different human activities at sea, including biodiversity protection (DIRECTIVE 2014/89/UE). One-step in marine spatial planning is mapping biodiversity and the ecosystem services it delivers, together with understanding its drivers, in order to be able to forecast its evolution under

different development scenarios (Bailey 2010).

During the last 20 years, in the Mediterranean Sea, several research programs focused on obtaining spatial information on marine habitats and the distribution range of marine species in order to begin implementing marine spatial planning (Gerovasileiou *et al.* 2019). In this region, marine species inventories, however, have generally been restricted to a limited number of locations, often in Marine Protected Areas (MPAs), although recent citizen science projects attempted to expand this spatial scale (Bramanti *et al.* 2011, Ponti *et al.* 2019). Ecological niche modelling (ENM) can be a powerful complementary tool to refine species

habitat mapping. It currently relies on correlative approaches, which are based on the assumption that the suitable environment of a species controls its range and allows one to project species distribution range in the geographical space using environmental predictors (Sillero *et al.* 2021). Early ENMs used statistical modelling such as Generalised Linear Modelling (GLM, McCullagh *et al.* 1983) relating presence-absence to environmental descriptors. Machine learning models that use presence-only data (Maxent, Phillips *et al.* 2006, Random Forest, Breiman 2001), together with satellite imaging boosted ecological niche modelling applications at large spatial scale. The performance of various ENM algorithms have been systematically compared, showing species-related performance and suggesting no single algorithm outcompetes others strengths and limitations (reviewed in Guisan *et al.* 2017). An important conclusion about the performance of ENM is their sensitivity to the number of environmental predictors used (Sainz-Villegas *et al.* 2022). The ensemble averaging of multiple ENM outputs has been proposed to improve species spatial distribution mapping (Achu *et al.* 2021). Still, this improvement blurs the explanation link of the environmental predictors. Among ENMs, the machine-learning model Maxent is one of the most tested. It displays ecologically meaningful responses to environmental predictors and is less data (presence-only) demanding than GLM, its statistical counterpart that requires more extended datasets with presence and absence (Elith *et al.* 2006, Wisz *et al.* 2008, Sainz-Villegas *et al.* 2022). Specifically, using presence-only data avoids the assumption that the absence of a species means habitat unsuitability, when in fact it could be the result of either disturbance or ineffective dispersal (Jarnevich *et al.* 2015). The most commonly reported sampling biases are geographic biases - such as partial distributions and environmental biases - leading to either unreliable results or to niche estimates with underrepresented subsets of the population environmental space (Sillero *et al.* 2021). The pressures on the habitats can also bias model construction. Indeed, the absence of a species can result from anthropogenic pressures irrespective of the characteristics of the habitat to support this species, that should be disregarded in ENM appli-

cations. Highly protected zones within MPAs might therefore be preferred locations for presence observations in order to limit confusion between predictors of the ecological niche and habitats degraded by direct anthropogenic pressures. In the present study, we aimed to develop an ecological niche model for the most common gorgonian species in the NW Mediterranean, using an extensive dataset of presence observations covering a wide range of environmental conditions in the NW Mediterranean Sea, inside and outside of MPAs.

Gorgonians (octocorals with an internal skeletal axis, McFadden *et al.* 2022) are among the most emblematic species of the Mediterranean Sea, playing an essential role in shaping and structuring the biological diversity on the rocky temperate marine habitats (Gili & Coma 1998, Otero *et al.* 2017). Among the Mediterranean gorgonian species, five are common in the rocky communities of shallow (0-50 m) coastal waters (Weinberg 1979a): *Eunicella singularis* (Esper, 1791), *Leptogorgia sarmentosa* (Esper, 1789), *Eunicella cavolini* (Koch, 1887), *Paramuricea clavata* (Risso, 1827) and *Corallium rubrum* (Linnaeus, 1758). The five species have a wide Mediterranean Sea distribution but also extend (except *E. cavolini*, López-González, 1993) to the Atlantic Ocean on the coasts of Morocco (Zibrowius *et al.* 1984, Grasshoff, 1977) and Portugal (Cúrdia *et al.* 2013, Boavida *et al.* 2016).

In the Mediterranean Sea, they have mainly been studied in the northwestern basin (Carpine & Grasshoff 1975, Cocito *et al.* 2002, Gori *et al.* 2011a, Di Camillo *et al.* 2018) although their distributions extend to the Tunisian and Sicilian coasts of the Ionian basin (Canessa *et al.* 2023), in the Aegean Sea (Salomidi *et al.* 2009, Vafidis, 2009, Çinar *et al.* 2018) and the Sea of Marmara (Topçu & Öztürk 2015). No populations were found in the easternmost part of the Mediterranean basin but this absence could be due to a lack of data (Otero *et al.* 2017). *Paramuricea clavata*, *C. rubrum* and *E. cavolini* have a wide bathymetric distribution from 10 m to more than 1000 m water depth (Grinyó *et al.* 2016, Knittweis *et al.* 2016, Carugati *et al.* 2022).

In this study, we analysed the spatial distribution of *E. singularis*, *L. sarmentosa*, *E. cavolini*, *P. clavata* and *C. rubrum* in their shallow bathyme-

tric range (10-50 m) in the eight major rocky habitat spots of the French Mediterranean coasts (spanning 3.5 degrees of longitude, from Toulon to Cerbère), using spatialized inventories performed on a regular grid (< 800 m). The ecological niche and distribution area of suitable habitats were modelled with Maxent software to identify the main environmental predictors that influence the spatial distribution of the five species. We tested the sensitivity of the fitted model to a reduced number of occurrences and their geographic position, and specifically assessed whether sampling limited to the five highly protected areas of MPAs in the study area (which can be considered exempt from the main direct anthropogenic pressures impacting gorgonians such as anchoring, nets, scuba diving, etc.) would affect the model predictions.

MATERIALS AND METHODS

Study area

The study area extends along 450 km of the French Mediterranean coastline, from Toulon to Cerbère, and from the coast down to the 50 m isobath. Along this coastline, the seabed mainly consists of a soft bottom with a few small patches of rocky habitat between 76 (VLR) and 3123 hectares (AGM). A set of eight sites (National Park of Port Cros: PNPC, Parc Marin de la Côte Bleue: PMCB, Aigues-Mortes: AGM, Aresquiers: ARES, Agde: AGD, Valras: VLR, Leucate: LEU, Côte Vermeille: CVM, Fig. 1) within this fragmented rocky habitat were investigated to assess the niche preferences of the five species: *Eunicella singularis*, *Eunicella cavolini*, *Paramuricea clavata*, *Leptogorgia sarmantosa* and *Corallium rubrum*. Of the eight sites, five include a high protection zone (according to the classification of Horta e Costa *et al.* 2016: Fully protected) with different surface areas (PNPC: 1232 ha, AGM:100 ha, AGD: 310 ha, CVM: 69 ha, PMCB: 289 ha, Fig. S1, Supplementary Material).

Data collection

Species distribution: The population density of the five species was assessed between 2013 and 2020 in 696 evenly spaced geo-referenced locations covering the hard bottom habitat in all sites except in PNPC where locations were selected based on the known presence of the species *P. clavata* (Guizien *et al.* 2022, <https://cardobs.mnhn.fr/>, Fig. 1 and Fig. S1, Supplementary Material). The spacing between sampling points varied from 100 m to 800 m, depending on the bathymetrical steepness of each zone. Using a regular grid in order to avoid the bias of preferential sampling enabled us to estimate that no spatial auto-correlation was observed (Legendre *et al.* 2002, Merckx *et al.* 2011). For the five species, a strong nugget effect (Carrasco 2010) at eight meters

distance was observed on empirical variograms (Fig. S2, Supplementary Material). In each georeferenced location, all individuals taller than 2 cm (i.e., older than one year) were counted either by direct visual census of autonomous scuba divers or on photographs in four to nine quadrats of 1 square meter. These surface areas were larger than the minimal area of 2 square meters established for shallow water Octocorallia communities (Weinberg 1978). The georeferenced database of the five species presence and absence was implemented in a GIS software (© QGIS).

Habitat suitability model

The Maxent model: Maxent is a machine learning software recognized as the most efficient method in species distribution modelling when presence-only species records are available (Elith *et al.* 2006, Hernandez *et al.* 2006, Hernandez *et al.* 2008, Bargain *et al.* 2018). Natively, Maxent estimates Receiving Operator Curve, more explicitly called Relative Occurrence Rate (ROR, Merow *et al.* 2013) which is defined as the probability for an observed individual to be found in a set of environmental conditions by maximising its entropy probability distribution (i.e., the most spread out or closest to the uniform, Phillips *et al.* 2006). Hence, Maxent is a niche or habitat suitability model whose predictions, although represented on a spatial grid, relate to environmental predictors only, disregarding spatial location arrangements except if geographical predictors are explicitly included in the set of environmental descriptors (Elith *et al.* 2011). The Maxent version 3.4.4 was used. Setting up the Maxent model requires defining a background grid on which environmental predictors are known, features, regularization multipliers, sampling biases, output types and diagnostics outputs (see Supplementary Material for details). The quality of Maxent fits is primarily assessed by the Area Under the ROR Curve (AUC). The AUC ranges from 0 (fit worse than random) to 1 (perfect fit). Below 0.5 the prediction is not better than at random, and above 0.7 the prediction is reliable (Zweig & Campbell 1993).

Data on marine habitats were retrieved from CARTHAM website (<https://geo.data.gouv.fr>) and the background grid covered the only hard bottom habitat where the five octocoral species exclusively dwell. Gridding the hard bottom habitat in the eight sites resulted in a set of 8942 georeferenced 100 m x 100 m cells (background grid), according to availability of environmental predictors data (Fig. S1). Overlaying the 8942 background grid cells with the 696 sampling sites (mentioned above) resulted in 444 grid cells where both environmental predictors were defined and the five species presence/absence was assessed. Across the 444 grid cells, *Corallium rubrum* was present in 16 grid cells, *P. clavata* in 18 grid cells, *E. cavolini* in 31 grid cells, *L. sarmantosa* in 62 grid cells and *E. singularis* in 255 grid cells. The implementation of habitats and masks was carried out in © QGIS software. The environmental predictors were selected within all available geomorphological and hydrological potential predictors, to which were added latitude and longitude as explicit

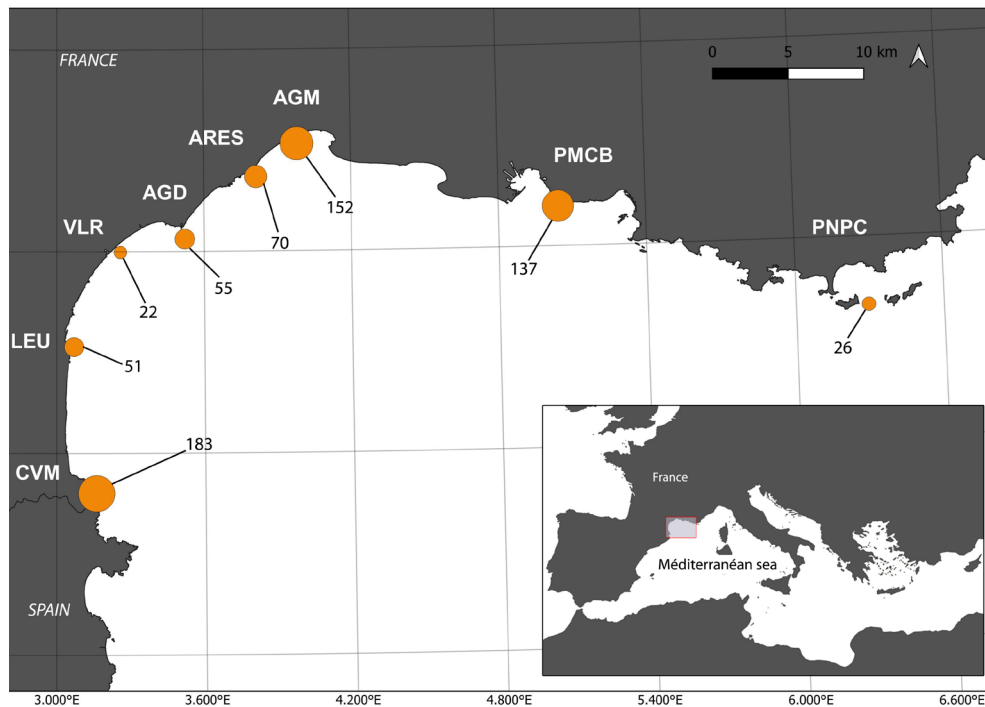


Fig. 1.- Map displaying the number of sampling locations in each of the eight sites of the NW Mediterranean Sea. Black rectangles highlight the sampled sites. The diameter of the orange circles is proportional to the number of sampling points at each site.

predictors in some modelling test.

Geomorphological predictors were calculated from raw bathymetrical dataset that combined three-dimensional point seedlings from Airborne Laser Imaging Detection and Ranging System (LIDAR) for water depth less than 15 m (LIT-TO3D, www.diffusion.shom.fr) and multibeam echosounder for water depth larger than 15 m, except in PNPC where LIDAR data extended down to 40 m water depth. LIDAR data was collected between 2014-2015, except in the CVM zone where it was obtained in 2011. Multibeam echosounder data was assembled from multiple collection periods between 2005 and 2018 and from different operators (Seaview for AGM, ARES and VLR, CEFREM for CVM, Semantic TS for AGD, Copethec and Mesuris for PMCB). Density of LIDAR data decreased from 10 to 0.1 points per square meter, when water depth increased. Density of multibeam echosounder data decreased from 25 to 4 points per square meter. Seven geomorphological metrics commonly used for habitat predictive modelling were computed for each cell of the background grid. These seven metrics describing the sea-floor topography were: bottom depth (unit: m), slope (unit: angle degrees), roughness (unit: m), rugosity (no unit), terrain ruggedness index (TRI, unit: m), eastness (no unit) and northness (no unit) orientation (Wilson *et al.* 2007, see Supplementary Material for details). These metrics were calculated using all raw bathymetrical data within a 20 m x 20 m slab located in the centre of each cell of the background grid. Hydrological predictors included turbidity and tempe-

rate, which were measured at the sea surface by satellite imaging (Aquamodis - level-3 global browser: <https://oceancolor.gsfc.nasa.gov/l3/>), and bottom temperature and flow speed, simulated by a coastal ocean circulation model (Briton *et al.* 2018). Satellite observations came from the Moderate Resolution Imaging Spectroradiometer (MODIS) on board the AQUA satellite. Data was acquired in 36 spectral bands at three resolutions (250, 500 and 1000 m) and covered all the globe in one to two days. Sea Surface Temperature (SST, unit: °C, 4 µm night time) and turbidity (unit: m⁻¹, Diffuse attenuation coefficient at 490 nm) monthly averages at 4 km spatial resolution were extracted between 2003 and 2018 for Turbidity and 2006 and 2018 for SST over the extent of the background grid. For each predictor (SST, Turbidity), the pluri-annual mean was calculated and interpolated at each background grid point using the `griddata` Matlab function with the default linear option. Although Chlorophyll-a concentration is one of the MODIS products (Chla, unit: mg.m⁻³, OCx Algorithm), it was not considered because data was missing in the PMCB site. However, Chlorophyll-a concentration significantly correlated with turbidity in all other sites (Spearman determination coefficient $R^2 = 0.68$, $p < 0.0001$). Bottom temperature and flow speed came from hourly outputs of a high-resolution SYMPHONIE 2015 regional circulation simulation, which ran from January 2010 to December 2012 (80 to 300 m horizontal resolution in the eight sites, Briton *et al.* 2018, Vissenackens *et al.* 2023). Monthly statistics of flow speed (tenth and ninetyth

percentiles, unit: m.s-1) and temperature (tenth and ninetieth percentiles, unit: °C) hourly values at the bottom were computed. Those statistics were then interpolated in each cell of the background grid. As the horizontal resolution at the coast of the ocean circulation model ranged from 80 m to 350 m and water depth was thresholded to 2 m (Briton *et al.* 2018), the area where ocean circulation was computed was not perfectly overlapping the background grid depicting the rocky habitat near the coast. In this shadow area, ocean variables are undefined and interpolating values on the background grid thus required to define temperature and flow speed values at the coastline. Flow speed values were set to 0 at the coastline, meaning total dissipation of flow energy at the coast. Sea bottom temperature values were set to their average along the 2 m isobath of the ocean circulation model. Inter-annual bottom flow speed and temperature minimum (Umin and Tmin) and maximum (Umax and Tmax) at each cell of the background grid were then calculated, taking the minimum and maximum values of monthly tenth and ninetieth percentiles, respectively, along the three years. Among those environmental predictors, uncorrelated ones were selected in order to avoid any identification confusion in Maxent due to their redundancies. Correlations between all possible pairs of predictors were calculated across the background grid range using Spearman's determination coefficient (Fig. S3, Supplementary Material), together with Principal Components Analysis (Fig. S4, Supplementary Material). When the determination coefficient of a pair of predictors exceeded 0.7, and their first and second components on the PCA plot were close, one of the two predictors was discarded (Sillero *et al.* 2021, Fig. S4). Among the 13 potential predictors, 10 were not correlated: five geomorphological predictors (Depth, Slope, Rugosity, Eastness and Northness) and five hydrological predictors (Umax, annual average SST, Tmin, Tmax and annual average Turbidity). For each of these 10 predictors, their statistical distribution was assessed across the five species' presence location sets, across the entire background grid and on restrictions of the background grid to each of the eight sites to evidence potential sampling bias. All data processing was performed using in-house © Matlab routines.

Diagnosics of model fitting

Model best fit was the average of five replicated probability maps fitted using all predictors under the cross-validate option. For each replicated fit, the presence data was partitioned randomly (random seed option) into a training dataset used to fit the model (75 % of presence data) and a testing dataset (25 % of presence data). AUC on the training and testing datasets and the relative contribution of each predictor to the replicated fit were calculated to assess the quality of the fit. The contribution of each predictor in defining each of the five replicated fits was ranked using a jackknife cross-validation procedure (Phillips *et al.* 2006). To do so, for each of the five presence data partitions, replicated fits were constructed using either a predictor on its own or all predictors except the one in question. Gains and losses of AUC compared to the

AUC with all predictors were calculated with and without each predictor, respectively.

