



OPEN Beyond biodiversity: the role of *Paramuricea clavata* forests in shaping taxonomic and functional diversity

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Marine animal forests, composed of sessile suspension feeders such as gorgonians are known to host rich communities that support important ecosystem functions and services. These habitats are undergoing severe loss due to multiple pressures, with potential cascading effects on associated communities and their functions that remain poorly understood. To address this critical knowledge gap, we used fine-scale data to assess the role of *Paramuricea clavata* forests in supporting biodiversity and ecosystem functioning at multiple locations, in the Gulf of Naples (Italy, Western Mediterranean Sea). Through functional trait analysis, we compared taxonomic and functional diversity of benthic assemblages inside and outside *P. clavata* forests and investigated the loss of traits as a consequence of forest loss. Analyses revealed significant enhancements in both taxonomic and functional diversity within *P. clavata* forests, with observed increased species and functional richness. Trait-based investigations revealed a higher abundance of colonial heterotrophic species within forests, while outside, assemblages were dominated by low-longevity autotrophs, suggesting that *P. clavata* modifies environmental variables creating unique ecological conditions that favor specific traits. β -diversity measurements demonstrated increased compositional and functional turnover inside forests, indicating that *P. clavata* provides more available niches, supporting the replacement of species and functions. Our findings offer insights into how marine animal forests can structure marine communities, with broader implications for understanding biodiversity loss in changing marine ecosystems.

Keywords Habitat-forming species, Functional ecology, Functional traits, β -diversity, Turnover, Biodiversity loss

Biodiversity loss is pervasive across ecosystems, with expected large-scale ecological consequences for ecosystem functioning and the benefits nature provides to people¹. A major concern is the loss of habitat-forming species, as these species directly regulate resource availability for themselves and for other species. In terrestrial ecosystems, this process has been well-documented, with clear evidence that widespread deforestation drastically reduces biodiversity, alters species composition and traits, and has significant effects on various ecological processes². On the contrary, the response of marine ecosystems to the loss of habitat-forming species is less documented. Recent studies have indicated that marine macroalgal forests and seagrass meadows are at risk^{3–5}, with alarming projections for their future⁶. Dramatic loss in the biomass and biodiversity of ‘animal forests’ has also been globally observed^{7,8}, driven by multiple stressors ranging from local to global changes⁹. Marine animal forests are three-dimensional structures formed by habitat-forming sessile invertebrates (e.g., gorgonians, corals, sponges, bivalves). These assemblages are referred to as “forests” because, similar to terrestrial forests, they create complex architectures that enhance biodiversity and support multiple ecosystem functions⁷. Their role in providing food, protection and nursery to associated species, as well as contributing to the blue carbon budget, is

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well recognized¹⁰. As habitat-forming organisms, their loss is expected to reduce biodiversity and erode species traits, thereby disrupting important ecosystem services¹¹.

The Mediterranean Sea, considered a miniature ocean¹², is a hotspot of biodiversity although one of the most impacted areas of the planet, where pollution, overfishing, habitat destruction and non-indigenous species introductions are producing large effects at basin scale^{13,14}. Climate change is severely affecting Mediterranean marine ecosystems faster than expected¹⁵, with unprecedented loss of habitat formers¹⁶. In this respect, the red gorgonian *Paramuricea clavata* Risso (1827) is a paradigmatic example. It is a long-lived species listed as a vulnerable species by the International Union for Conservation of Nature¹⁷, with slow growth rate, annual recruitment and late reproductive maturity, making it particularly vulnerable to human disturbances^{18,19}. It shows a large-scale distribution²⁰ living at depths ranging from 15 to 200 m^{21,22} and contributing to the three-dimensional complexity of coralligenous outcrops, a Mediterranean biogenic temperate reef formed by complex and heterogeneous benthic assemblages^{23,24}. *Paramuricea clavata* has a critical functional role since it is supposed to stabilize key environmental variables such as light intensity, water flows, and sedimentation rate, thus supporting the settlement and growth of underlying benthic species^{25,26}. Several studies have reported mass mortality events of *P. clavata* populations due to increasing frequency of extreme climate events (marine heat waves, storms) together with other sources of disturbance such as fishing pressure, ultimately affecting population's resilience and resistance^{9,27–31}, also highlighting dramatic effects on the associated community^{9,32,33}. However, despite the importance of this species, the consequences of its loss are still little known. Fine-scale data are needed to understand if the presence of *P. clavata* forests support assemblages with distinct structural and ecological features able to provide specific ecosystem processes and associated ecosystem services. Such data are essential for predicting future changes in this habitat and the associated ecological processes under global change scenarios, addressing the long-term ecological implications of the decline and loss of *P. clavata*.

Simultaneously exploring taxonomic and functional aspects of biodiversity provides a more comprehensive understanding of community assembly processes³⁴. Functional traits are those specific characteristics or attributes of organisms that influence their performance and interactions within ecosystems³⁵. The unique combination of traits in a species identifies a functional entity (FE) meaning that species that share the same traits belong to the same FE^{36,37}. The functional trait approach presents some limitations, since there is a deep gap of knowledge on trait-specific information which leads to their simplification by scientific experts. Nevertheless, this methodology can provide valuable insights to complement taxonomic analyses, allowing for a robust assessment of how changes in species composition translate into functional shifts. This approach is particularly useful in species-rich communities, such as coralligenous assemblages, where directly measuring the full range of ecosystem functions is often unfeasible^{38,39}. Recently, trait-based indices have been introduced as an alternative 'functional' approach to assess the relationships between biodiversity and human disturbances. Indeed, these indices have the potential to determine the cause of change in systems by investigating the type of traits affected. Evidence showed that human pressures selectively remove a distinct suite of traits from assemblages^{40,41}. For instance, ocean acidification reduces the abundance of calcifying species with three-like morphologies, favoring low-lying fast-growing species across different habitats⁴². Also, mechanical disturbances such as bottom trawling have negative effects on large epifaunal filter feeders, which are more sensitive to physical stressors, and are replaced by mobile infaunal scavengers^{43,44}. This approach enables to identify the most vulnerable traits to a specific source of disturbance, allowing to anticipate potential shifts in ecosystem properties.