Another ranking of the contribution of each predictor to a model was performed using a permutation procedure. For each predictor in turn, the values of that predictor in the training presence and background data were randomly permuted. The model was then re-evaluated on the permuted data, and the resulting drop in training AUC indicated that the model depended heavily on that predictor. Values were normalized to give percentages.

Finally, predictor response curves showed how the predicted probability of presence varied with the environmental predictor holding all other environmental predictors at their mean sample value.

Sensitivity of model fitting to potential sampling bias when reducing presence number and their geographical extent

Sampling bias can appear when some environmental conditions are preferentially sampled without respecting the proportion of their availability (Merow *et al.* 2013). Sampling bias can be the result of geographical or temporal preference in presence exploration or limited presence exploration. Such bias is evident for rare species whose niche is limited in space and requires intensive exploration to be observed. For more widely distributed species, sampling bias is less evident. Among the five gorgonian species, we selected the two species with highest occurrence: *E. singularis* (255 presence data) and *L. sarmentosa* (62 presence data), and assessed the sensitivity of the AUC when decreasing the number of presence data in the training dataset (while still keeping the full presence dataset for testing). To do so, we followed two procedures. In the first procedure, the model was fitted using three training datasets, gradually reducing the number of presence data to an evenly distributed selection of 50 %, 20 % and 10 % of the initial training dataset. In the second procedure, the model was fitted using a reduced number of presence data in the training datasets, corresponding to geographical restrictions to six sites for *E. singularis* (CVM, AGD, ARES, AGM, PMCB and PNPC) and two sites for *L. sarmentosa* (CVM and AGD). For both species, a training dataset including only presence recorded within MPAs was tested as well. Finally, sampling bias was also tested by testing if adding the geographical coordinates as predictors altered the model fitting.

Model performance assessment

The performance of two models, one using the full presence dataset and the other only using the presence observed in MPAs, was compared. Model performance was assessed by contrasting the habitat suitability predictions with observed presences and absences for each of the five species (the 444 grid cells subset mentioned above). For each species, regional habitat suitability maps were built after binarizing the mean of the five replicated probability maps. This was done by applying a threshold probability above which the habitat was considered as suitable and below which it was considered as unsuitable. Due to varying prevalence among the five species, the threshold probability

was not set to a fixed value but instead was determined using one of the 11 possible thresholding procedures advised in Maxent (Phillips *et al.* 2017). Deciding which threshold to use is a major consideration when minimizing false positives and false negatives. Its choice, however, is often arbitrary in studies (Liu *et al.* 2005). Among these threshold procedures, the 10percentile_training threshold (maximum error rate of 10 % on the presence points of the learning data) was arbitrarily selected because it displayed the highest threshold values when compared to other thresholding procedures, resulting in the less extended and hence more conservative habitat suitability maps (Fig. S5, Supplementary Material).

RESULTS

Species presence observations and sampling bias assessment

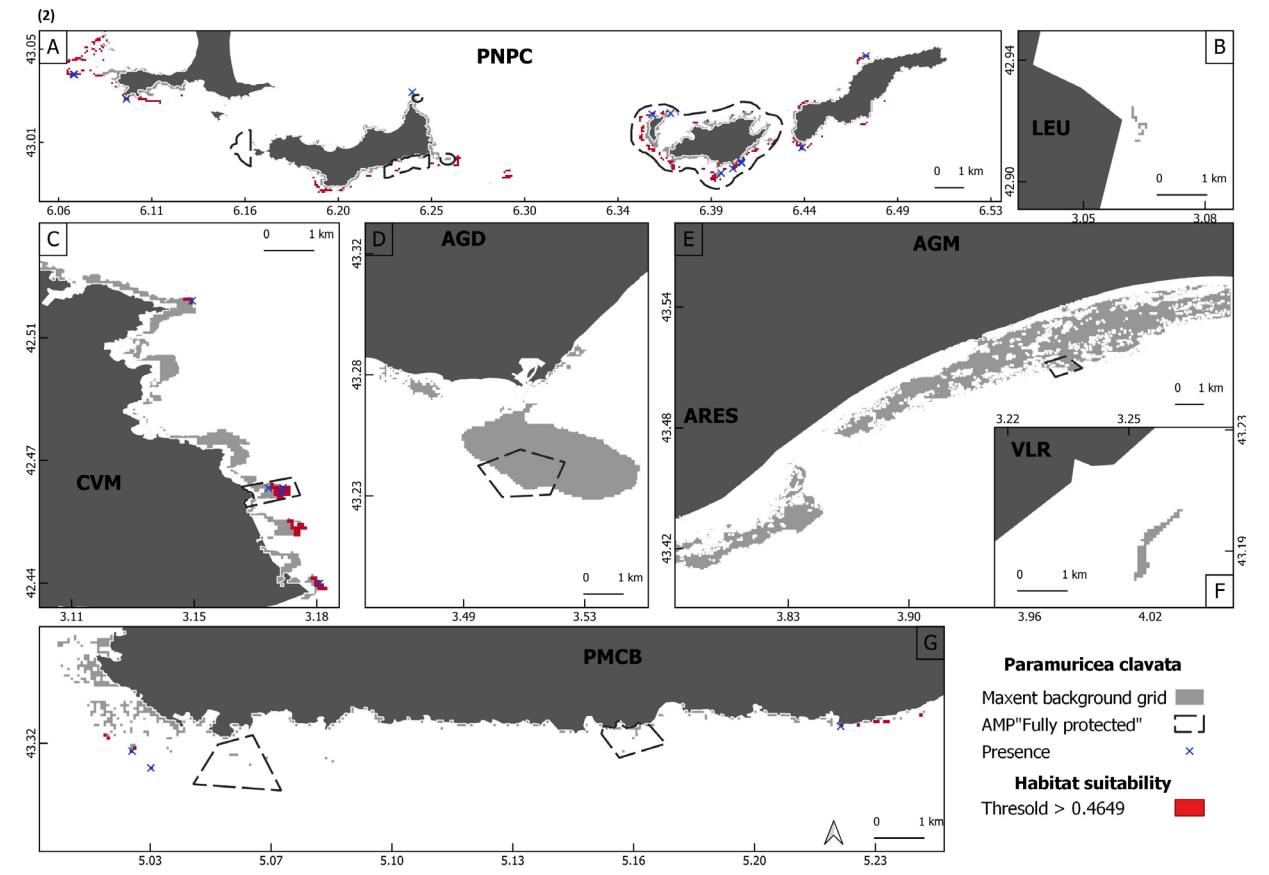
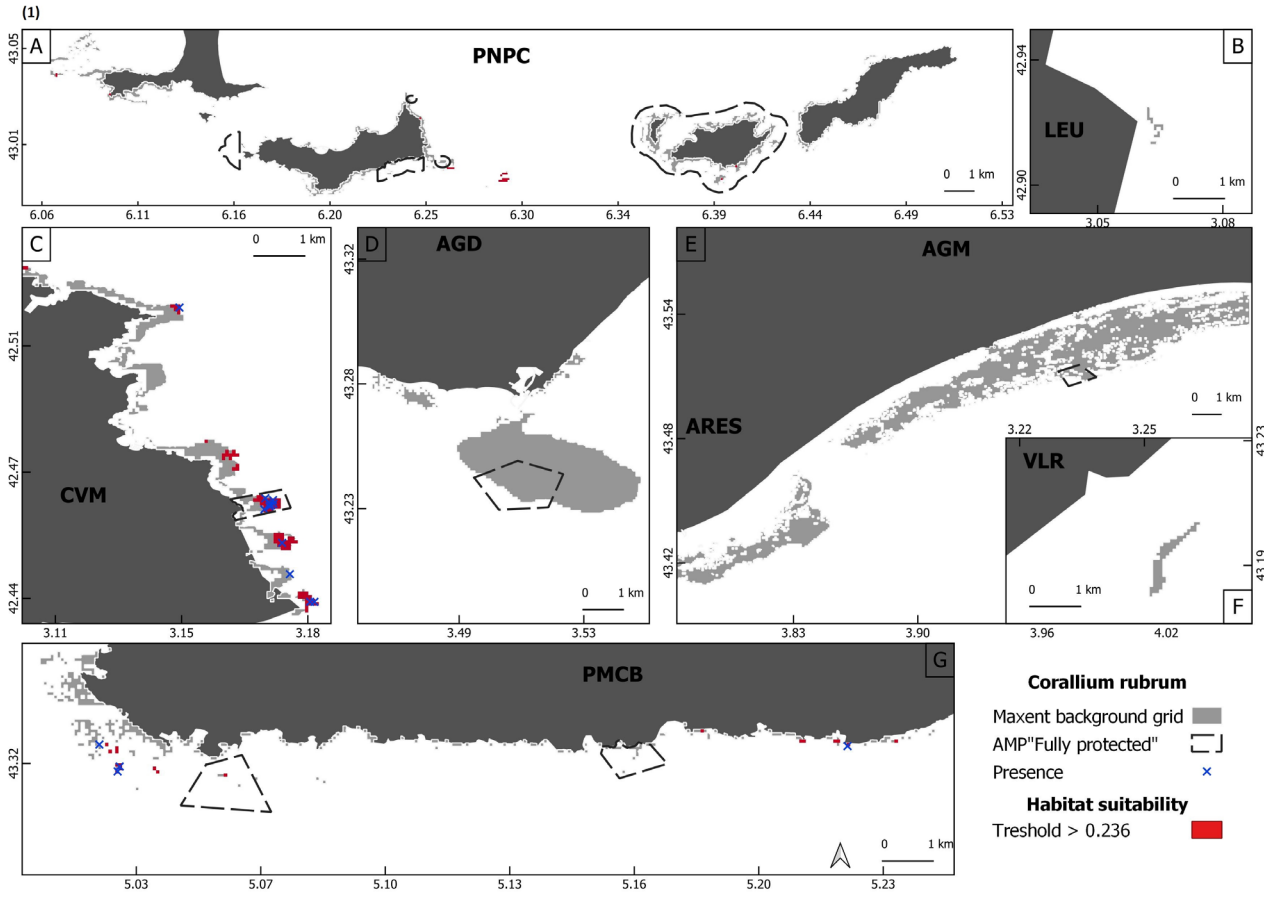
In the study area, the observed spatial distribution varied among the five species. Some species were distributed in a limited number of clustered locations: *C. rubrum* with 39 presences in only two sites (CVM and PMBC, Fig. 2), *P. clavata* with 61 presences in three sites (CVM, PMBC and PNPC, Fig. 2) and *E. cavolini* with 87 presences distributed in three sites (PNPC, PMCB and ARES, Fig. 2). *Leptogorgia sarmentosa* was detected everywhere except in PNPC and AGM (Fig. 2). Finally, *E. singularis* was the most widespread species with 479 presences, distributed in all eight sites (Fig. 2). The species most frequently associated together were *E. singularis* and *L. sarmentosa* (17.9 %). Boxplots that represented the environmental predictors range, showed marked differences among the eight sites forming the background seascape (Fig. 3). Depth range was doubled in four out of the eight sites, extending down to 40 m depth (CVM, AGD, PMCB and PNPC, Fig. 3D). Larger slope (Fig. 3E) and rugosity (Fig. 3F) were found in two (PNPC and CVM) out of the eight sites, while seabed orientation (eastness and northness) displayed a similar range of variability in all eight sites (Fig. 3G and H). Higher annual average SST (Fig. 3B) characterized one of the eight sites (PNPC) while Tmin at the seabed (Fig. 3J) clearly separated two sites (PMCB and PNPC) where temperature at the seabed did not fall below 11 °C from the other six sites where minimum temperature was below 10 °C. Lower annual average turbidity (Fig. 3C) characterised three out of the eight sites (CVM, PMCB and PNPC). Finally, larger Umax at the

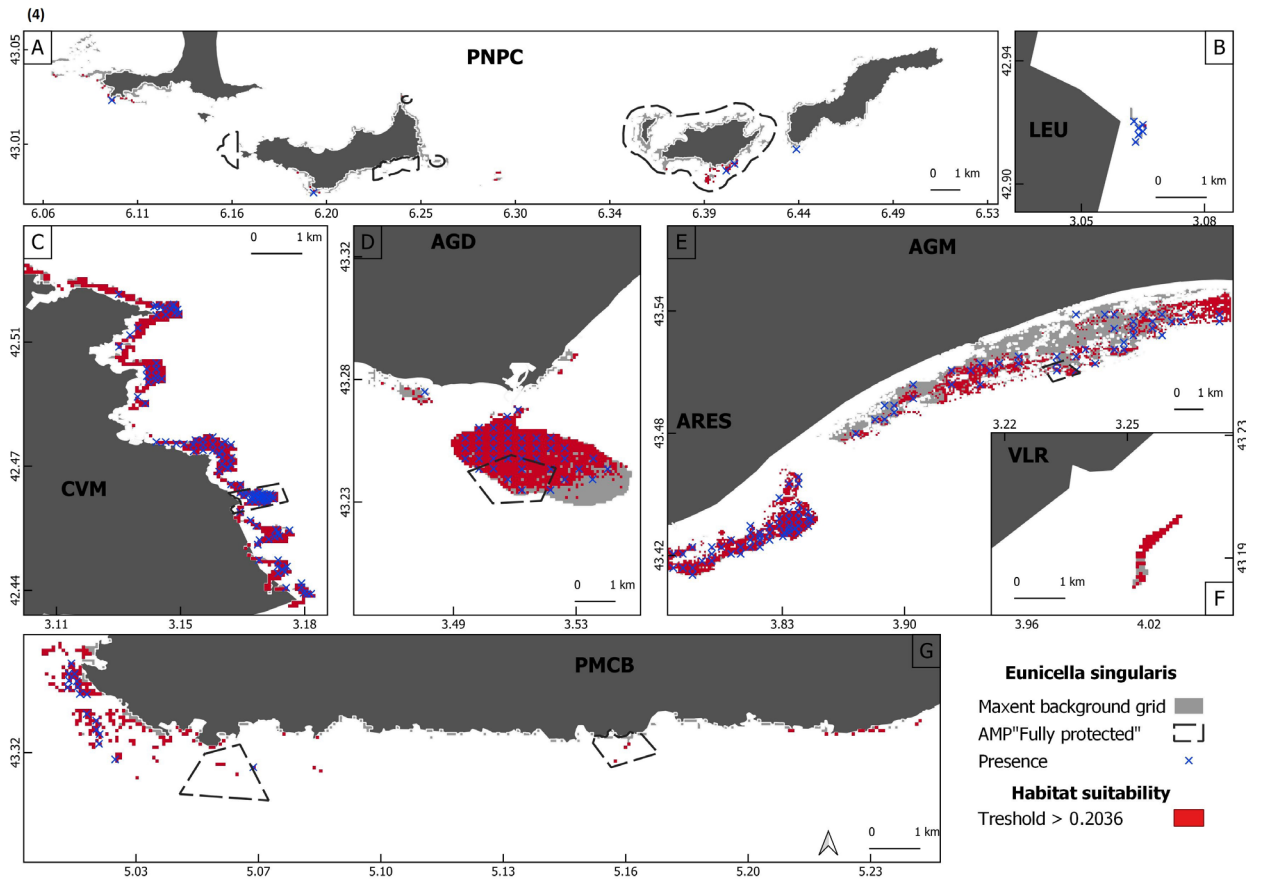
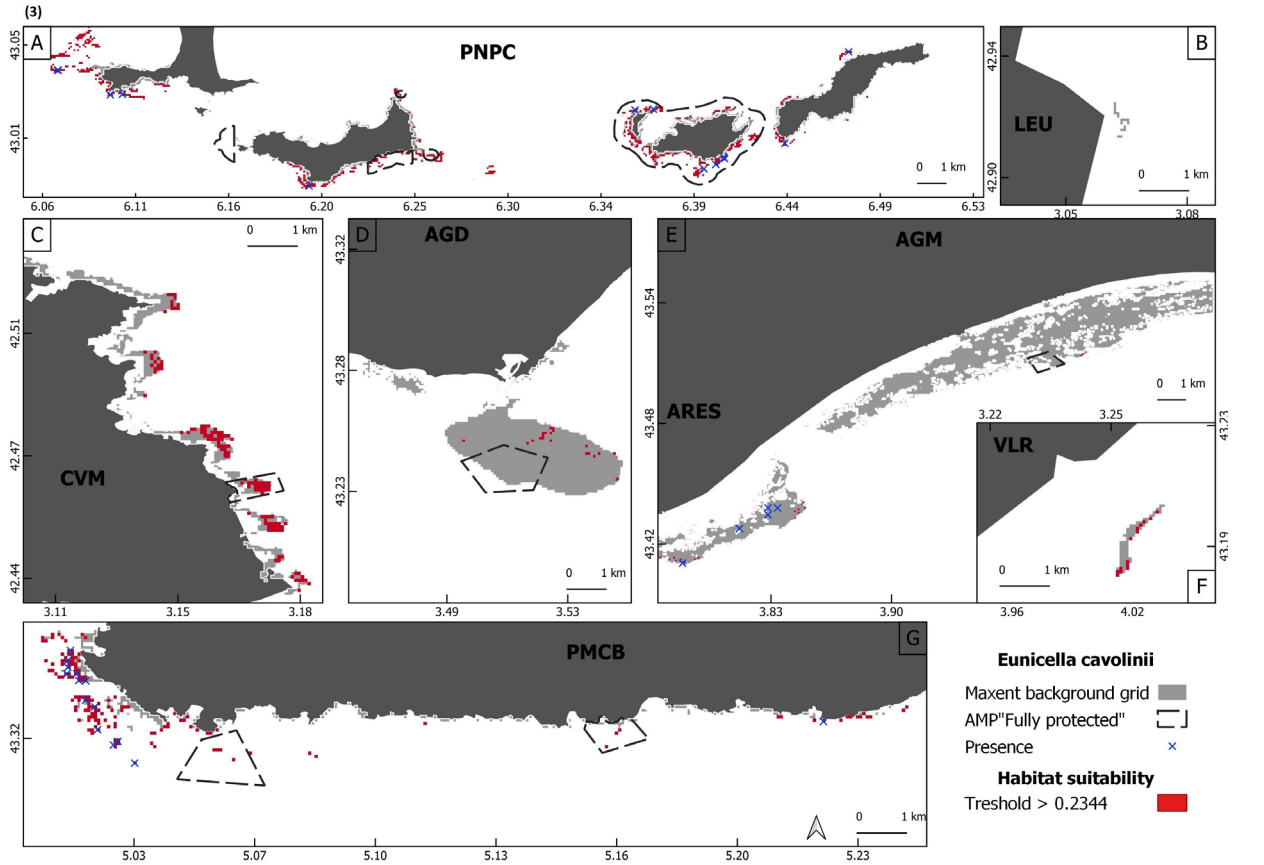
seabed was frequently found in five out of the eight sites (LEU, VLR, AGD, ARES, AGM, Fig. 3A).

Boxplots that represented environmental predictors range at species presence locations, indicated some differences between species as well. Umax at the seabed was consistently higher at presence locations of *C. rubrum* than of any other species (Fig. 3A). Annual average SST was consistently lower at presence locations of *C. rubrum*, *E. singularis* and *L. sarmentosa* than of *P. clavata* and *E. cavolini* (Fig. 3B). Annual average turbidity was higher at presence locations of *E. singularis* and *L. sarmentosa* (Fig. 3C), while slope was slightly higher at presence locations of *P. clavata* than at other species presence locations (Fig. 3E).

Environmental predictors contribution to habitat suitability prediction

Different environmental predictors contributed to the Maxent model best fit of each gorgonian species in the Gulf of Lion (when only using the presence locations found in the background grid, Fig. 2). For *C. rubrum*, annual average turbidity and Umax at the seabed contributed more than 90 % to the suitable habitat prediction (both in percent contribution and permutation percentage, Table I). Response curves for these two predictors indicated species preference for current speeds greater than 0.25 m.s-1 and annual average turbidity lower than 449 m-1 (Fig. S7, Supplementary Material), suggesting that the ecological niche of *C. rubrum* extends beyond the variation range of those predictors within the background grid of the study. Annual average turbidity and Umax were the predictor that, when absent, most decreased the AUC in the model, and Umax at the seabed contributed the most to the AUC in stand-alone mode, followed by depth and rugosity (Jackknife's test, Fig. S6, Supplementary Material). For *P. clavata*, Tmin at the seabed, depth, and Umax combined contributed more than 70 % of the prediction of suitable habitat (percentage contribution, Table I). However, Tmin at the seabed contributed only 6.3 % in permutation percentage (Table I), reflecting the similarity in the range of values of values of Tmin in the background and in





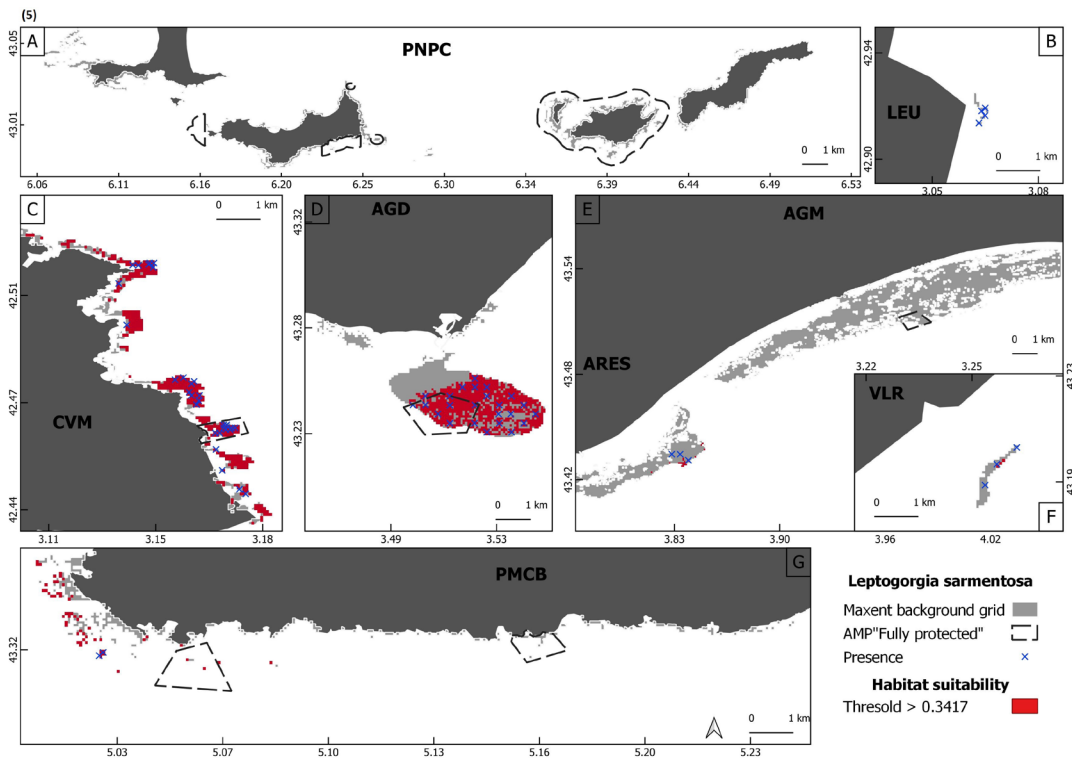


Fig. 2.- Presence location (Blue cross) and regional habitat suitability maps predicted using Maxent with all predictors except geographical ones (Latitude / Longitude) (Grid red). The habitat suitability map was calculated by applying a threshold value to obtain binary maps (0-1) from Cloglog output. A: close up PNPC, B: close up LEU, C: close up CVM, D: close up AGD, E: close up AGM and ARES, F: close up VLR, G: close up PMCB. Maps for : (1) *C. rubrum*, (2) *P. clavata*, (3) *E. cavolini*, (4) *E. singularis*, (5) *L. sarmentosa*.

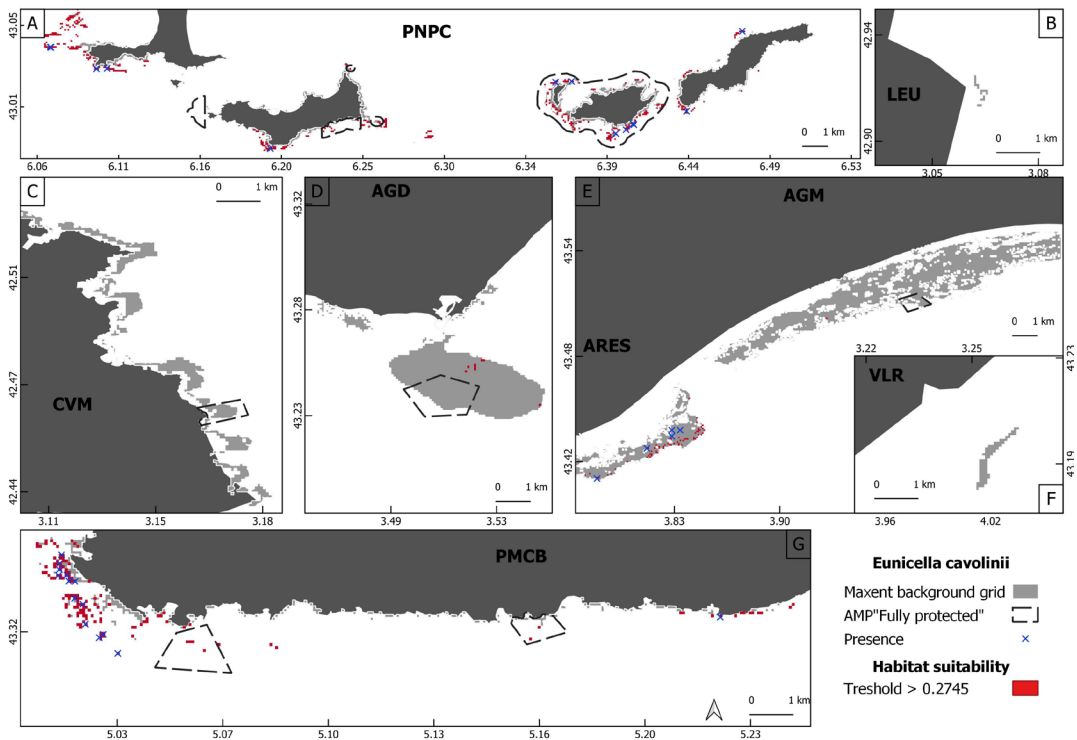


Fig. 2a: Presence location (Blue cross) and regional habitat suitability maps predicted using Maxent for *E. cavolini* with all predictors including geographical ones (Latitude / Longitude) (Grid red). The habitat suitability map was calculated by applying a threshold to obtain binary maps (0-1) from Cloglog output. A: close up PNPC, B: close up LEU, C: close up CVM, D: close up AGD, E: close up AGM and ARES, F: close up VLR, G: close up PMCB. Maps for : (1) *C. rubrum*, (2) *P. clavata*, (3) *E. cavolini*, (4) *E. singularis*, (5) *L. sarmentosa*.

	Variables	Percent contribution (%)	Permutation importance (%)
<i>C. rubrum</i>	Umax	70.5	43.3
	Annual average turbidity	22.9	49.8
<i>P. clavata</i>	Tmin	31.1	6.3
	Depth	26.1	35.7
	Umax	14.6	18.8
	Annual average turbidity	6.2	19.5
<i>E. cavolini</i>	Tmin	32.5	45.2
	Rugosity	30.3	33
	Depth	20.3	13.4
<i>E. singularis</i>	Rugosity	25.9	13.6
	Annual average SST	22.3	24.6
	Depth	14.6	26.9
	Annual average turbidity	13.7	12.8
<i>L. sarmentosa</i>	Depth	37	6
	Rugosity	21.6	20.2
	Annual average SST	10.7	10.3
	Annual average turbidity	10.7	17.8
	Tmin	5.2	18.9
	Tmax	4.2	20.2

Table I.- Significant environmental predictors identified for each species by Maxent without geographical predictors (Latitude / Longitude). Percent contribution and permutation importance

P. clavata presence sites (Fig. 3B). Conversely, annual average turbidity was ranked fourth in the percentage of contribution but second in percentage of permutation (19.5 %, Table I), reflecting the narrow range of annual average turbidity in *P. clavata* presence sites compared to the annual average turbidity range in the background (Fig. 3C).

Response curves of the four predictors indicated species preference for Tmin larger than 8.2 °C, water depth greater than 24 m, current greater than 0.14 m.s-1, and annual average turbidity less than 357 m-1 (Fig. S8, Supplementary Material). All response curves displayed a plateau when they reached high values, suggesting the

species can distribute beyond 50 m water depth and into stronger currents. Depth was the predictor that most decreased the AUC when it was not included in the model (Jackknife's test, Fig. S6).

For *E. singularis*, rugosity, annual average SST, depth contributed similarly to the suitable habitat prediction (both in percent contribution and permutation percentage, Table I), summing to 75 %. Response curves indicated that *E. singularis* had a preference for annual average SST ranging from 15.7 and 17.5 °C, water depth ranging from 4 to 40 m, with a characteristic niche-like optimal range of values and rugosity larger than 1 (Fig. S10, Supplementary Material). Annual average SST was the predictor that most decreased the AUC when removed from the model and contributed the most to the AUC in stand-alone mode (Jackknife's test, Fig. S6).

For *L. sarmentosa*, depth, rugosity, annual average SST and turbidity contributed the most to the suitable habitat prediction, summing up to 80 % in percent contribution. However, in permutation percentage, Tmin and Tmax at the seabed replaced annual average SST and depth, whereas together with rugosity and annual average, turbidity contributed to 77 % (Table I). Response curves indicated that a preferential habitat for *L. sarmentosa* would be in areas with water depth greater than 15 m, a temperature below 22.8 °C, in terrain with rugosity between 1 and 5 and with an annual average turbidity of between 204 and 600 m⁻¹ (Fig. S11, Supplementary Material). No predictor had a significant influence on AUC when they were absent from the model. Annual average SST, depth and Tmax contributed the most to stand-alone model AUC (Jackknife's test, Fig. S6).

In summary, annual average turbidity was identified as a significant environmental predictor for four species. Maximum flow speed at the seabed was the primary predictor for two species out of five. Among geomorphological predictors, rugosity and depth contributed to suitable habitat modelling, while no orientation predictors did. Despite rugosity being a significant predictor for all species, ru-

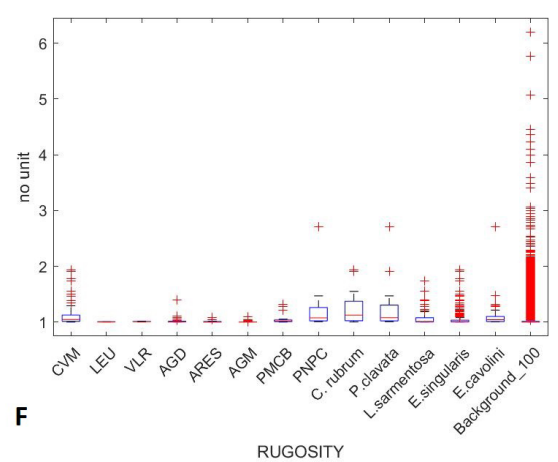
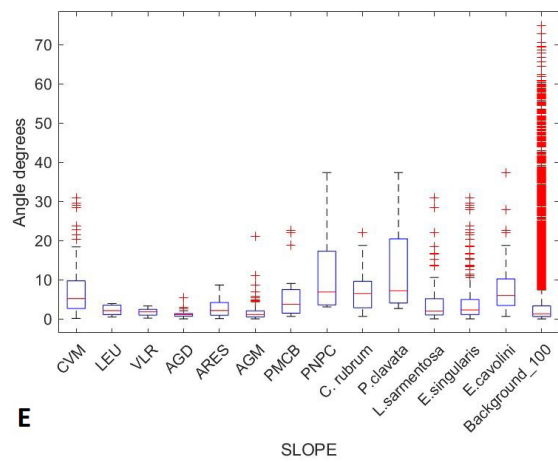
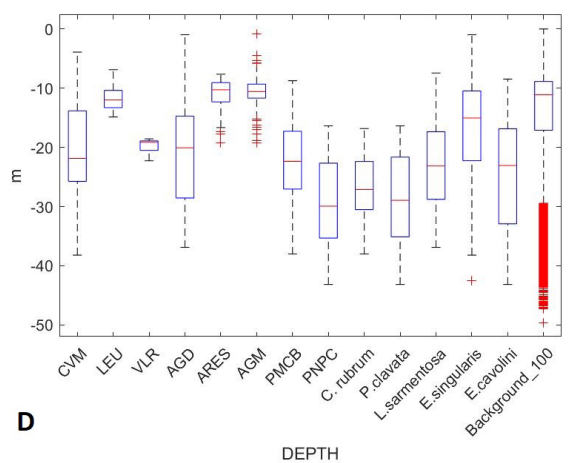
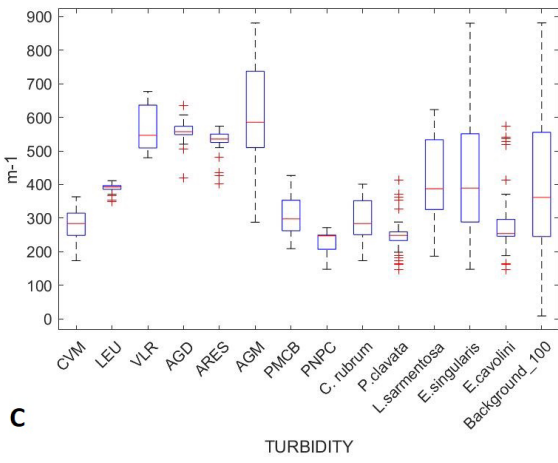
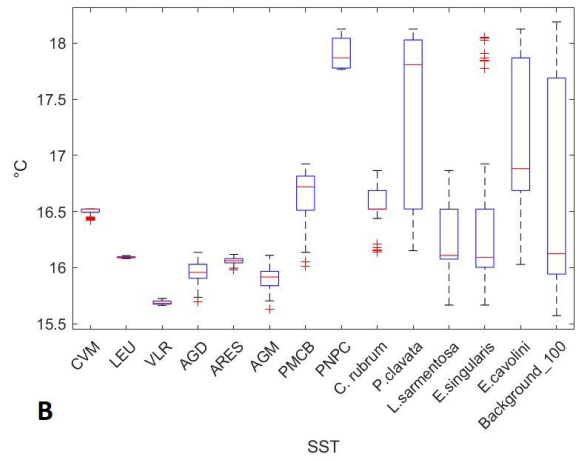
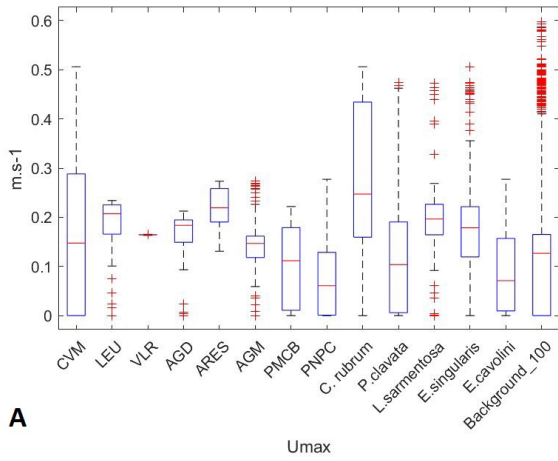
gosity response curves did not indicate much discrimination between the species, as any rugosity larger than 1 was favourable. Minimum flow speed at the seabed did not contribute to the suitable habitat modelling of any of the five gorgonian species.