For a broader understanding of the ecological role of *P. clavata*, assessing β -diversity is essential, given that the structural complexity created by habitat-forming species has been shown to influence not only species richness and community composition, but also spatial heterogeneity⁴⁵. β -diversity refers to the variations in species composition across different habitats or spatial units and it can be partitioned into species replacement (turnover) and richness differences (nestedness), allowing to disentangle the contribution of the two mechanisms to the heterogeneity and thus to understand causes generating differences in species composition⁴⁶. While research has primarily focused on compositional β -diversity, combining both compositional and functional β -diversity provides a valuable lens for understanding how ecological processes and human pressures drive species assemblages and distribution, offering key insights for species conservation and ecosystem management⁴⁷. Because functional traits reflect species' adaptive strategies to the environment, numerous studies on functional β -diversity have rapidly emerged over the past two decades. These studies show contrasting turnover and nestedness-resultant components of compositional and functional β -diversities⁴⁸ and demonstrate the importance of considering the multifaceted nature of biodiversity when examining community assembly³⁴.

The ecological value of *P. clavata* forests has been recently explored. Gómez et al.³³ reported that marine heatwaves (MHWs) can affect the functional structure of *P. clavata* dominated assemblages with detrimental consequences on several ecological processes and their associated ecosystem functions. However, comparisons with coralligenous assemblages outside the forests also revealed that *P. clavata* can mitigate the effects of warming by helping to maintain the original assemblage structure dominated by macroinvertebrates⁹. Complementing this, Teixidó et al.³² documented that the removal of *P. clavata* facilitates the proliferation of opportunistic turf-forming algae and invasive species. Taken together, these findings emphasize the ecological relevance of *P. clavata*, and highlight the need to better understand how forest presence or absence shapes taxonomic and functional diversity. To address this, we used fine-scale data with non-destructive sampling methods from replicated locations in the Gulf of Naples (Italy, Western Mediterranean Sea), including species presence-absence, abundance, and functional traits. This comprises 'effect' traits (e.g. morphological, physiological and phenological features of species involved in the understory) and 'response' traits (e.g. reproductive and dispersal strategies). We applied a combined taxonomic and functional framework to test the hypothesis that the understory assemblages associated with *P. clavata* forests exhibit distinct features compared to coralligenous assemblages where the forests are absent.

Material and methods

Study area

The Gulf of Naples (Italy) is a semi-enclosed embayment nestled within the southeastern Tyrrhenian Sea (Mediterranean Sea). The basin is South-West oriented and bounded by the islands of Procida and Ischia to the North, and by the island of Capri and the Sorrento peninsula to the South. Six locations were sampled based on prior knowledge of the presence of *P. clavata* forests within the operative depth of recreational scuba diving of 30–40 m. The selected locations were: Ischia (Punta Sant'Angelo) (1) and Procida (Punta Pizzaco) (2), located within the Regno di Nettuno Marine Protected Area (MPA); Scoglio Penna (3) and Scoglio del Veruce (5), within the Punta Campanella MPA; Banco di Santa Croce (6), protected area by Ministerial Decree of June 15, 1993 and the location of Capri (4), out of protection boundaries (Fig. 1). For each location, we randomly selected two sites (A, B) about 50 m apart from each other, with similar slope and exposition. At each site, data were collected under two conditions: an “inside” condition, located within *P. clavata* forests, and an “outside” condition, corresponding to adjacent areas of the same coralligenous outcrop where the forests were naturally absent. This design allowed for a direct comparison of benthic assemblages under similar environmental settings, differing only in the presence of the habitat-forming gorgonian.

Data collection

Data collection has been carried out during scuba diving surveys in summer 2022. Within each site, photographic sampling of benthic assemblages was conducted using ten random replicate quadrats of 25×25 cm for each of the two conditions: inside *P. clavata* forest and in the adjacent zone outside *P. clavata* forest. Additionally, six random quadrats of 50×50 cm were used in each site within the forest to count *P. clavata* colonies to later assess the density and population structure (Fig. S1). Each *P. clavata* colony inside these quadrats has been measured with a ruler from the base to the end of the farthest tip⁴⁹. The sampling unit of six 50×50 cm was chosen following Linares et al.¹⁹, who found the stabilization of the s.e.m. (standard error of the mean) as a proportion of the mean colony density with a sample size of 1.5 m^2 . The photographic sampling was carried out using Canon G7X Mk III in Nauticam housing and equipped with two strobe lights. The percentage cover of sessile organisms was quantified using the photoQuad_v1_4 software⁵⁰. During the analysis, each species has been identified at the