Model performance and assessment of the predictions

Average AUC varied from 0.805 to 0.985 (less than 5 % standard deviation) across the five species, indicating good to very good fit when learning and testing against replicated random partitions of the presence dataset (Table II). Maps of suitable habitats that were obtained after applying a thresholding occurrence probability to the five replicated models were highly congruent, with less than 10 % of locations where the five replicated model predictions disagreed for all species except *E. singularis* (20 %). Maps of suitable habitats derived from the model best fit demonstrated correct predictions when compared to both presence and absence observations for all species (Fig. 2). However, model accuracy was lower for *L. sarmentosa* and *E. singularis* (70 % and 73 %, respectively) compared to *P. clavata* and *C. rubrum* (92 % for both, Table II). In all species, TSS differed significantly from a random contingency table (Chi-square, $P < 0.0001$), with values larger than 0.8 for *C. rubrum* and *P. clavata* and larger than 0.4 for *L. sarmentosa* and *E. cavolini* (Table II). For *E. singularis*, TSS was only 0.33 due to low specificity (43 % of absence correctly predicted, Table II), reflecting the large extension of the suitable habitat predicted for the species.

Assessing the influence of the number and location of presence observations used for model training on model fit goodness

Sensitivity of model AUC to the number and location of presence observations used for model training was carried out on *E. singularis* and *L. sarmentosa* because these species have the widest spatial coverage and the greatest number of presences. For *E. singularis* and *L. sarmentosa*, model AUC remained within the range of variation of the five replicated model AUCs when the training dataset was reduced by 90 % and 80 %, respectively,



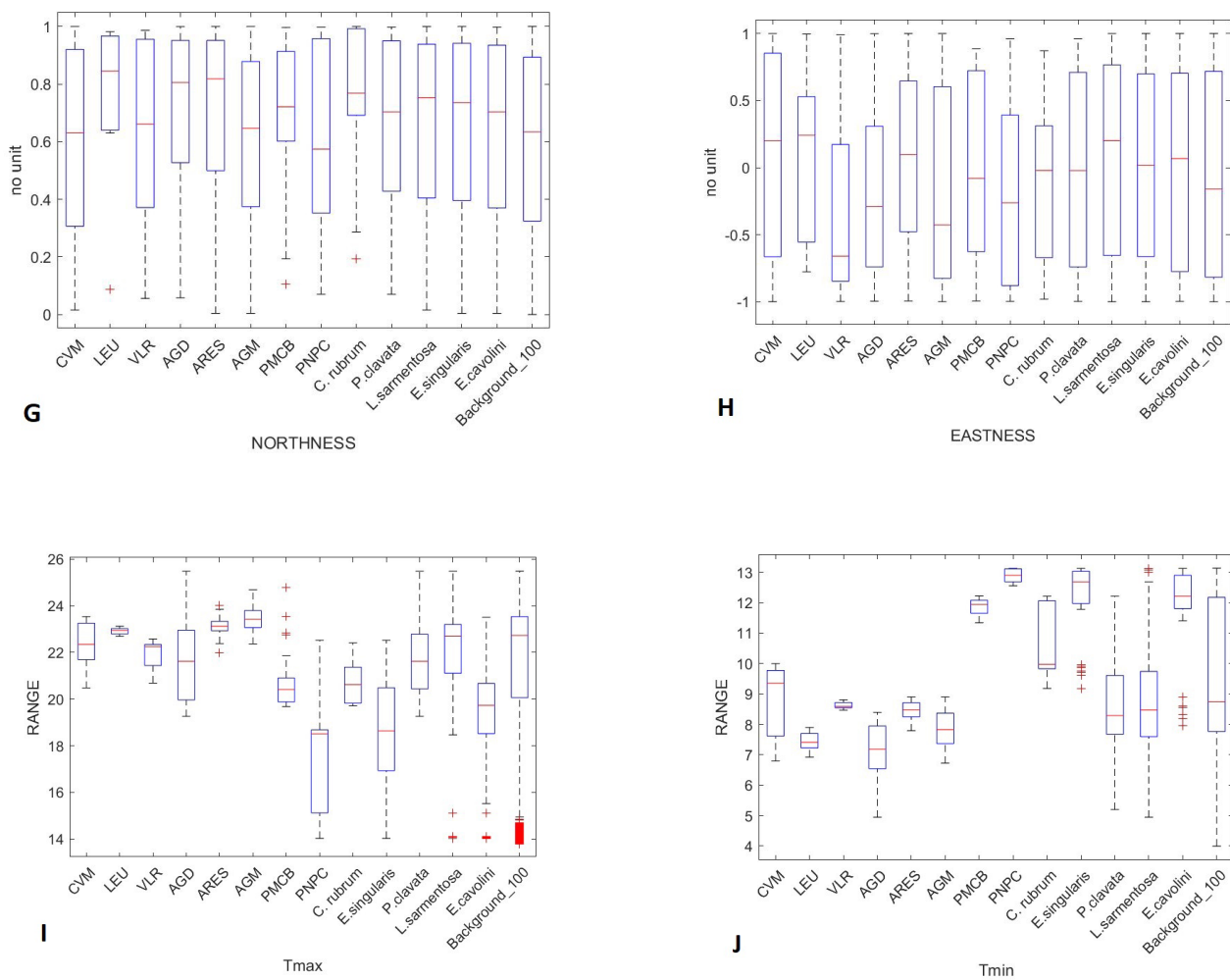


Fig. 3.- Boxplot of the 10 non-correlated predictors (except latitude and longitude) in the 8 sites (CVM, LEU, VLR, AGD, ARES, AGM, PMCB, PNP), at species presence locations and in the 100 m x 100 m background grid (A: Umax, B: Annual average SST, C: Annual average turbidity, D: Depth, E: Slope, F: Rugosity, G: Northness, H: Eastness, I: Tmax, J: Tmin).

as long as the presence observations of the training dataset were spread in all eight sites (Fig. 4). In contrast, model AUC decreased sharply when the number of presence observations was reduced by the same amount but restricted to a single site only (Fig. 4). When the number of presence observations was restricted to the highly protected zones of the area, model AUC decreased to 0.7 for *E. singularis* and 0.85 for *L. sarmentosa*. This decrease was, however, less than if the same number of presences had been taken from a single site, although slightly more than if it was evenly distributed in all eight sites. When the number of presence observations was restricted to the highly protected zones of the area, model AUC decreased only for *P. clavata* and

L. sarmentosa (0.93 and 0.88 respectively, Table II) while accuracy decreased for *P. clavata* and *E. singularis* (0.89 and 0.59 respectively, Table II). Adding latitude and longitude as environmental predictors, the mean test AUC was unchanged for *E. singularis*, *P. clavata* and *C. rubrum*, increased for *E. cavolini* (0.955) and decreased for *L. sarmentosa* (0.890). For *L. sarmentosa*, adding geographical predictors altered predicted distributions, resulting in the loss of a site where the species has been observed (ARES, Fig. S12, Supplementary Material). In contrast, the suitable habitat map for *E. cavolini* reduced considerably. Suitable habitat area was removed in CVM and reduced in AGD and AGM with improvement of accuracy

	AUC test	Standard deviation of AUC test	Accuracy	Sensitivity	Specificity	TSS
<i>C. rubrum</i>	0.985(0.989)	0.005(0.019)	0.92(0.94)	0.90(0.65)	0.92(0.95)	0.82(0.60)
<i>P. clavata</i>	0.974(0.936)	0.007(0.023)	0.92(0.89)	0.88(0.88)	0.92(0.89)	0.81(0.77)
<i>E. cavolini</i>	0.932(0.965)	0.045(0.037)	0.82(0.94)	0.90(0.10)	0.81(1)	0.71(0.1)
<i>E. singularis</i>	0.805(0.932)	0.034(0.034)	0.73(0.59)	0.91(0.44)	0.43(0.85)	0.33(0.29)
<i>L. sarmentosa</i>	0.911(0.886)	0.027(0.073)	0.70(0.78)	0.77(0.57)	0.68(0.83)	0.45(0.40)

Table II.- Assessment of the performance of model and predicted suitability maps trained on 80 % of presence data with all predictors except geographical ones (Lat / Lon) : AUC test average and Standard deviation of 5 models, Accuracy, Sensitivity: true positif, Specificity: true absence. Values in *Italic* between parentheses corresponds to model trained on sampling in AMPs only.

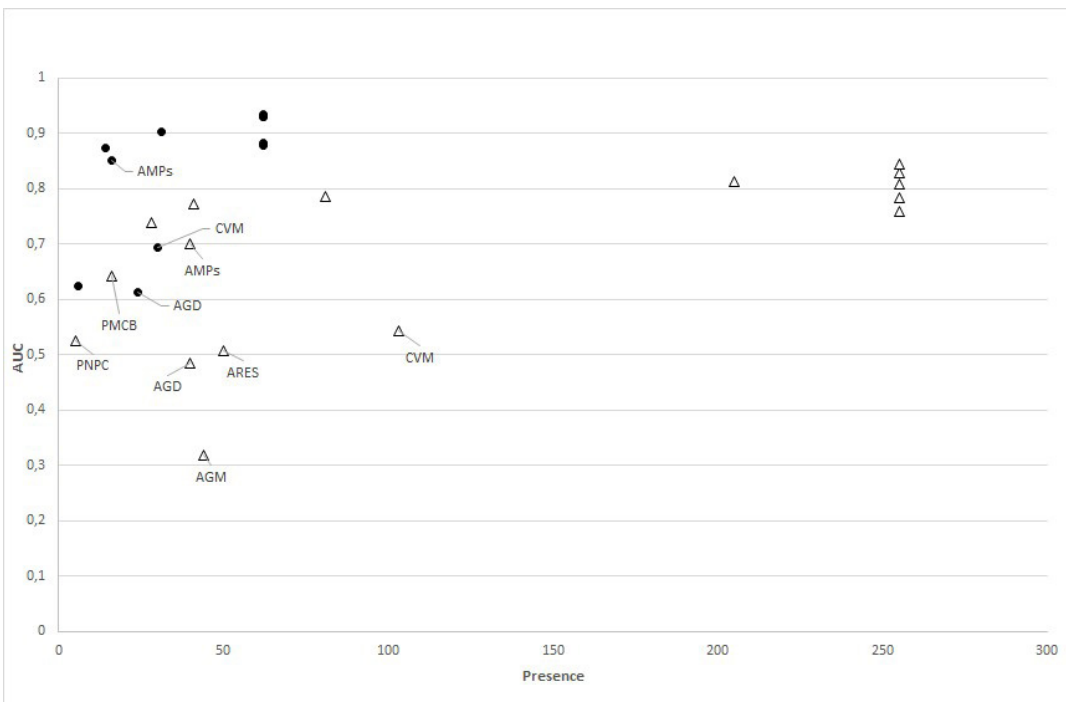


Fig. 4.-. Assessment of the AUC for *Eunicella singularis* (white triangle) and *Leptogorgia sarmentosa* (dark cercle) varying the number of presence data (down to 7 % of the initial training dataset) and their location in the training dataset while keeping the full presence data in the testing dataset.

cy and TSS (0.96 and 0.86 respectively, Fig. 2a)

DISCUSSION

The observations between 10 and 50 m depth across eight rocky sites spanning 450 km along the Gulf of Lion (NW Mediterranean Sea) confirmed that between 2013 and 2020 the hierarchy in the occurrence of the five gorgonian species was similar to the one previously reported

in the same area (Weinberg 1979a, Linares *et al.* 2008, Gori *et al.* 2011a). *Eunicella singularis* was the most abundant gorgonian species with a wide continuous spatial distribution. *Leptogorgia sarmentosa* was the second most abundant gorgonian species, with scattered distribution. *Paramuricea clavata* and *Corallium rubrum* displayed highly patchy wwdistributions and their presence was observed in three rocky sites only. *Eunicella cavoli-*

ni was the least abundant species, with marginal presence in a few spots in three rocky sites. Occasional associations between the five gorgonian species were observed but were not frequent, similar to previously reported between 0 and 70 m deep south of the Gulf of Lion (Gori *et al.* 2011a).

Depth is a primary factor that has been used as a basis for the classification of marine benthic communities strata in the shallow environment as both light availability and agitation at sea decrease with depth (Perez 1961, Odum 1971). In the present study, depth was identified as a significant predictor for all species distribution except *C. rubrum*, but with different responses according to the species. For *P. clavata*, model predictions corroborated a clear preference for deeper habitats as already mentioned in the literature (Rossi 1959, Boavida *et al.* 2016, Fourt *et al.* 2017). Similarly, model predictions for *E. singularis* which is the only one of the five species hosting *Symbiodiniaceae* (Forcioli *et al.* 2011) indicated a preference for shallow areas which might be related to more abundant light (between 4 and 32 m). Nevertheless, a different morphotype of the species with extremely low densities of *Symbiodiniaceae* has been reported beyond 50 m depth (Theodor 1969, Gori *et al.* 2011b). Moreover, limited gene flow between the shallow and deep morphotypes was reported at local scale (Costantini *et al.* 2016). These observations suggest that care should be taken when delineating habitat suitability for this species, as different morphotypes may likely display different niches, hence related to different environmental predictors. For *L. sarmentosa*, depth response excluded depth less than 15 meters while the species has been observed in shallow harbour areas (Betti *et al.* 2018). For *E. cavolini*, depth response only excluded very shallow environments (shallower than 5 m) in the present study area while in the locality of Kornati (Adriatic Sea), an *E. cavolini* population was found at 5 m depth (Sini *et al.* 2015). For *C. rubrum*, depth was not identified as a significant predictor, despite the fact that the species preference for low light has been largely reported (Laborel 1960, Weinberg 1979b, Bianconi *et al.* 1988). The presence of *C. rubrum*

at shallow depth might, instead, be related to the smaller size of this species, compared to the other four gorgonian species, which would allow the colonization of small dark crevices (Laborel 1961, Rossi *et al.* 2008). In any case, despite the fact that depth is easier to measure than light, direct measurement or modelling of light intensity should be undertaken to improve the delineation of the species niche (Betti *et al.* 2018).

Seabed orientation is another proxy for light availability that has been used in several studies at small spatial scales (Glasby 2000, Knott *et al.* 2004). However, in the present study, orientation was never indicated as a significant predictor. This was, most probably, due to the fact that orientation and light availability are not correlated over a large spatial scale.

Beyond depth, the five gorgonian species have been associated with different preferences in geomorphological features. *C. rubrum* is mainly found in overhangs or in caves (Cau *et al.* 2016). *P. clavata* and *E. cavolini* are reported to dwell in steep rocky coralligenous bottoms, such as drop-offs, and they are often found in association (Carugati *et al.* 2021). *L. sarmentosa* have been associated to the detritic-muddy benthic community, characterized by boulders and abundance of organic matter (Carpine & Grasshoff 1975, Mistri & Ceccherelli 1993). *E. singularis* has been mainly reported on well-lit horizontal and sub-horizontal rocky bottoms (Gori *et al.* 2011a, Weinberg & Weinberg 1979). The latter two species have also been reported on gently sloping rocky bottoms (Gori *et al.* 2011a). High-resolution terrain mapping thanks to airborne and satellite remote sensing has been an essential data source for determining species habitats when using predictive habitat modelling in terrestrial environments (Wilson *et al.* 2007). However, high-resolution seabed mapping over large areas is still rare. Vessel-mounted multibeam acoustic sounders can provide seabed maps with a vertical resolution at the cm scale and horizontal resolution at the tens of cm scale, but only over a small surface (Dolan *et al.* 2008). The cost of acquiring such data has limited its implementation, which is why predictive ha-

bitat modelling studies at sea have focused on specific environments such as canyons (Guinan *et al.* 2009, Bargain *et al.* 2018, Lo Iacono *et al.* 2018). The present study aimed to fill a gap in predictive habitat modelling for shallow marine environments by leveraging advances in bathymetry remote sensing using airborne laser imaging between 2011 and 2015 (LITTO3D, www.diffusion.shom.fr), which enabled mapping shallow areas along hundreds of kms of coastline, with submetric spatial resolution similar to that of vessel-mounted acoustic sounders. Yet, our choice to average geomorphological predictors over a 20 m x 20 m square slab to match the decametric scale at which the species presence was detected, no longer enabled to detect rapid terrain variations with a single variability predictor. This is a frequent dilemma when quantifying geomorphological variations (Wilson *et al.* 2007), and it explains why we kept two variability predictors (slope and rugosity), as in combination they have the potential to differentiate flat tilted bottom (low slope and large rugosity) from trenches (large slope and low rugosity) and walls (large slope and large rugosity). However, model predictions did not indicate slope as a significant environmental predictor for any of the five species, while steep slope is often assumed to favor benthic invertebrates colonisation, as the sediment deposition on flat surfaces could limit their recruitment (Laborel & Vacelet 1961, Bianconi *et al.* 1988, Glasby 2000). In particular, walls, which have often been described as the ecological niche of *P. clavata*, may not actually be (Cocito *et al.* 2002, Boavida *et al.* 2016, Ponti *et al.* 2019). This result is consistent with the presence of very big colonies of *P. clavata* on horizontal rocks in the deep environment (Grinyó *et al.* 2016). Reversely, rugosity differentiated the suitable habitat of *L. sarmentosa*, which is mainly found at a moderate rugosity, and that of other two species (*E. singularis* and *E. cavolini*) which can be found on a wider range of rugosity. Yet, the LIDAR resolution did not enable the detection of crevices smaller than 0.1 m in which *C. rubrum* colonies are often found (True 1970, Virgilio *et al.* 2006). This may explain why rugosity was not identified as a significant predictor for this latter species in the present study.