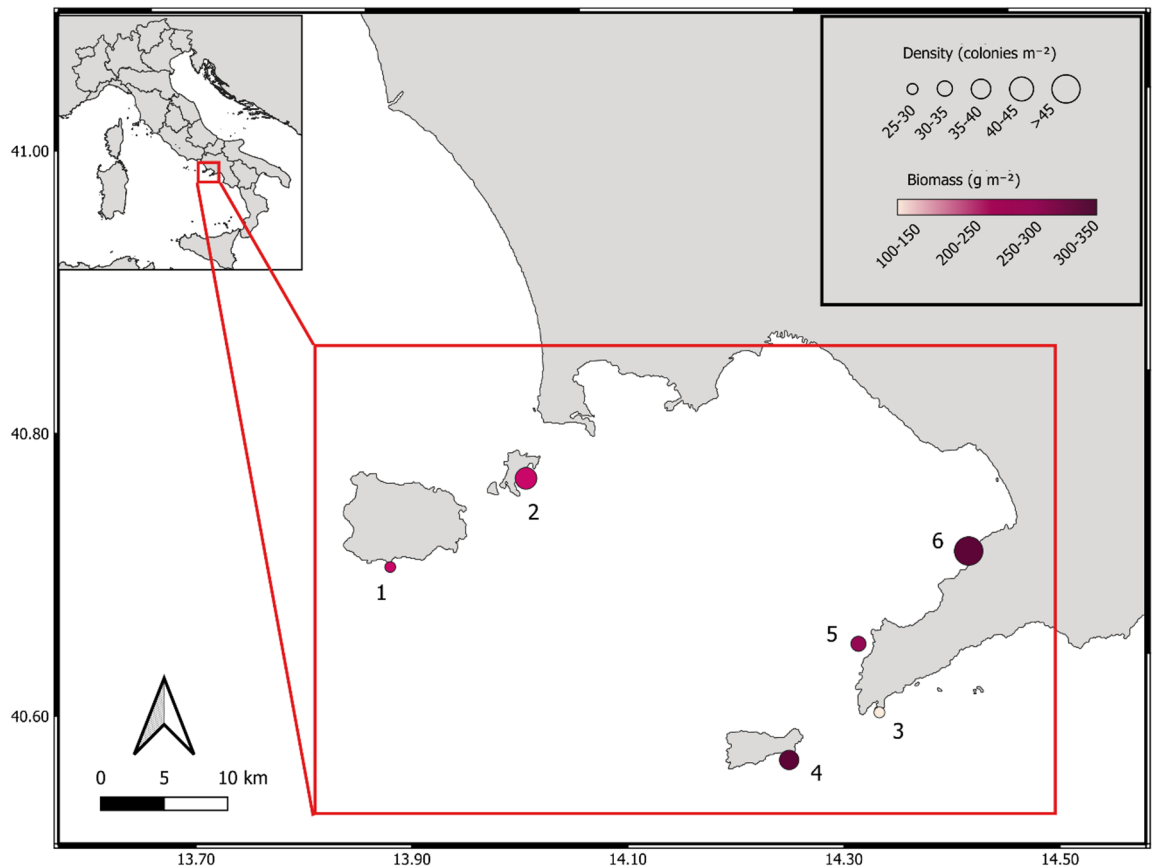


Fig. 1. Distribution of the sampling locations in the study area. Mean density and biomass values of the *P. clavata* colonies are represented as different circle colours (biomass) and dimensions (density). The map was produced with QGIS1 3.28.6 (QGIS.org 2023. QGIS Geographic Information System. Open-Source Geospatial Foundation Project. <http://qgis.org>).

lowest possible taxonomic level, and when specific identifications were not feasible, a morpho-functional group has been assigned (e.g. algal turf, encrusting sponge)

Data analysis

Paramuricea clavata forests

To assess the health status of *P. clavata* forests, mean height, density and biomass of each colony have been calculated for each site. Height measurements collected in the field were used to estimate the biomass using the relationship $B = 0.002H^{2.61}$ reported by Coma et al.⁵¹ and corrected by Linares et al.¹⁹, where B is the biomass of dry weight (DW) in grams and H is the colony height in centimeters. The population structure was assessed by assigning each colony to a size class with a 10 cm range and then reporting the frequency distribution of each size class across sites. Size distribution was analysed using the skewness and kurtosis coefficients^{19,52}. Skewness measures the asymmetry of a distribution relative to its mean. When skewness is significant, it indicates that the distribution is not symmetrical. Positive skewness suggests a predominance of smaller size classes within the population, while negative skewness indicates a dominance of larger size classes. Kurtosis assesses the sharpness of the distribution's peak around its central mode. A significant kurtosis value implies that the distribution has longer tails compared to a normal distribution, reflecting a higher prevalence of certain size classes within the population. Skewness and kurtosis coefficients are considered significant if the ratio of their values to their standard error exceeds ± 2 ⁵². Population dynamics of marine animal forests are regulated by density-dependent processes determined by the intra-specific competition for the use of resources such as food and space⁵³. As a result of this self-thinning mechanism, pristine populations are characterized by large colonies, while in young or impacted populations, high densities of small colonies saturate the space^{19,54}. We assessed the relation between biomass and density of *P. clavata* populations through a linear regression with the log–log model as a descriptor of population conservation status¹⁹.

Experimental design and statistical analysis

Statistical analyses were performed to test the hypothesis that *P. clavata* forests have the role of increasing local compositional and functional diversity. To test these hypotheses, we applied a multifactorial design consisting of three factors: *location* (Lo, 6 levels, random), *site* (Si, 2 levels, random and nested in location) and *condition* (Co, 2 levels, fixed and orthogonal), with $n = 10$ replicates. The analyses are described in the sections below.

Functional traits and trait space Twelve categorical and/or ordinal functional traits were identified on the base of previous studies on benthic assemblages^{37,47}: (1) Morphology, (2) Coloniality, (3) Maximum longevity, (4) Size, (5) Epibiosis, (6) Energetic resource, (7) Major photosynthetic pigment, (8) Feeding strategy, (9) Potential of asexual reproduction, (10) Growth rates, (11) Defences, (12) Propagules (Table S1). Functional traits were subdivided into categories, and, for each species, a category was assigned based on the relative trait determined through extensive bibliographic research (see supplementary materials) on biological and ecological characteristics of that species (Table S1). Subsequently, each species has been classified into a functional entity (FE) (i.e. groups of species with unique combinations of functional traits)³⁶. Functional richness (FRic) measures the amount of functional trait space occupied by a community and thus reflects the diversity of functional traits present. It has been calculated as the percentage of the volume of the multidimensional trait space occupied by all FEs within the functional space⁴⁰. To create this multidimensional trait space, a Principal Coordinates analysis (PCO) was performed on FEs, based on a Gower dissimilarity matrix, chosen since it allows the analysis of mixed types of data while giving them equal weight⁵⁵. Seven PCO dimensions were selected, based on the lowest mean squared-deviation index value (mSD = 0.0022), to ensure a faithful representation of trait-based differences between species. The coordinates of each FE obtained by PCO were used to calculate the FRic for each quadrat sampled during the study to assess statistically significant differences between conditions, and across sites and locations. These coordinates were also used to visualize spatial differences in functional richness. To examine differences in trait composition and abundance between conditions in space, Community-Weighted Mean (CWM) was calculated as the average of trait values for species at each quadrat weighted by the relative abundance of each species possessing that trait. CWM of traits is a valuable index for evaluating shifts in mean trait values within communities due to environmental selection for specific functional traits^{56,57}.

Univariate and multivariate analyses A three-way univariate analysis of variance was carried out using PERMANOVA based on Euclidean distances of untransformed data⁵⁸ to assess differences in species richness, in the number of FEs and in the FRic, between conditions and across locations and sites. Each term was tested using a maximum of 999 permutations. To investigate the effect of *P. clavata* on taxonomic and functional structure, and on CWM, multivariate analyses were performed using PERMANOVA based on the Bray–Curtis resemblance matrix calculated on fourth root transformed data to reduce the differential between dominant and rare species. The PERMANOVA test was performed since it is less sensitive to dispersion than other tests like ANOSIM and the Mantel test, particularly in balanced designs, and it was therefore considered appropriate for testing our hypothesis⁵⁹. Each term was tested using a maximum of 999 random permutations of the appropriate units⁶⁰, and post-hoc pairwise comparisons were carried out when significant results were obtained. Non-metric multidimensional scaling ordinations were carried out on species cover and CWM to display differences on taxonomic and functional diversity between conditions at the scale of sites, since the interaction term Si(Lo) × Co was found significant in the both multivariate analysis. Statistical analysis has been performed using PRIMER software (v.7).

Compositional and functional β -diversity According to Villéger et al.⁶¹, compositional β -diversity is calculated as the ratio of species not shared between assemblages relative to the total number of species. By analogy, the

functional β -diversity is quantified as the ratio of FEs not shared between assemblages relative to the total FEs. Compositional and functional β -diversity analyses, based on the Jaccard resemblance matrix on presence/absence transformed data, were carried out within condition and partitioned into the two β -diversity component nestedness and turnover. Turnover can be defined as the degree of species replacement reflecting the selective differentiation of species among assemblages because of environmental sorting^{46,62}. Nestedness, on the other hand, refers to the hierarchical arrangement of species assemblages, wherein species present in species-poor habitats are subsets of those found in species-rich habitats, often in response to environmental gradients⁶³. A permutational analysis of multivariate dispersion (PERMDISP)⁵⁴ was then carried out to test statistically significant differences in the heterogeneity between conditions (In vs. Out). PCO based on Jaccard dissimilarity matrix was performed for compositional and functional β -diversity and its components to visualize differences in terms of community heterogeneity between conditions. FRic, CWM and β -diversity analyses were performed using the R functions from the 'FD', 'tripack', 'geometry', 'matrixStats' and 'betapart' R package (R v 3.4.1, R development Core Team, 2017)^{64–68}.

Results

Paramuricea clavata forests

The mean *P. clavata* density ranged from 27.3 ± 5.9 to 65.3 ± 27.1 colonies/m² \pm s.e.m., while lowest and highest biomass values corresponded to 100.1 ± 89.7 and 505.4 ± 367.7 g dry weight/m² \pm s.e.m., respectively (Table S2). Small colonies (< 10 cm) were the most represented in all investigated populations (Fig. 2). Skewness coefficients confirmed that most of the size class distributions (11 out of 12 sites) were significantly positively skewed, indicating the prevalence of small size classes. The kurtosis coefficients also showed a significantly positive trend in 10 out of the 12 sites, highlighting the presence of long tails in the size class distribution of *P. clavata* colonies among sites. *Paramuricea clavata* populations showed no significant relationship between density and biomass ($R^2 = 0.097$; $p = 0.3234$) (Fig. S2).

Species richness, functional entities and functional richness of the understory assemblages in presence and in absence of the forests

A total of 80 species/taxa were found in the understory assemblages and grouped in 63 FEs. Univariate analyses revealed the effect of the presence of *P. clavata* forests on the number of species ($p = 0.012$), the number of functional entities (FEs, $p = 0.009$) and functional richness (FRic, $p = 0.042$) (Table S3). Differences were consistent across sites and locations for the three variables (Table S3). More specifically, the mean number of species and FEs were higher inside the forests (9 ± 0.3 s.e.m.; 8.8 ± 0.2 s.e.m. respectively) compared to the areas outside (7.5 ± 0.2 s.e.m.; 7.5 ± 0.2 s.e.m. respectively). Additionally, FRic was also higher inside the forests at all locations (Fig. 3), with a higher mean value (20.9 ± 3.2 s.e.m.) compared to the areas outside (9.3 ± 2.1 s.e.m.).