In addition to geomorphological predictors, hydrological predictors have been put forward as factors regulating the presence of gorgonians. Among these, turbidity is one of the trickiest ones. Sea water turbidity remotely sensed from satellites can indicate sedimentation due to river inputs or to resuspension by waves (Liew *et al.* 2009, Gohin *et al.* 2020). High sedimentation is considered a limiting factor for most gorgonians, potentially due to the burial and/or clogging of recently settled polyps, except for *L. sarmentosa* (Otero *et al.* 2017). The preference of *L. sarmentosa* for turbid seawater would be related to a morphological adaptation, with higher polyp density per branch and finer branches, which reduced surface would limit sediment accumulation in turbid environmental conditions, compared to the other four gorgonian species (Cocito *et al.* 2002, Rossi *et al.* 2011). In the present study, annual average turbidity was identified as a significant predictor for all species, except *E. cavolini*. As expected, *C. rubrum* and *P. clavata* were limited to a lower range of turbidity than *L. sarmentosa*, which displayed a well delimited range of intermediate turbidity preference (Weinberg 1979a, Mistrì 1995). For *E. singularis*, the response to turbidity was unclear with tolerance to both high and low annual average turbidity, which may reflect the presence of two morphotypes in the 10-50 m depth range (with high or low densities of Symbiodiniacea, Gori *et al.* 2012). Indeed, preference for turbid environments can be expected in heterotrophic suspension feeders and might be related to food requirements while preference for less turbid environments can be expected in autotrophic species relying on photosynthesis. Apart from the broad differentiation between autotrophy and heterotrophy dominance in *E. singularis*, available environmental predictors did not enable distinguishing between living (zooplankton) and non-living (detritic) food preferences of the other four heterotrophic gorgonian species. Anyhow, *P. clavata*, *L. sarmentosa* and *C. rubrum* display only subtle differences in the proportions of their trophic sources (Ribes *et al.* 1999, Ribes *et al.* 2003, Tsounis *et al.* 2006). Sedimentation can be locally reduced by large

flow speed values (Hiscock 1983), another factor suggested as structuring the distribution of gorgonians (Leversee 1976, Weinberg 1979a) and deep-water corals (White *et al.* 2005). The five studied gorgonian species, being passive suspension feeders, are expected to have a preference for moderate to strong flow speed values (Gori *et al.* 2011a, La Riviere *et al.* 2021). The present study identified large flow speed as a significant predictor for *C. rubrum* and *P. clavata* only. For *P. clavata*, maximum flow speed at the seabed was a more significant predictor than slope and roughness to describe the spatial distribution of the species, pointing to the need of adding hydrodynamical predictors to avoid confusion of factors when analysing the drivers of species distribution. Indeed, flow speed at the seabed is not commonly used as a predictor in ecological niche models (but see Bargain *et al.* 2018). It should be more systematically tested, taking care of using hydrodynamical simulations at adequate spatial resolution near the coast (Sciascia *et al.* 2022). Finally, temperature appeared as a significant environmental predictor for four of the five species, but never highlighting a limitation due to large maximum temperature at the seabed, while temperature positive anomalies during heat waves have been related to mass mortalities in *E. singularis* (Ezzat *et al.* 2013), *E. cavolini* (Turicchia *et al.* 2018) and *P. clavata* (Coma *et al.* 2006, Bally & Garrabou 2007, Garrabou *et al.* 2009). Although *L. sarmentosa* is found even inside ports, its tolerance to high water temperature (up to 29°C) was not evidenced either (Betti *et al.* 2018, personal observation). Actually, temperature predictors for *E. singularis* and *L. sarmentosa* (annual average SST < 17.5°C) and *P. clavata* and *E. cavolini* (Tmin > 8°C) reflect large-scale geographical separation between the colder Gulf of Lion sites and the warmer NPC and PMCB sites. In summary, identifying the environmental predictors for marine species still faces limitations and will require both increasing the number of presence observations to deeper areas and including more explicitly factors such as local temperature, light, turbidity, sedimentation, chlorophyll-a and flow speed. To this aim, the use of high-resolution flow

and primary production simulations would be an invaluable tool for ecological studies, as they allow projecting the distribution of the future suitable habitats under climate change scenarios and test the deep refugia hypothesis (Bongaerts *et al.* 2010).

Nevertheless, despite the above-mentioned limitations, Maxent performance was reasonable for *E. singularis* and very good for the other four species (according to AUC rating, Hanley & McNeil, 1982, Sainz-Villegas *et al.* 2022). Accuracy of predicted suitable habitat maps was good to excellent for all species (according to TSS rating, Araujo *et al.* 2011). The good sensitivity of the models for *E. singularis* and *L. sarmentosa* with few but regularly distributed observations of presence, confirms, on the one hand, that spatial autocorrelation was avoided and that large AUC values could not be attributed to model overfitting (Radosavljevic & Anderson 2013) and, on the other hand, that a low number of presence data is sufficient to obtain a good performance of the models as long as a wide geographical area is encompassed, to avoid local specificities in the training process. Moreover, in shallow environments, characterised by a diversity of environmental conditions and strong human impacts, we advocate for the use of highly protected zones (such as the Gulf of Lion MPAs) as sentinel sites, to allow for the monitoring of the factors shaping the benthic species distribution in coastal areas. Indeed, gorgonians like other benthic species are known to be sensitive to mechanical impacts such as mooring (Broad *et al.* 2020), bottom fishing (Bavestrello *et al.* 1997, Betti *et al.* 2020), scuba diving (Coma & Zabala 1994) and even direct exploitation (i.e., *C. rubrum* for jewellery, Tsounis *et al.* 2010, 2013). The small number of presence observations obtained from MPA species inventories, that were repeated over time, provided enough data to understand the habitat preferences of these species. In the present study, true absence and presence data used to assess model accuracy were defined based on repeated inventories obtained between 2013 and 2020 and avoided bias due to transient population accidents (*E. singularis* was observed in 2013 but was absent in 2019 in some locations). Given the

slow population dynamics of gorgonians, impacted populations may take several years to recover (Linares *et al.* 2007). The reiteration of sampling over a decade is thus essential, in order to assess the ENMs performance. In addition, an observation strategy that is focused on a limited number of locations would improve the measurement of predictors by equipping sentinel sites with adequate sensors (e.g., seawater temperature measurement networks such as T-MEDNet, Garrabou *et al.* 2019).

For all species, the suitable habitat prediction area was greater than the original inventory range. As explained above, suitable habitat is, by definition, expected to extend beyond actual presence observations and repeated observations in time are key to the fine tuning of true absence data and model accuracy. However, in the present study, *E. cavolini* displayed very reduced presence distributions compared with its predicted suitable habitat. Moreover, geographical predictors improved the prediction. A phylogenetic study suggested that differentiation between *E. singularis* and *E. cavolini* may have resulted from isolation in different refugia during the last glaciation (Aurelle *et al.* 2017). In such a hypothesis, the absence of *E. cavolini* from the Gulf of Lion (except one locality in ARES) could be attributed to a quicker re-colonisation of the shallow suitable habitat by *E. singularis* after the last glaciation, pointing to relatively recent geographical vicariance. Since then, the two species seem to have evolved towards diverging environmental niches as their predictors differ. However, beyond the definition of a suitable habitat based on environmental predictors (potential niche), actual species distribution (realized niche) ultimately depends on efficient colonisation or demographic connectivity, which is a complex process that operates at multiple spatial and temporal scales, resulting from species demography (population size, structure, fecundity) and larval dispersal (larval release timing, ocean currents, pelagic larval duration, larval behaviour). Scenarios using spatially explicit metapopulation modelling can help testing the respective role of the suitable habitat and demographic connectivity in explaining observed species spatial distribution (Guizien *et al.* 2014).

To conclude, habitat suitability models

differentiated the niche of *P. clavata*, *C. rubrum*, *E. singularis*, *E. cavolini* and *L. sarmentosa* using seven predictors (maximum flow speed at the bottom, annual average turbidity, depth, minimum and maximum temperature at the bottom, sea surface temperature and rugosity). The spatial distribution of the suitable habitat was predicted with excellent accuracy and sensitivity for the five species in the Gulf of Lion. These models can be used to complement the delineation of *P. clavata*, *C. rubrum*, *E. singularis*, *E. cavolini* and *L. sarmentosa* suitable habitat in the shallow rocky bottoms in other regions along the Mediterranean basin and guide marine spatial planning, including the establishment of new MPAs. However, observations in the mesophotic and aphotic environments should be added to the models developed here for the shallow rocky bottoms and should be updated to incorporate more direct predictors (i.e., light, sedimentation rate, organic matter content) in order to avoid the use of proxies such as depth and turbidity.

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Authors contributions

SB and KG conceived the study, SB and LB carried out sampling and SB did the statistical analysis. All contributed to

manuscript writing.

Conflict of interest disclosure

The authors of this preprint declare that they have no financial conflict of interest with the content of this article.

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SUPPLEMENTARY MATERIALS

ECOLOGICAL NICHE MODELLING OF FIVE GORGONIAN SPECIES WITHIN THE SHALLOW ROCKY HABITAT OF THE FRENCH MEDITERRANEAN COAST

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MATERIALS AND METHODS

Background grid, features and regularization

The background grid is a set of georeferenced locations where probability of species presence will be predicted (a priori with uniform species prevalence is defined) using known environmental predictors values. Probability distribution of environmental predictors in the background grid is compared to probability distribution of environmental predictors in presence locations in order to build response curves. Hence, Maxent background grid actually defines the range of environmental conditions on which features will be fitted in order to maximize the species entropy probability distribution, resulting in different features and Maxent predictions according to background extent (Phillips *et al.* 2006).

Various feature classes can be combined in Maxent to build response curves from environmental predictors. Theoretically, a niche model is expected to be unimodal, pointing at species optimum, which will be best represented by quadratic features. Sometimes the niche range, however, might be truncated in the presence observations. In this situation, linear or hinge features will be more appropriate (Merow *et al.* 2013). Multiplying feature class possibilities may lead to confusion because complex features created by Maxent are often highly correlated. From a statistical perspective, when Maxent is used to infer environmental drivers of species distribution, feature classes should be selected with parsimony. From a machine-learning perspective, testing as many feature classes as possible will increase the predictive accuracy of the presences. In this study, we adopted an in-between perspective, selecting the default «auto-features» mode of Maxent. This mode increases the tested feature classes to the number of presence observations with all feature classes for more than 80 presence observations (Phillips & Dudík 2008). This mode enabled Maxent to accommodate the different number of presence observations, which varied from 16 for *C. rubrum* to 255 for *E. singularis*. Regularisation dampens predictors coefficients when predictors variance is high at the presence locations. Regularisation multiplier was varied from 0.1 to 5 and the AUC of model fit varied by less than 1 %. Hence, it was set to its default value of 1.

Output type

The ROR is the relative probability that a cell is contained in a collection of presence samples, assuming individuals have been randomly sampled in the environmental space. A logistic transformation of ROR enables one to predict the probability of presence-only, assuming the the probability of presence in the suitable habitat, namely the prevalence, is 50 % (Phillips & Dudík 2008). Due to the difficulty of checking this assumption, the safer interpretation of Maxent predictions was using ROR as a habitat suitability index (Merow *et al.* 2013). As an alternative, the cloglog value calculated as $1 - \exp(-c \cdot r)$ has been proposed in order to estimate the probability of presence (c is the exponential of the entropy of the Maxent distribution and r is the ROR value, Phillips *et al.* 2006). In the present study, however, model fit was not sensitive to the species prevalence value in the background grid. The latter varied from 20 to 80 % and resulted in average AUC (Area Under the Curve) variation of 0.6 %. Therefore, prevalence was kept to its default value of 50 % and the cloglog output type was selected.

Definitions of the seven geomorphological predictors

All predictors were computed on the 20 m x 20 m slab located at the center of the 100 m x 100 m grid cell.

- Local bottom depth was the average of bathymetric data within the slab.
- A local digital terrain model (DTM) was defined by fitting all bathymetric data within the slab with a bivariate quadratic equation (1) (Evans 1980):

$$Z = aX^2 + bY^2 + cXY + dX + eY + f \quad (1)$$

with Z being the bottom depth deviation from the mean bottom depth and X,Y the cartesian distance to the center of the background grid cell and a, b, c, d, e and f are the fitting parameters that defines the local DTM and vary at each background grid cell. These parameters are subsequently used to calculate the slope, roughness, rugosity, ruggedness and orientation. • Slope was defined as the average slope of the local DTM:

$$S = \arctan(\sqrt{(d^2+e^2)})$$

- Roughness is length scale (in meter) defined as the difference between the maximum and minimum depth in the slab which is consistent with the hydraulic roughness parameter used to estimate seabed friction by Nikuradse (1933).
- Rugosity is a topological metric (no unit) defined as the ratio between the surface of the local DTM and the slab surface. By definition, rugosity is larger or equal to 1.
- Terrain ruggedness index was defined as the average of absolute deviation of the local DTM with respect to the mean depth.
- Eastness and northness orientation were defined as the sinus and cosinus of the DTM aspect, the latter being defined as:

$$\text{Aspect} = \text{atan}(d/e)$$

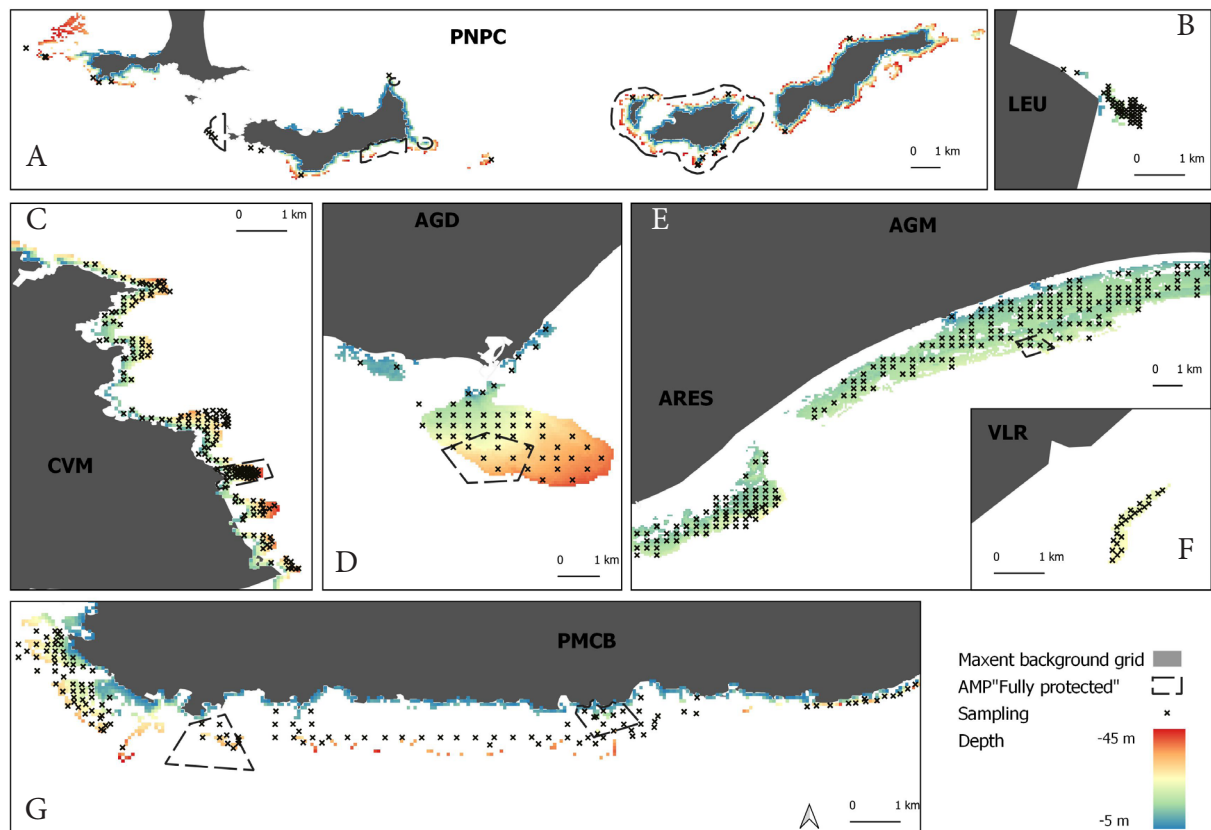


Fig. S1.- Sampling location and bathymetry in each of the eight sites. A: close up PNPC, B: close up LEU, C: close up CVM, D: close up AGD, E: close up AGM and ARES, F: close up VLR, G: close up PMCB.

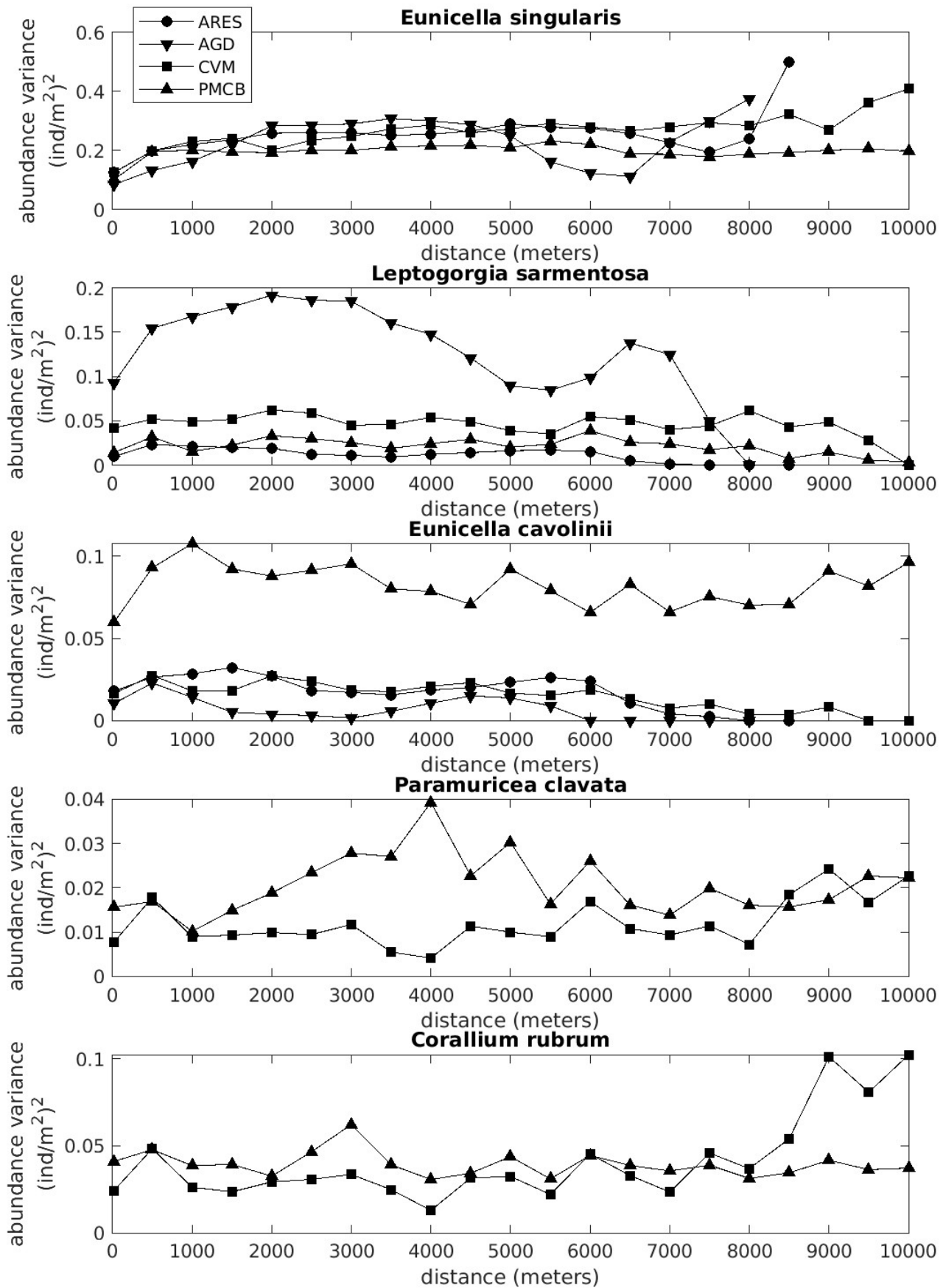


Fig. S2.- Empirical semivariograms describing the spatial variability of each of the five gorgonian species in the four sites (ARES, AGD, CVM, PMCB) where the species was present. Semivariograms were computed using population spatial density measured in the 696 evenly spaced geo-referenced locations covering the hard bottom habitat and clustering them by distance classes ranging from 500 meters to 10 kms, with a step of 500 meters increment. The nuggwvet of the semivariogram was computed using replicated measurements made at less than 20 meters distance, around each geo-referenced location (Guizien *et al.* 2022).