Structural and functional changes in the understory assemblages in presence and in absence of the forests

The multivariate analysis showed a significant effect of the forests on the taxonomic structure of the understory assemblages, differing across sites ($p = 0.001$) (Table S4). However, a posteriori pairwise comparison indicated

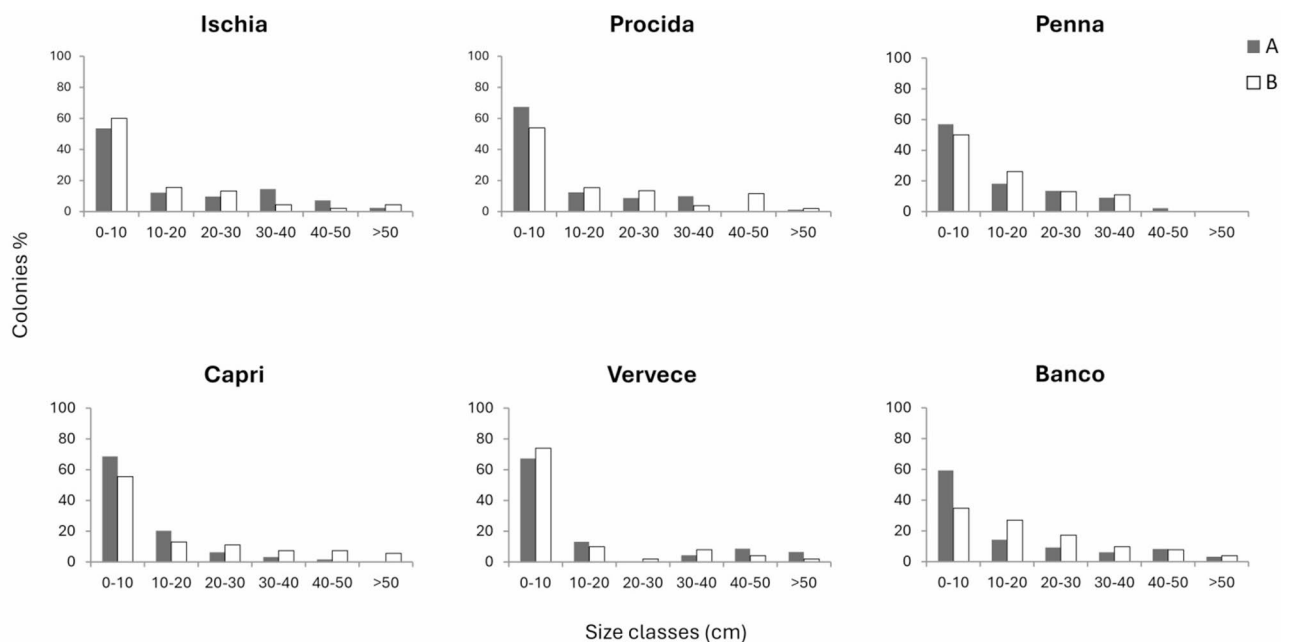


Fig. 2. Population structure. *Paramuricea clavata* population structure for site (A) (grey) and (B) (white) of assessed locations.

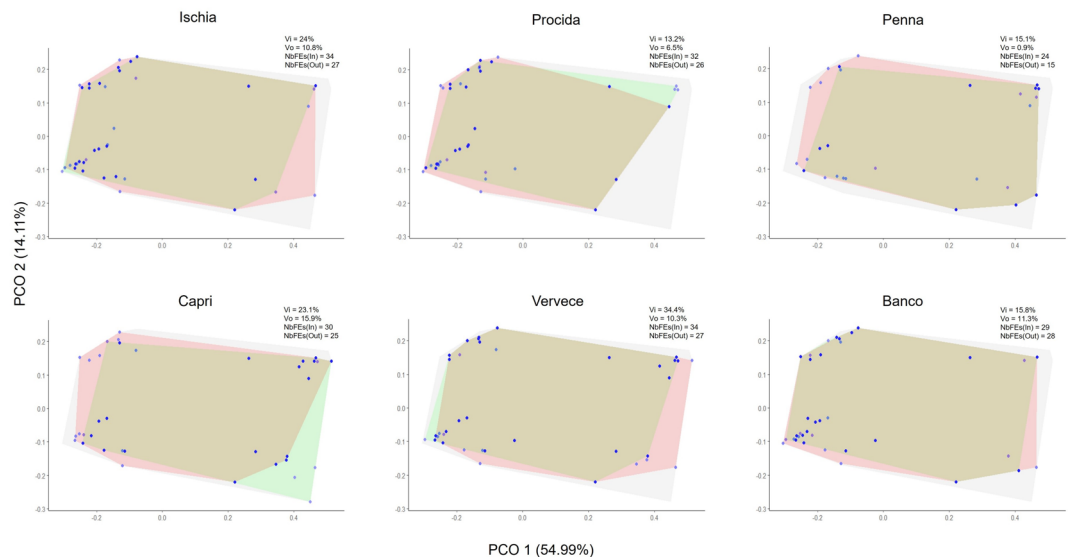


Fig. 3. Functional richness inside (red) vs. outside (green) the forests. The total functional space given by all FEs found in this study is represented as a grey polygon. The yellowish area corresponds to the overlap between the grey, red, and green polygons. Vi: volume inside the forest; Vo: volume outside. NbFEs(In): total number of FEs within the location inside the forest; NbFEs(Out): total number of FEs within the locality outside the forest. The axes PCO1 and PCO2 represent the first two dimensions of the 7D functional space and cumulatively explain 69.1% of the total variance.

that in almost all sites (11 out of 12) a distinct community structure inside vs. outside the forest was found (Table S5). Differences between conditions were mainly driven by the presence of invertebrates (e.g. *Smittina cervicornis*, *Crambe crambe*, *Pleraplysilla spinifera*, *Schizomavella mamillata*) inside the forests, while algal species (e.g. Dictyotales and the non-indigenous species *Caulerpa cylindracea* and *Lophocladia trichoclados*), mulilage and sediments were more abundant outside the forests (Fig. S3). A significant effect of the forests was also found on the functional structure of the assemblages in terms of composition and relative abundance of FEs, changing across sites ($p=0.001$) (Table S4), and a posteriori pairwise comparisons show significant differences in 10 out of the 12 sites included in the analysis (Table S5).

The multivariate analysis showed a significant effect of *P. clavata* forests on the CWM of traits but not consistently across sites ($p=0.012$) (Table S6). A posteriori pairwise comparisons indicated that the functional traits differed between conditions in 7 out of 12 sites (Table S7). The nMDS analyses carried separately for each site reported that colonial heterotrophs, with lecitotrophic larvae, physical and chemical defenses and non-photosynthetic pigments were more abundant within the forests, while low-longevity autotrophs that mainly use phycoerythrin pigments (e.g., rhodophytes) and reproduce via spores characterized communities outside the forests (Fig. S4).

Patterns of heterogeneity in the understory assemblages in presence and in absence of the forests

PERMDISP analysis revealed significant differences in community heterogeneity between the two conditions, both at the compositional and functional level ($p=0.001$) (Fig. 4). More specifically, a higher heterogeneity was observed inside the forests than outside for both compositional (0.683 ± 0.001 s.e.m. inside vs. 0.612 ± 0.002 s.e.m. outside) and functional (0.673 ± 0.001 s.e.m. inside vs. 0.610 ± 0.002 s.e.m. outside) analyses (Fig. 5), suggesting less variability in the patterns of distribution of benthic assemblages in absence of *P. clavata*. The breakdown into the two components revealed that turnover is the major component of the pattern of heterogeneity for both conditions. Even in this case, the turnover component of community heterogeneity was higher inside the forests than outside, both at the compositional (0.555 ± 0.002 s.e.m. inside vs. 0.486 ± 0.002 s.e.m. outside) and functional level (0.546 ± 0.002 s.e.m. inside vs. 0.482 ± 0.002 s.e.m. outside) (Fig. 5) and both differences were statistically significant ($p=0.01$) (Fig. 4). In other words, rather than having a gain or loss of species/functions going from one condition to another, communities associated with *P. clavata* compose a distinct and heterogeneous assemblage from both compositional and functional point of view. A summary of the results obtained from univariate and multivariate analyses is reported in Table 1.

Discussion

The results of our study document that the presence of *P. clavata* forests favors the development of distinct benthic assemblages, characterized by higher species and functional richness compared to areas where forests are absent.

In temperate regions, the decline and loss of habitat-forming species in response to multiple anthropogenic stressors is increasingly documented⁶, with indications of the drivers behind these changes and their consequences

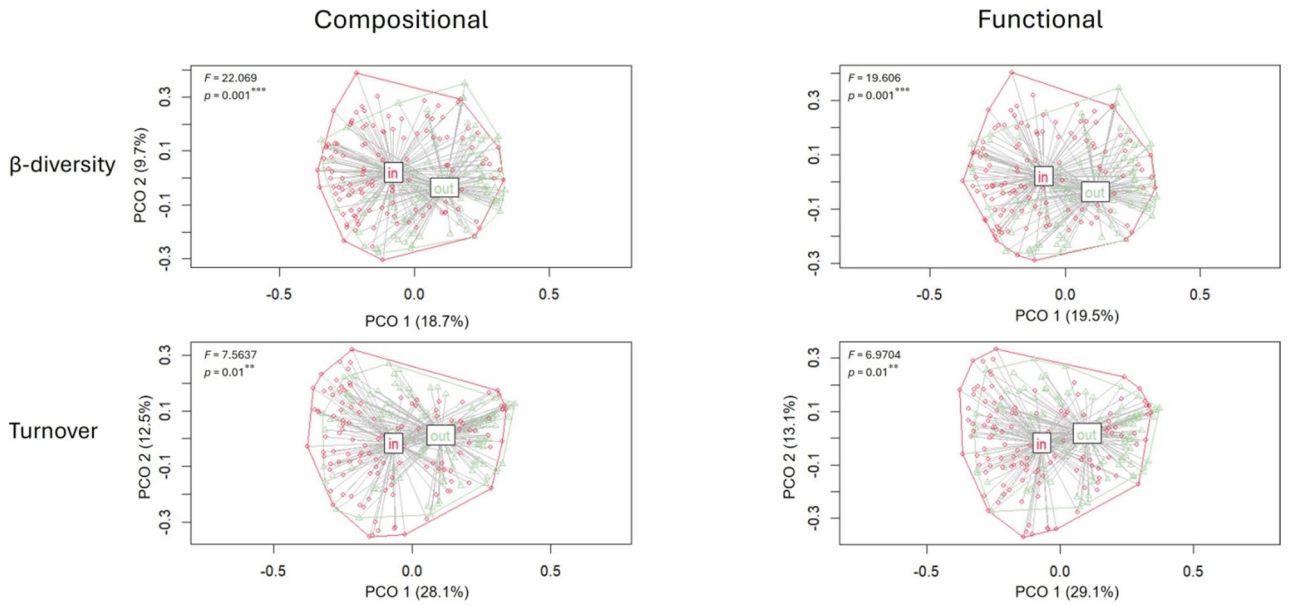


Fig. 4. Heterogeneity (PERMDISP) of the two conditions (In and Out). PCO based on Jaccard dissimilarity matrix of compositional and functional total β -diversity and the turnover component inside (red) vs. outside (green) the forests.