	0.02	0	0.03	0.03	0	0.18	0.23	0	0	0.06	0.33
2	1	0.77	0.74	0.87	0.21	0.11	0.13	0	0	0.26	0.04
	0.77	1	0.56	0.75	0.23	0.08	0.09	0	0	0.25	0.07
3	0.74	0.56	1	0.85	0.17	0.11	0.14	0	0	0.21	0.02
3	0.87	0.75	0.85	1	0.2	0.12	0.14	0	0	0.22	0.02
	0.21	0.23	0.17	0.2	1	0.18	0.2	0.01	0.01	0.27	0.02
3	0.11	0.08	0.11	0.12	0.18	1	0.74	0	0	0	0.03
3	0.13	0.09	0.14	0.14	0.2	0.74	1	0	0	0	0.09
	0	0	0	0	0.01	0	0	1	0.01	0.01	0
	0	0	0	0	0.01	0	0	0.01	1	0	0
3	0.26	0.25	0.21	0.22	0.27	0	0	0.01	0	1	0.31
3	0.04	0.07	0.02	0.02	0.02	0.03	0.09	0	0	0.31	1
1	0.14	0.08	0.15	0.14	0.15	0.01	0	0	0	0.45	0.18
TH	ROUGHNESS	RUGOSITY	SLOPE	TRI	TURBIDITY	Umax	Umin	NORTNESS	EASTNESS	SST	Tmax

Fig. S3.- Spearman determination coefficient among the 13 initials environmental predictors including latitude and longitude.

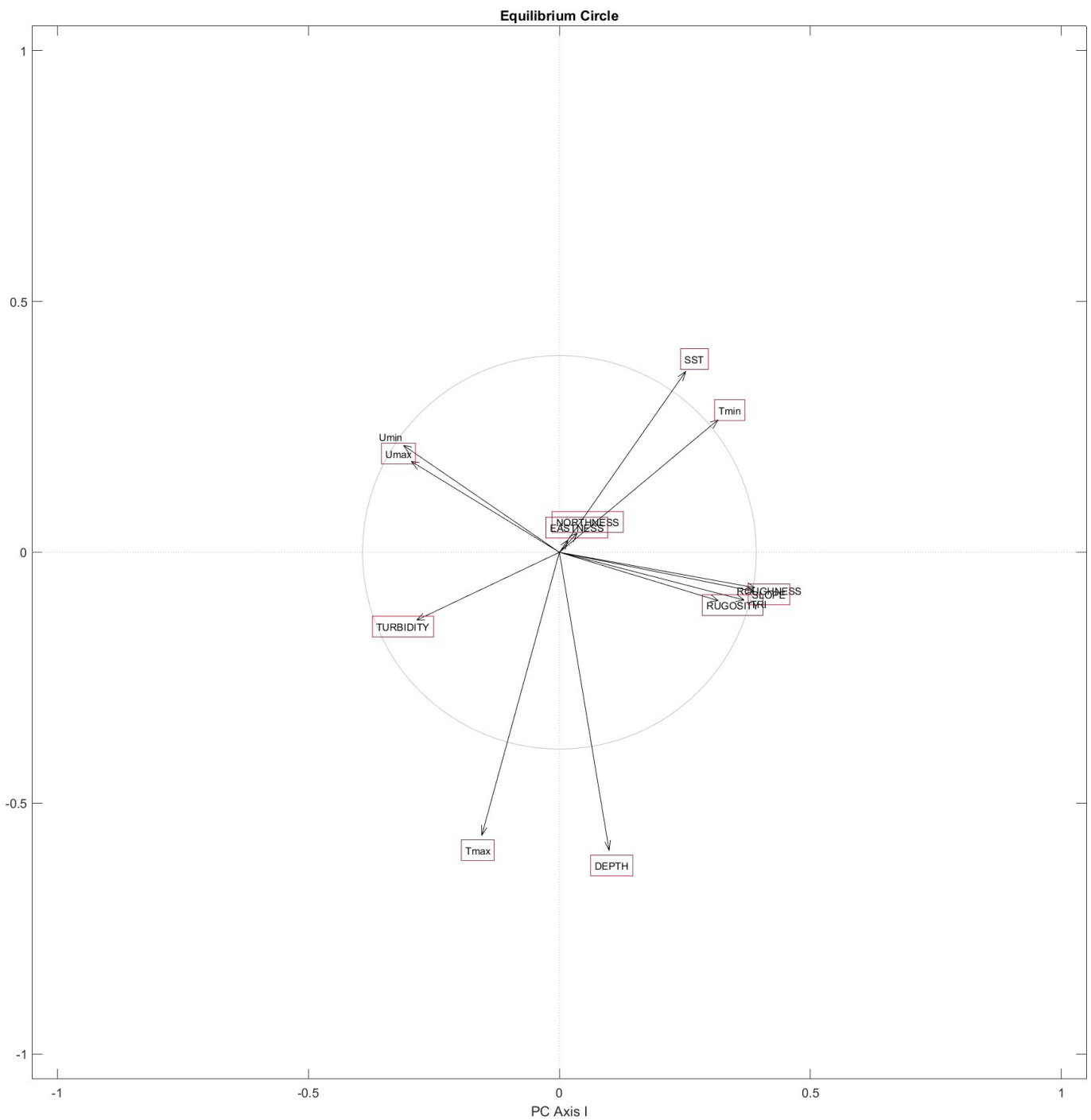
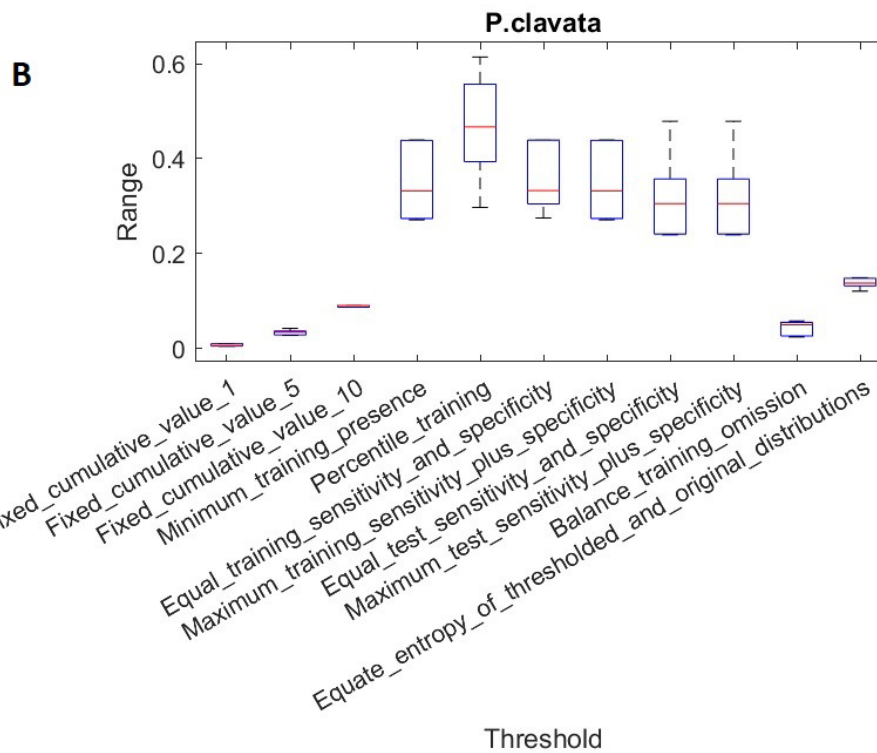
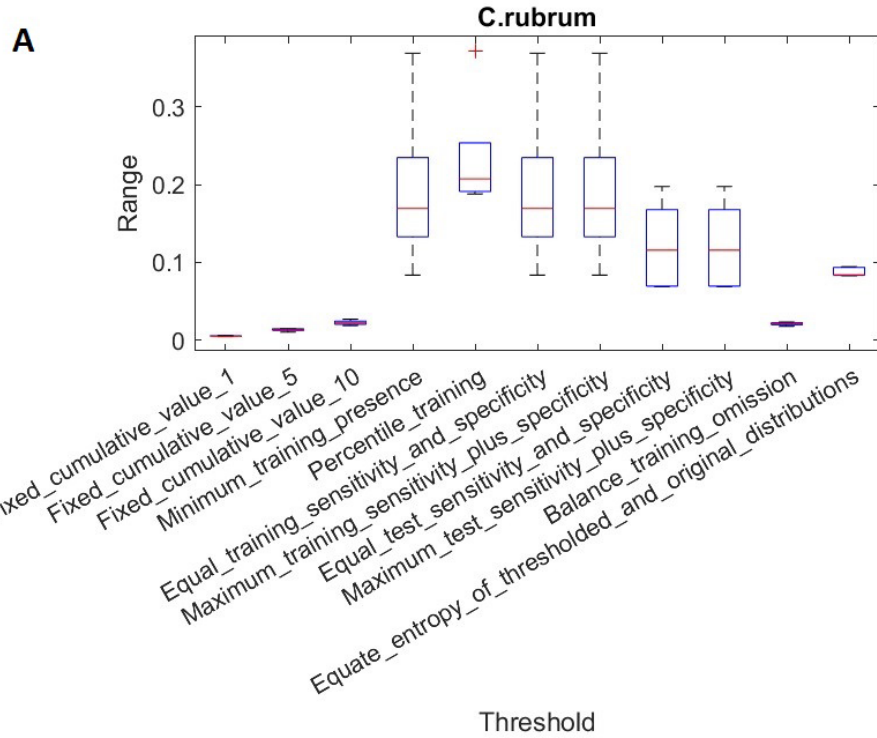
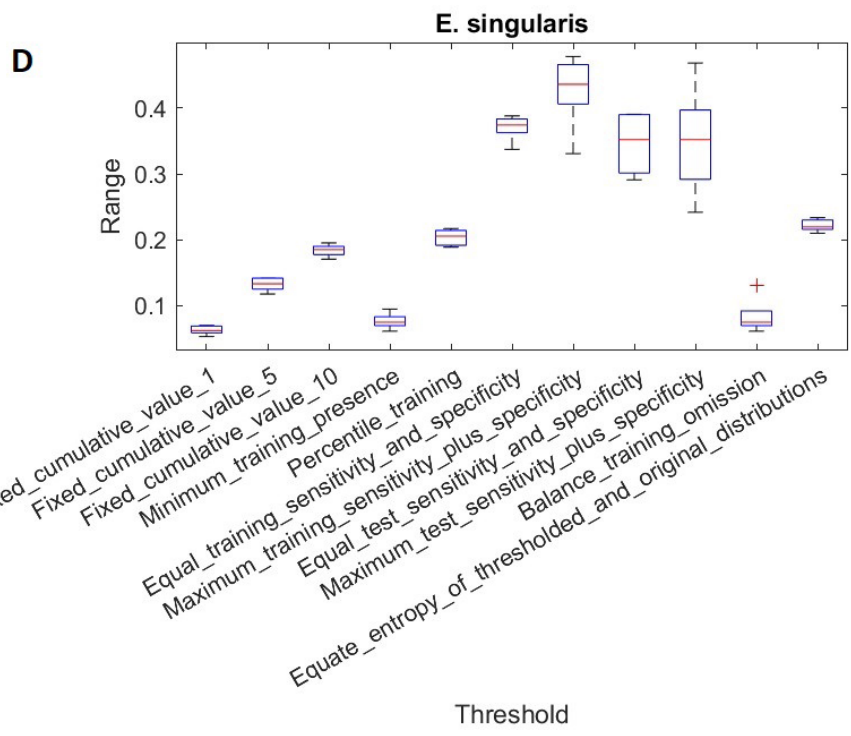
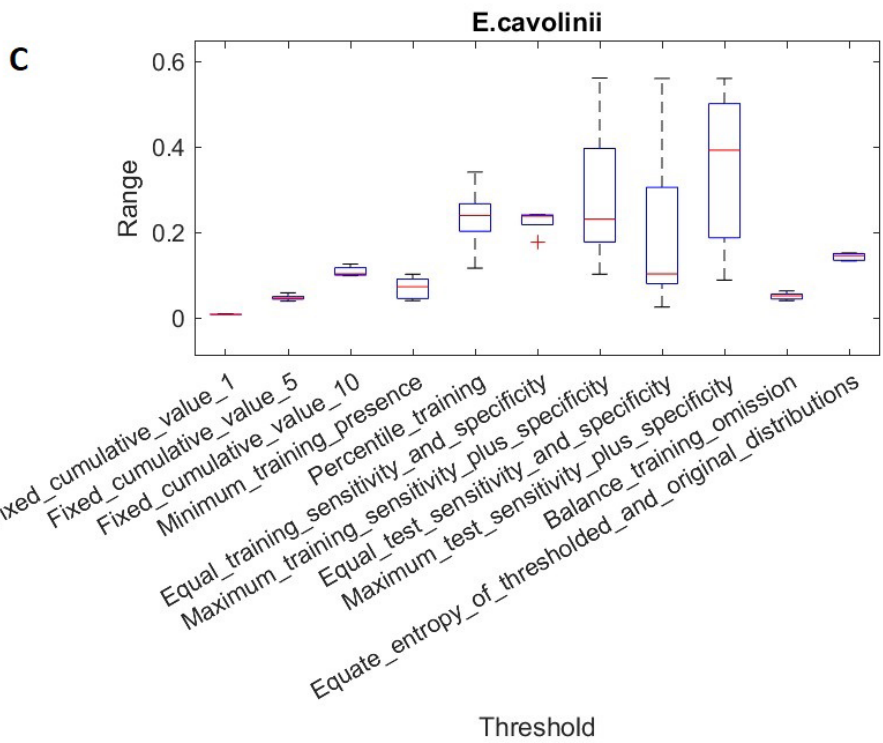


Fig. S4.- Principal Components Analysis on the 13 initials environmental predictors including latitude and longitude.





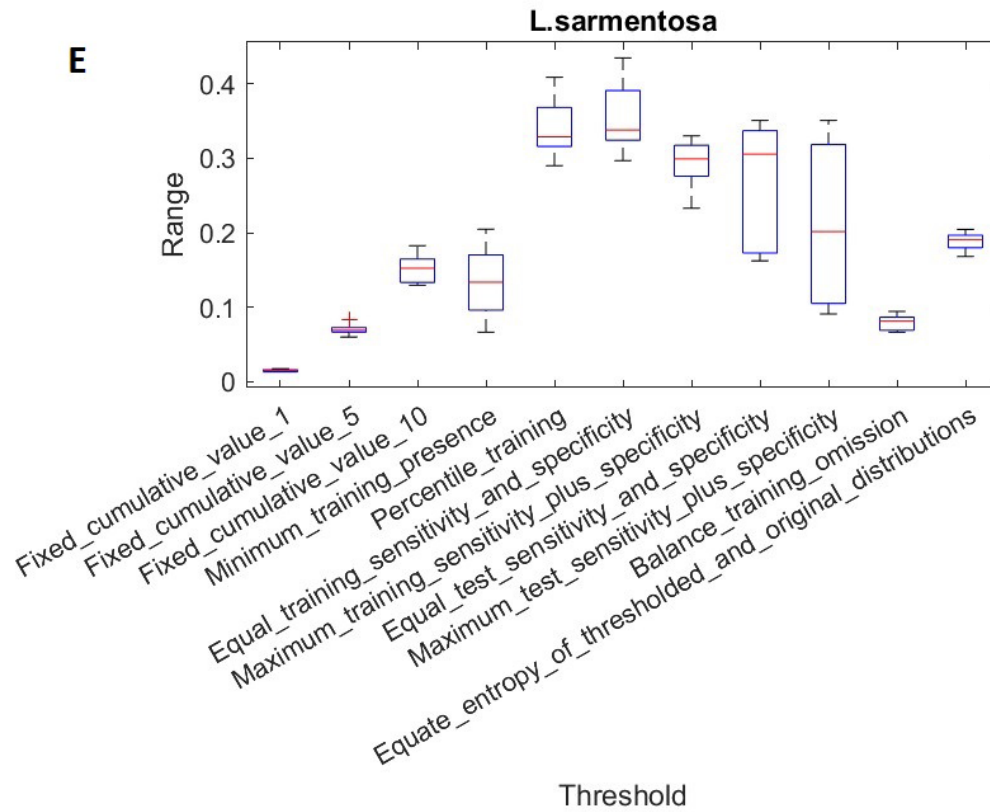
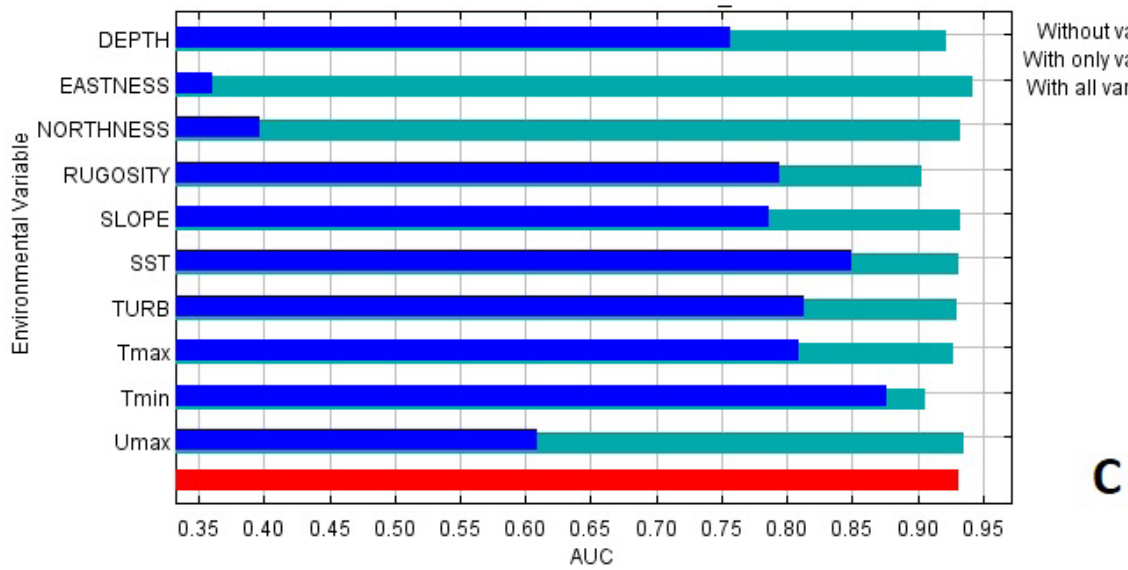
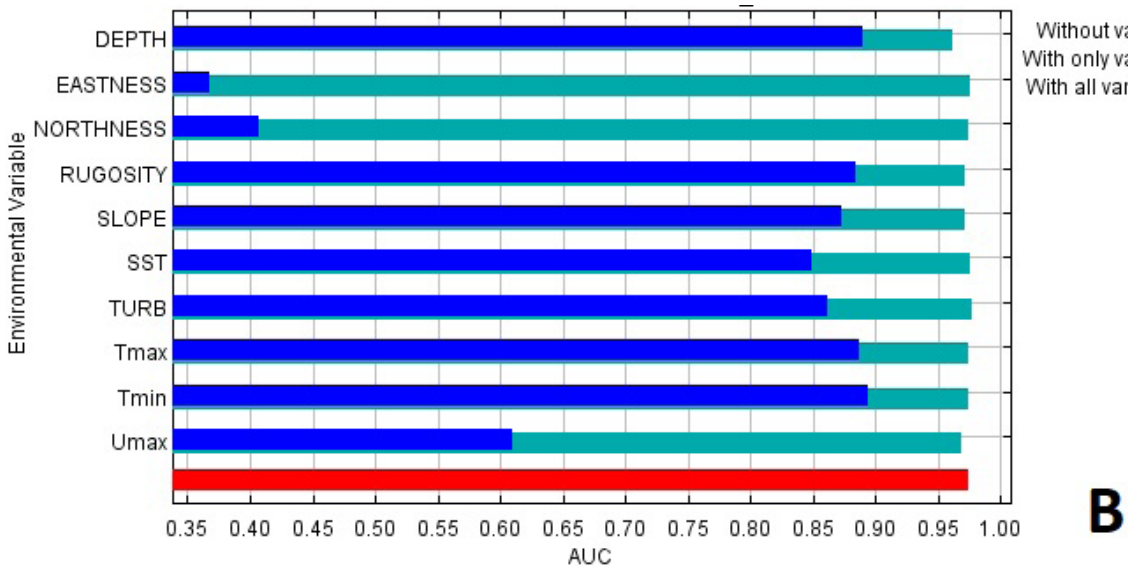
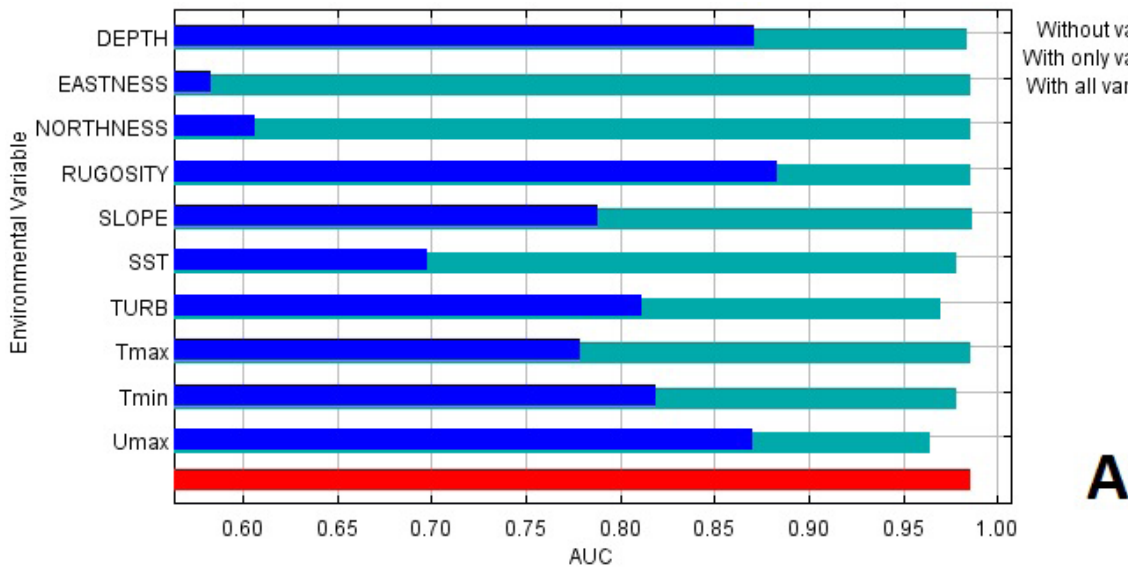


Fig. S5.- Threshold proposed by Maxent to predict suitability maps for various indices and for each species. Modelling with all predictors except geographical ones (Latitude / Longitude). A: *C. rubrum*, B: *P. clavata*, C: *E. cavolini*, D: *E. singularis*, E: *L. sarmentosa*.



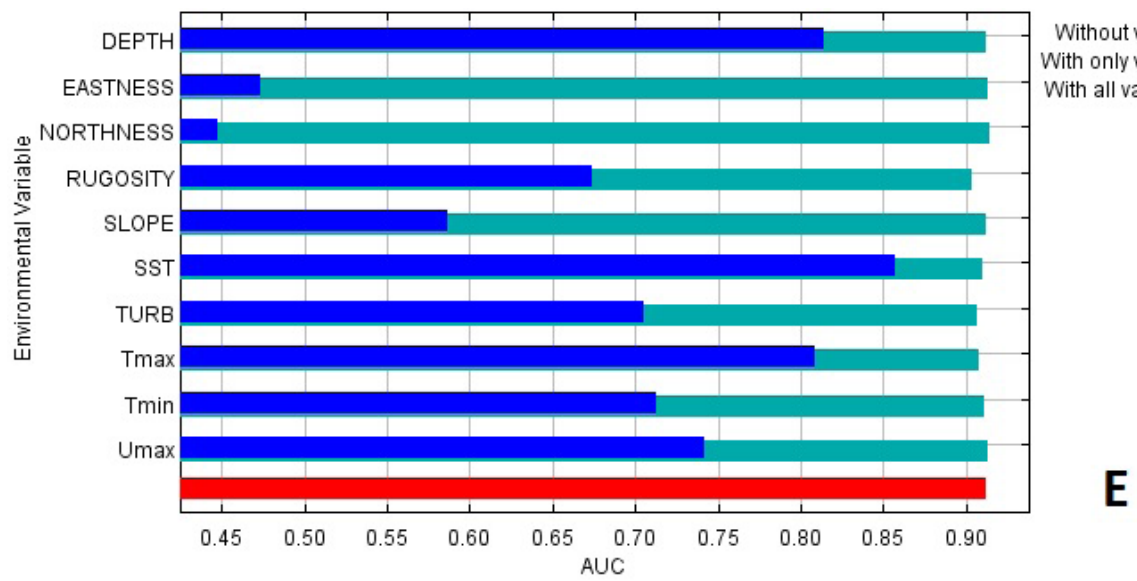
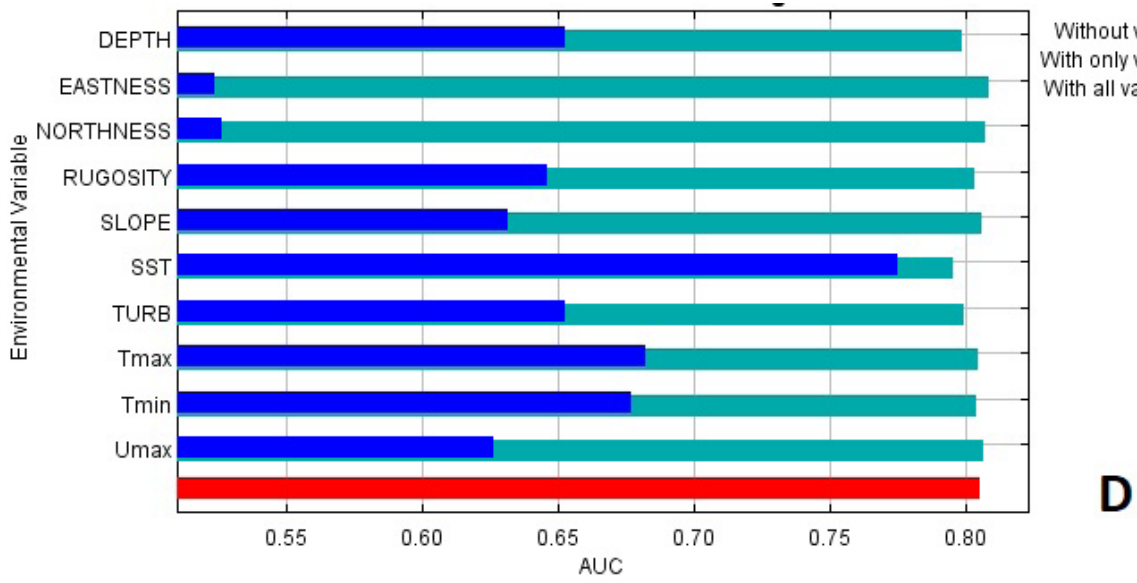


Fig S6.- Maxent AUC applying two Jackknife procedures across all predictors except geographical ones (Latitude / Longitude). A: *C. rubrum*, B: *P. clavata*, C: *E. cavolini*, D: *E. singularis*, E: *L. sarmentosa*. Without variable (light blue), with only variable (blue), with all variables (red). Variable is the same as predictor.

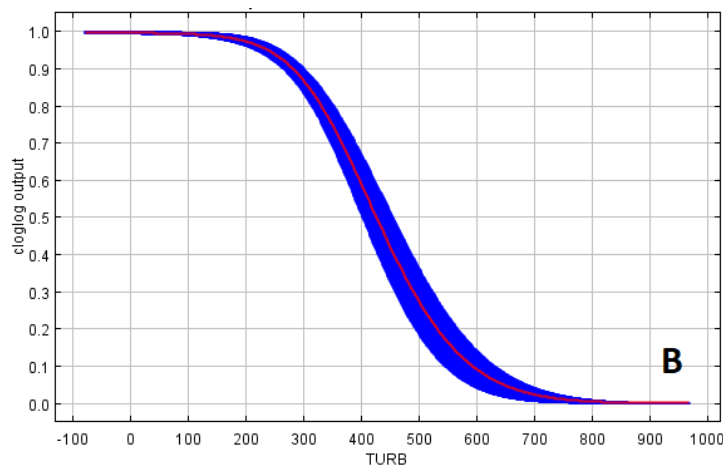
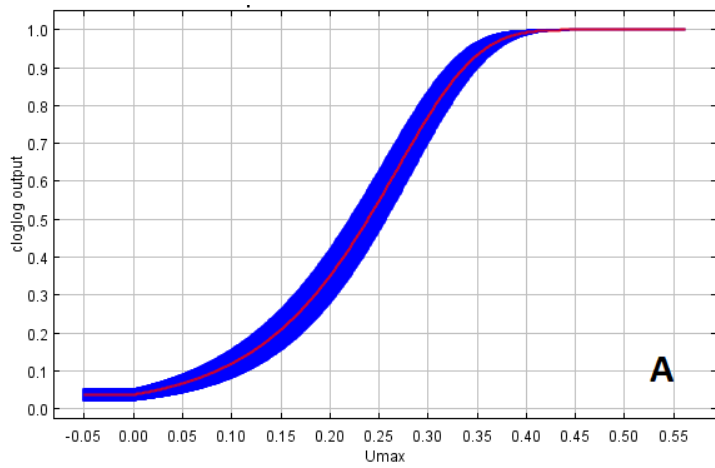
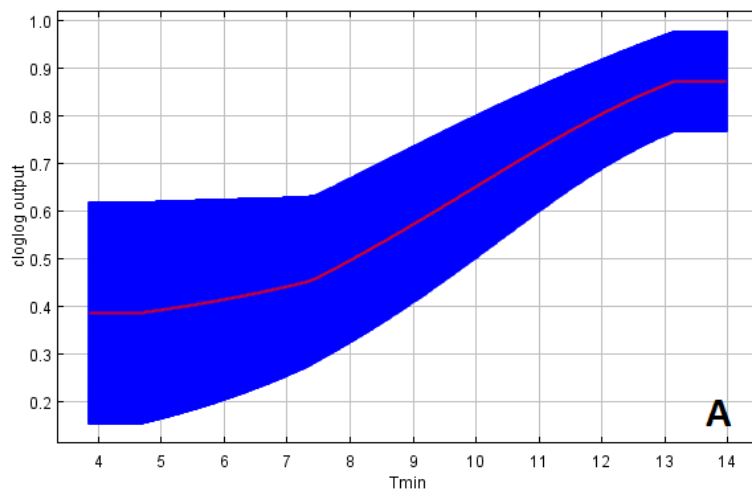


Fig. S7.- Response curves for the two significant predictors in the Maxent model for habitat suitability of *C. rubrum*: A: Umax, B: Annual average turbidity. The red lines indicate the mean values, while blue areas the standard deviation limits, resulting from 5 cross-validation model runs in cloglog output.



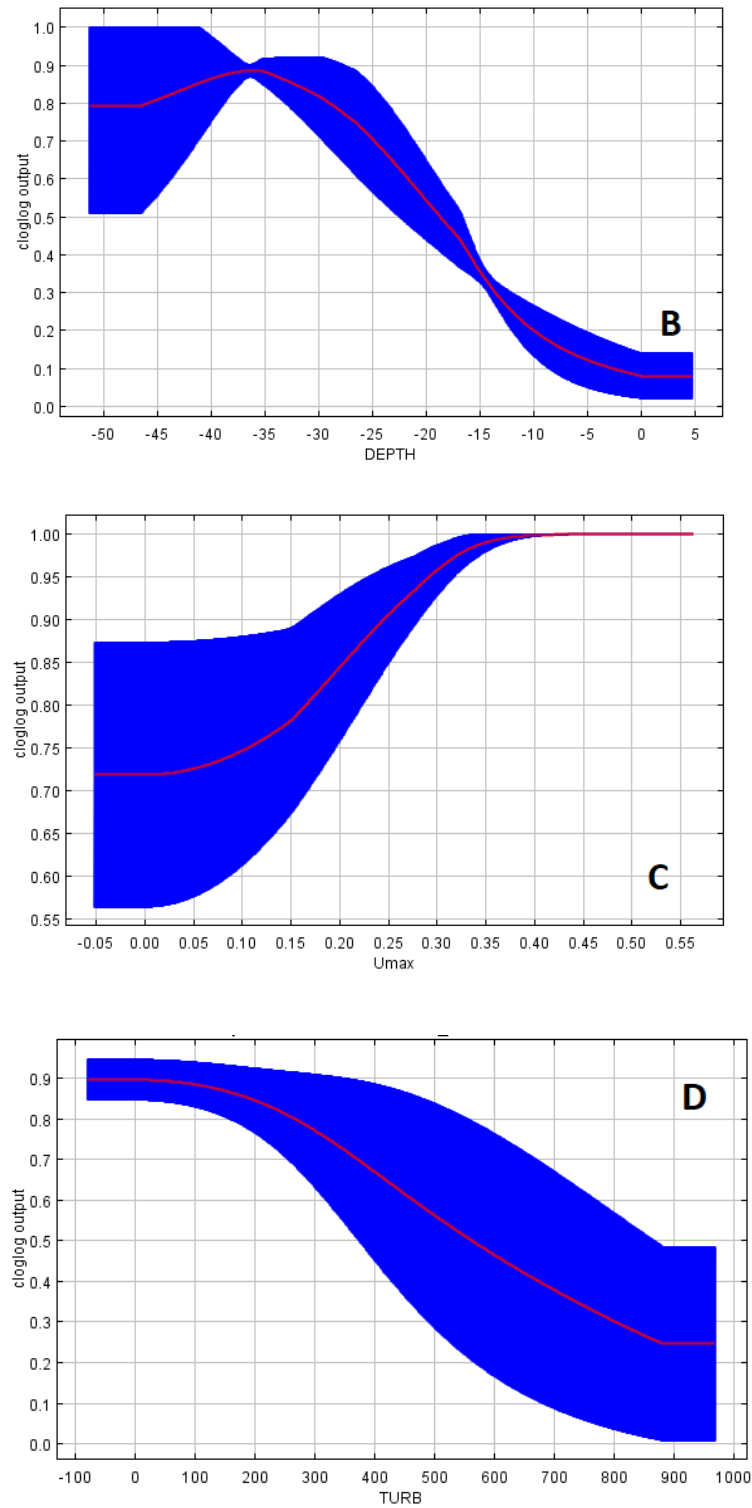


Fig. S8.- Response curves for the four significant predictors in the Maxent model for habitat suitability of *P. clavata*. A: Tmin, B: Depth, C: Umax, D: Annual average turbidity. The red lines indicate the mean values, while blue areas the standard