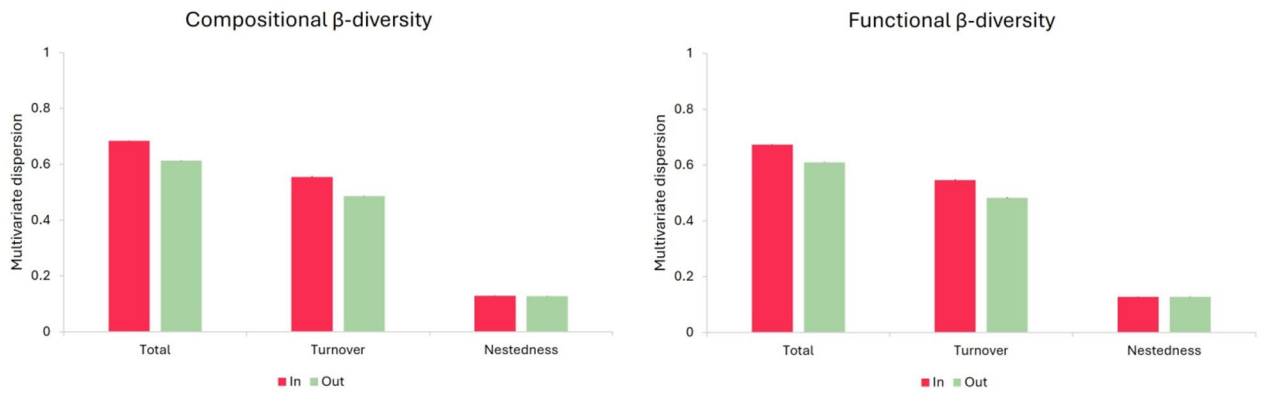


Fig. 5. β -diversity. Compositional and functional β -diversity within condition and their partitioning into the two components of turnover and nestedness. Error bars represent s.e.m.

	Sp	FEs	FRic	TS	FS	CWM
Lo						**
Co	*	**	*			
Si (Lo)	**	**	**			
Lo \times Co				*	*	
Si (Lo) \times Co				***	***	*
Transformation	None	None	None	FR	FR	FR

Table 1. Summary of statistical analysis results for each response variable. *Sp* species richness; *FEs* richness; *FRic* functional richness; *TS* taxonomic structure; *FS* functional structure; *CWM* Community-Weighted Mean; *FR* fourth root. Significant differences are shown as follow * $P < 0.05$. ** $P < 0.01$. *** $P < 0.001$.

in terms of species composition and relative abundance. Among habitat formers, marine animal forests are recognized as biodiversity hotspot for different communities including meiofauna, infauna, sessile and vagile species, epibionts and ichthyofauna^{7,69–73}. However, the limited quantitative knowledge about their distribution and functional role hampers our understanding of the underlying causes of their increasing loss and our ability to predict future changes, which could lead to regime shifts and alterations of associated biodiversity⁷⁴. By using fine-scale data, we described the *P. clavata* populations and investigated their role in shaping taxonomic and functional diversity. Within the assessed study areas, *P. clavata* populations were characterized by the dominance of small, non-reproductive colonies, and by significantly positive skewness and kurtosis values, indicating altered size structures that deviate from the bell-shaped patterns expected in undisturbed populations^{19,53}. In addition, the positive slope of the density–biomass relationship, which is typically negative in mature forests, suggests that the studied populations are likely experiencing ongoing stressors affecting subtidal assemblages¹⁹. The Gulf of Naples is indeed an urbanized coastal region, where multiple stressors such as fishing and climate-related events may affect the status of *P. clavata* forests^{13,75–77}.

Our study represents a baseline for this area of the Mediterranean Sea and expands current knowledge on biodiversity associated with these forests, which has been sparse and fragmented across very few Mediterranean regions, already documenting that the loss of *P. clavata* forests can lead to significant changes in recruitment patterns in the understory assemblages^{70,78} resulting in lower species diversity and richness^{9,41}. Systematic efforts and long-term monitoring focused on improving knowledge of marine animal forests and associated biodiversity are needed to quantitatively assess the status and the effects of different combinations of stressors across the basin. This information is particularly urgent, given that cnidarians are more affected than any other group by strong thermal anomalies, which are leading to mass mortality events at the Mediterranean scale¹⁶.

Our findings also show that the presence of the forest supports a different functional structure with more functional entities and higher functional richness and that results are consistent at both the scales of tens' meters and kilometres. A broader occupation of the functional trait space indicates a more diverse set of ecological roles and processes being supported under the forests. This role in influencing ecosystem functioning has also been documented for other marine animal forests such as coral reefs, sponge grounds, vermetid reefs, and oyster reefs, which regulate energy transformation and habitat provision by modifying local environmental variables⁷. Our functional analysis indicates that the ability of *P. clavata* to favor specific traits varies across sites. Spatial variability of local environmental factors and/or the different disturbance histories of assessed populations may explain these results. A fine-scale assessment of these drivers, within the framework of manipulative experiments, could help disentangle the effects of the structure and functioning of the forests from those of other local and global drivers influencing the system. However, some traits such as heterotrophy, coloniality and the presence of species with defined physical and chemical defenses are typically prevalent within the forests. Outside the forests, fast-growing and low-longevity primary producers replace heterotrophs, possibly altering ecological processes involved in energy fluxes, such as productivity and benthic–pelagic coupling, with cascade effects across biodiversity levels⁷⁹. This functional shift may affect important ecosystem services supported by marine animal forests, including carbon sequestration, food provision, genetic and medicinal resources, erosion prevention, and tourism^{7,10,80–83}. These patterns correspond to those reported for *P. clavata* and *Corallium rubrum* before and after the occurrence of marine heat waves, when fast-growing autotrophic species (e.g. algal turf, *Caulerpa cylindracea*) rapidly colonized the free spaces after mass mortality events at the expense of morphologically complex and long-lived heterotrophs³³. This evidence highlights the importance of marine animal forests in influencing environmental variables, creating unique habitat conditions that selectively favor the settlement of certain species^{25,84}. Additionally, the physical structure of *P. clavata* colonies has been observed to act as a filter for mucilage accumulation and to reduce water flow, limiting sediment resuspension and deposition^{26,78}. A higher abundance of mucilage, sediments and of non-indigenous species *L. trichocladus* and *C. cylindracea* was recorded outside *P. clavata* populations. This function is vital for preventing the establishment of invasive species such as *Caulerpa cylindracea*, which thrives in sediment-rich and disturbed conditions^{37,78,85}. Nevertheless, while the presence of *P. clavata* protects the associated community from the spread of the mucilage and the resulting negative aspects on the community itself, mucilage that remains attached to the colony can induce necrosis, generating a diffused oxidative stress in the entire *P. clavata* colony and affecting its physiological processes⁸⁶. The results, so far, are also in accordance with the biotic resistance hypothesis, which states that more diverse communities are more resistant to invasion due to the complementary use of resources by natives (complementary effects) or the higher probability of including highly competitive native species which limit the use of resources by invaders^{87,88}.

The specific environmental conditions created by *P. clavata* are also likely to drive the high heterogeneity observed within the forest, as it has been documented for other habitats dominated by different gorgonian species (i.e. *Acanthogorgia hirsuta*, *Swiftia pallida*, *Isidella elongata*, *Eunicella verrucosa*)⁴⁵. Our β -diversity analysis contributed to identifying *P. clavata* forests as hot-spots of temperate reef biodiversity, with greater total compositional and functional β -diversity inside the forest compared to adjacent zones. Outside the forests, the less heterogeneous patterns of distribution found include both compositional and functional β -diversity. Here, biotic homogenization could be driven by the presence of stressors such as invasive species, sediments and mucilage, leading to species loss and decrease of more vulnerable species (e.g. *Myriapora truncata*, *Smittina cervicornis*)^{89,90}. Thus, areas outside the forest reflect the characteristics of more disturbed conditions than assemblages associated with *P. clavata*, featured by the gain of species and functions which foster the recovery or resistance to disturbances like biological invasions^{91,92}. The breakdown into the two components revealed that species replacement dominates both compositional and functional β -diversity. Turnover in species composition translates into functional turnover when communities have low functional redundancy (i.e. low number of species performing similar functions)⁹³. The positive correlation between these complementary aspects of β -diversity may be related to the high variability of environmental conditions, which leads species to a differential

partitioning of resources, resulting in a high turnover of species performing different functions. This pattern has also been observed in shallow subtidal habitats, where environmental factors (e.g. temperature, light exposition, hydrodynamism) are highly variable and likely drive the species sorting⁴⁷. In contrast, environmental homogenization can induce functional nestedness even though compositional turnover represents the dominant component^{47,93}. In our study, the increase of the compositional and functional turnover inside the forests suggests that the presence of *P. clavata* drives the replacement of species and functions⁹⁴, possibly arising from differences in niche features between the two conditions³⁴. The high habitat complexity created by the forests allows the colonization by organisms with different ecological needs, compared to more simple habitats⁹⁵. Therefore, preserving the integrity of the *P. clavata* populations is crucial to ensure the maintenance of the habitat complexity which in turn is vital to support compositional and functional diversity. Extended time-series data would strengthen our results and further confirm the ecological role of the species in shaping local biodiversity over the long term and under global change scenarios.

The results of our study highlight the need to improve conservation efforts for *P. clavata* forests in order to protect their important role in enhancing not only taxonomic diversity but also ecosystem functioning, the latter often being neglected in conservation planning. Currently, only 18% of *P. clavata* potential habitat at the Mediterranean scale is under protection regimes²². Considering the increasing pressures related to climate change, local threats need to be removed to limit cumulative and synergistic negative impacts on animal forests and associated assemblages³⁰. Present efforts to achieve the 30% conservation target set by the new EU Biodiversity Strategy for 2030 should translate into specific strategies for the inclusion of *P. clavata* forests within Marine Protected Areas⁹⁶, an effective tool to protect this habitat and enhance its resilience⁹⁷. Quantitative information and fine-scale data on the distribution and status of *P. clavata* are also relevant for the Nature Restoration Law (NRL). The NRL constitutes the EU's long-term strategy to restore biodiversity and ecosystem services over the next decades⁹⁸, recently approved by the EU with the objective to halt and reverse biodiversity loss. However, the NRL sets very ambitious quantitative targets in terms of both the areas to restore and the timeframe for their restoration, considering the current poor knowledge about the distribution and status of several species and habitats across the EU, including *P. clavata* forests. Achieving these targets requires urgent development of knowledge, methodologies, tools, and best practices to monitor progress and ensure success. This work addresses these gaps to consolidate existing knowledge for a successful implementation of EU Directives.

Data availability

All data generated or analysed during this study are included in this published article (and its supplementary information files).

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Author contributions

S.F. and Al.Co. conceived the study; Al.Co., L.L., An.Ch., S.D., M.M., S.M.S.M., C.S. participated in the field work; L.L., Al.Co., S.F. and E.F. performed the data analyses; S.F., Al.Co. and L.L. led the writing of the manuscript with the contributions from E.F., An.Ch., S.D., M.M., S.M.S.M., C.S., P.S.; Al.Co. and L.L. are authors with equal contribution.

Declarations

Competing interests

The authors declare no competing interests.

Additional information

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