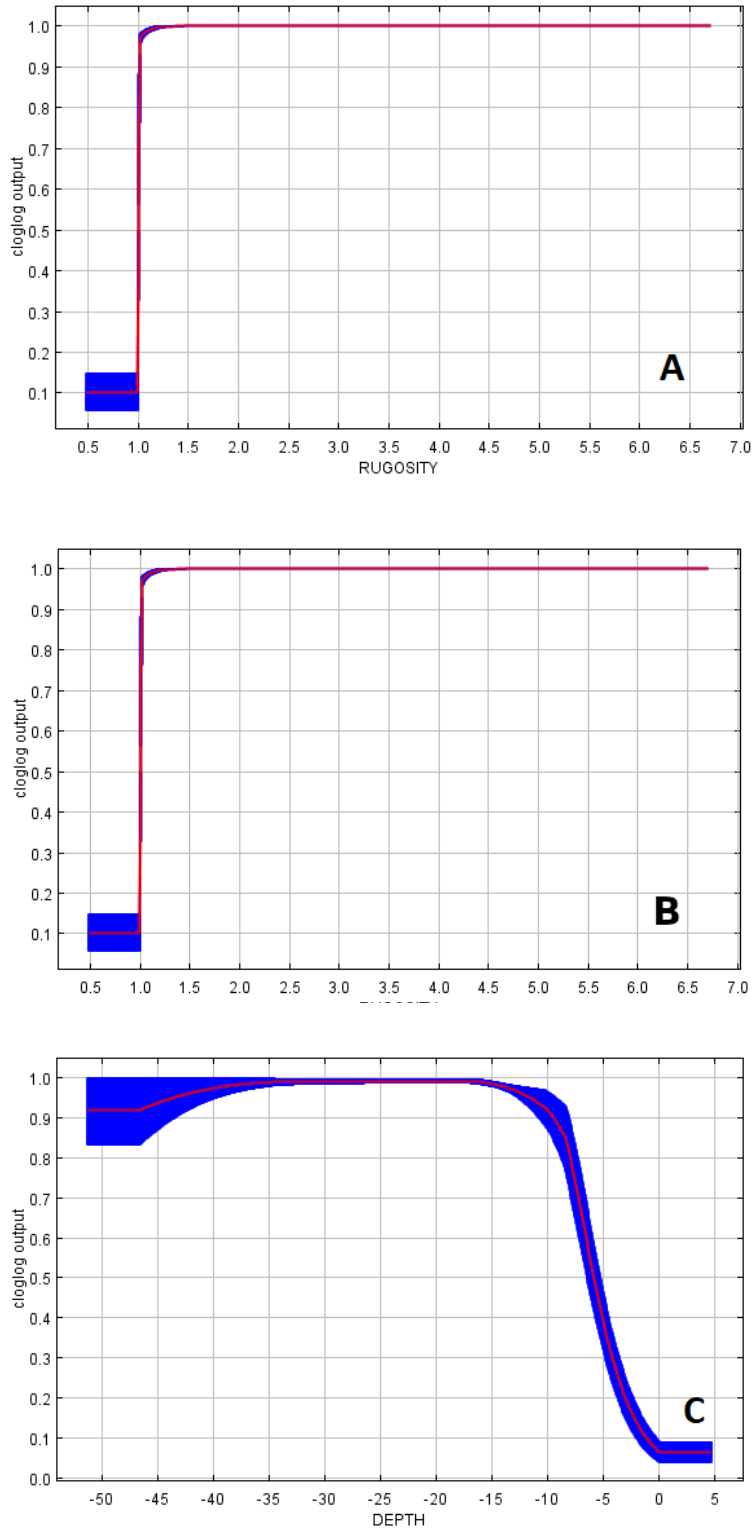
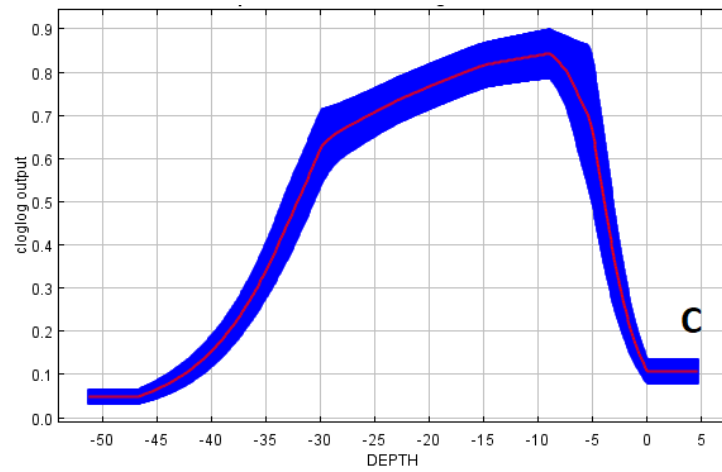
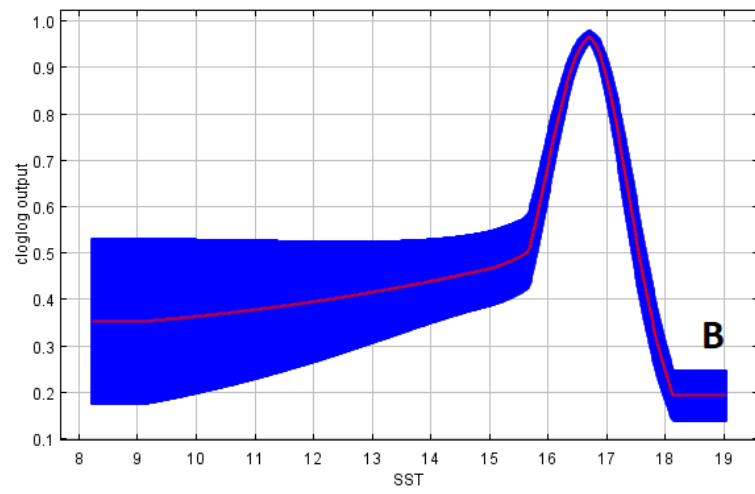
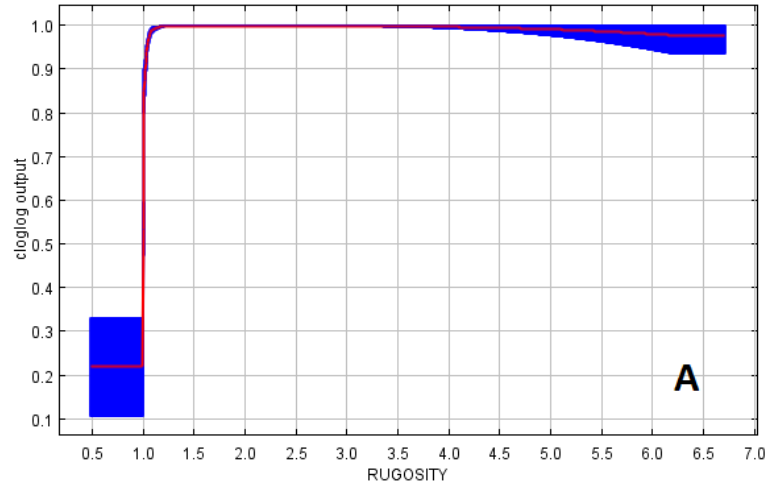


Fig. S9: Response curves for the three significant predictors in the Maxent model for habitat suitability of *E. cavolini*. A: Tmin, B: Rugosity, C: Depth. The red lines indicate the mean values, while blue areas the standard deviation limits, resulting from 5 cross-validation model runs in cloglog output.



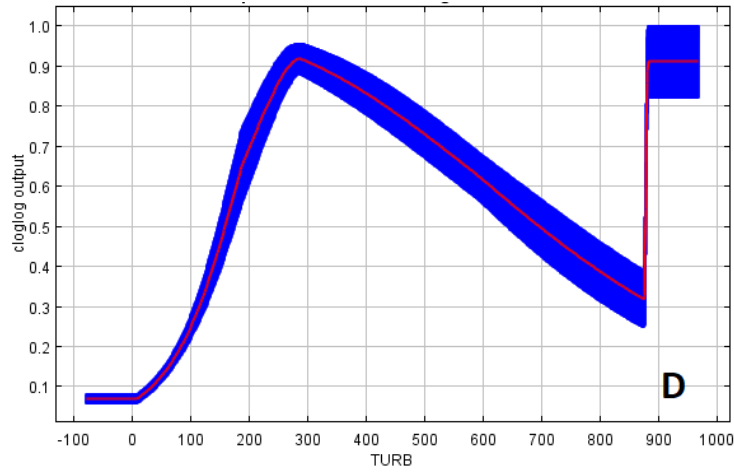
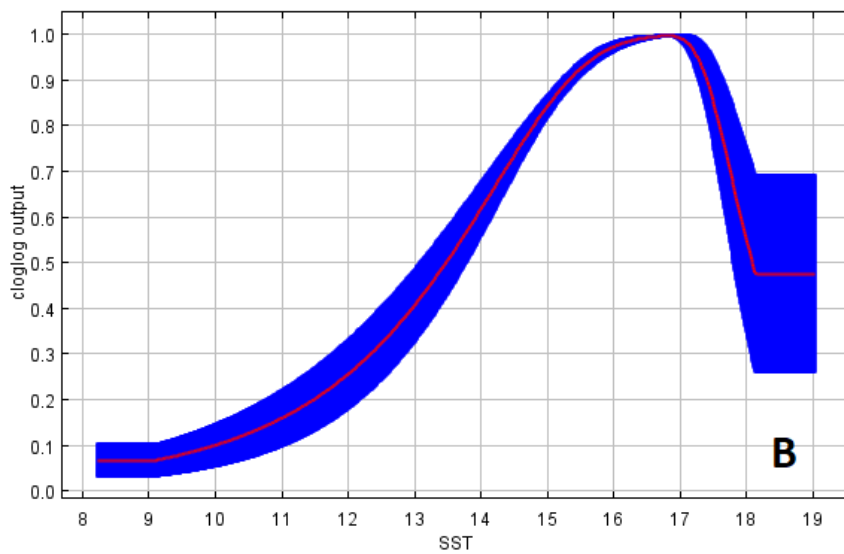
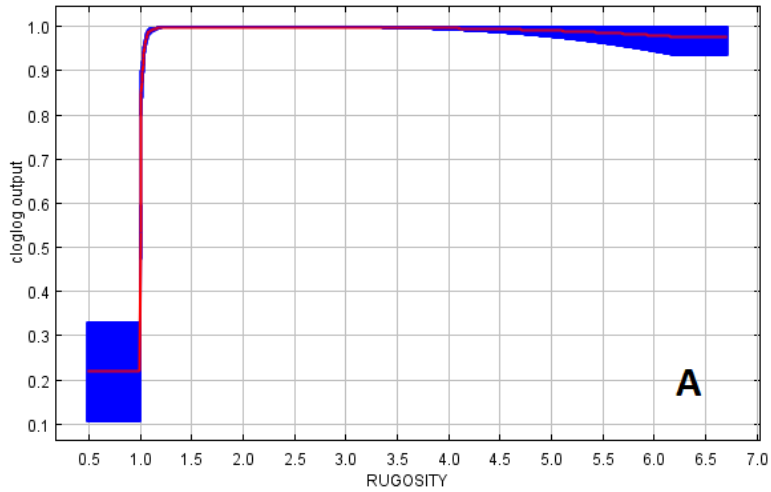
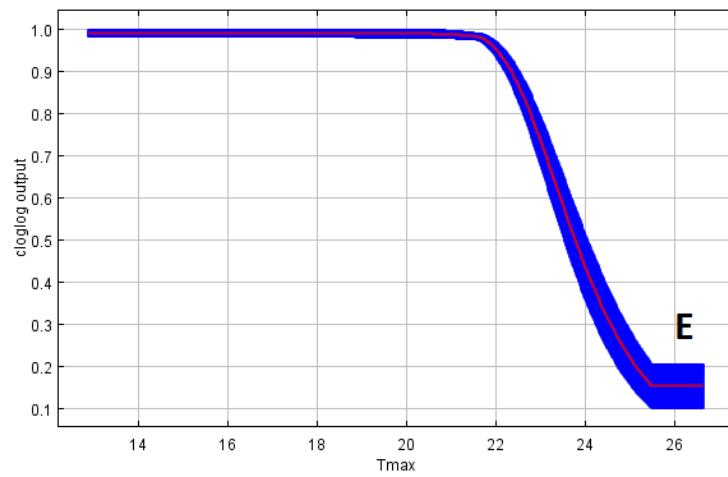
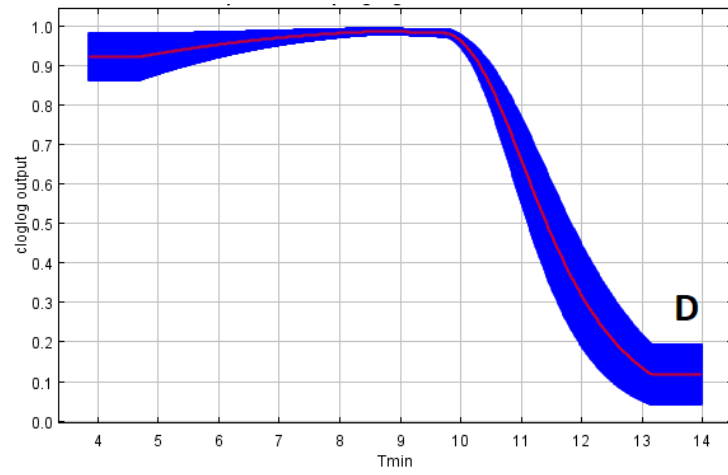
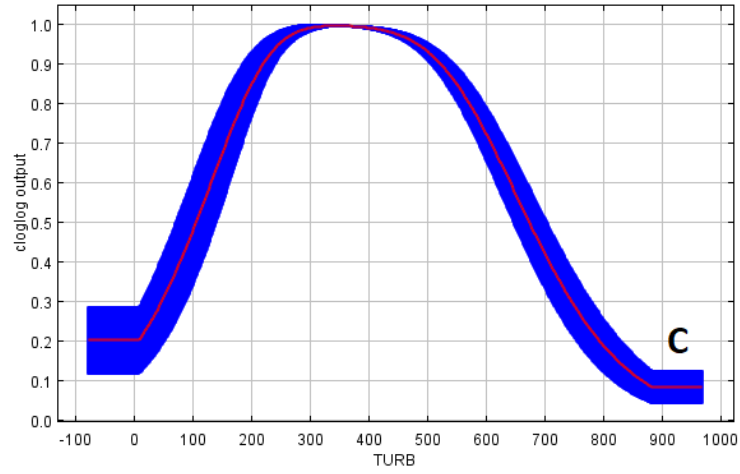


Fig. S10.- Response curves for the four significant predictors in the Maxent model for habitat suitability of *E.singularis*. A: Rugosity, B: Annual average SST, C: Depth, D: Annual average turbidity. The red lines indicate the mean values, while blue areas the standard deviation limits, resulting from 5 cross-validation model runs in cloglog output.





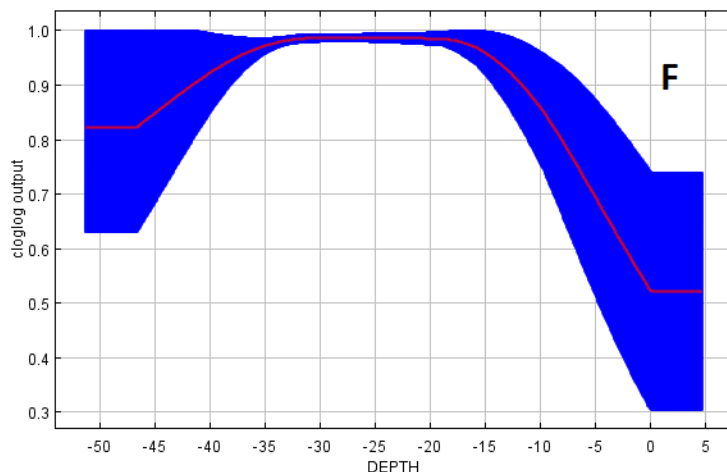


Fig. S11.- Response curves for the six significant predictors in the Maxent model for habitat suitability of *L. sarmentosa*. A: Rugosity, B: Annual average SST, C: Annual average turbidity, D: Tmin, E: Maximum bottom temperature Tmax, F: Depth. The red lines indicate the mean values, while blue areas the standard deviation limits, resulting from 5 cross-validation model runs in cloglog output.

	AUC	Variables	Percent contribution (%)	Permutation importance (%)
<i>C. rubrum</i>	0.9628	Umax	56.5	21.7
		Latitude	26.4	0
		Annual average turbidity	6.9	34.8
		Tmin	4.5	32.6
<i>P. clavata</i>	0.9731	Tmin	30.5	14.6
		Depth	27.9	32.2
		Umax	12.8	13
		Annual average turbidity	3.4	14.8
<i>E. cavolini</i>	0.9491	Rugosity	27.5	52.9
		Depth	20.5	14.8
		Latitude	5.8	18
<i>E. singularis</i>	0.8133	Latitude	56.3	16.4
		Longitude	11.7	23.2
		Depth	10.3	28.1
<i>L. sarmentosa</i>	0.9731	Longitude	38.7	65.6
		Rugosity	5.6	10

Table S1.- Significant environmental predictors identified for each species by Maxent with geographical predictors (Latitude / Longitude). Percent contribution and permutation importance.