

REDUCING THE DATA-DEFICIENCY OF THREATENED EUROPEAN HABITATS:
SPATIAL VARIATION OF SABELLARIID WORM REEFS AND ASSOCIATED FAUNA
IN THE SICILY CHANNEL, MEDITERRANEAN SEA

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1 ABSTRACT

2 Biogenic reefs, such as those produced by tube-dwelling polychaetes of the genus
3 *Sabellaria*, are valuable marine habitats which are a focus of protection according to European
4 legislation. The achievement of this goal is potentially hindered by the lack of essential
5 empirical data, especially in the Mediterranean Sea. This study addresses some of the current
6 knowledge gaps by quantifying and comparing multi-scale patterns of abundance and
7 distribution of two habitat-forming species (*Sabellaria alveolata* and *S. spinulosa*) and their
8 associated fauna along 190 km of coast on the Italian side of the Sicily Channel. While the
9 abundance of the two sabellariids and the total number of associated taxa did not differ at any
10 of the examined scales (from tens of centimetres to tens-100 of kilometres), the structure
11 (composition in terms of both the identity and the relative abundance of constituting taxa) of
12 the associated fauna and the abundance of several taxa (the polychaetes *Eulalia ornata*, *Syllis*
13 *pulvinata*, *S. garciai*, *Nereis splendida* and *Arabella iricolor*, and the amphipods *Apolochus*
14 *neapolitanus*, *Tethylembos viguieri* and *Caprella acanthifera*) varied among locations
15 established ~50-100 km apart. *Syllis pulvinata* also showed significant variation between sites
16 (hundreds of metres apart), analogously to the other syllid polychaetes *S. armillaris* and *S.*
17 *gracilis*, the nereidid polychaete *Nereis rava*, and the amphipod *Gammaropsis ulrici*. The
18 largest variance of *S. spinulosa*, of the structure of the whole associated fauna and of 56% of
19 taxa analysed individually occurred at the scale of replicates (metres apart), while that of the
20 dominant bio-constructor *S. alveolata* and of 25% of taxa occurred at the scale of sites. The
21 remaining 19% and the total richness of taxa showed the largest variance at the scale of
22 locations. Present findings contribute to meet a crucial requirement of any future effective
23 protection strategy, i.e., identifying relevant scales of variation to be included in protection
24 schemes aiming at preserving representative samples not only of target habitats and organisms,
25 but also of the processes driving such variability.

26

27 *Keywords:* Biogenic reef; ecosystem engineer; endangered habitat; *Sabellaria alveolata*;
28 *Sabellaria spinulosa*; spatial scale; species interaction; subtidal

29

30 **1. Introduction**

31 Organisms known as bio-constructors produce three-dimensional structures providing
32 habitats to a large diversity of sessile and mobile species (e.g., Jones et al., 1994). Besides the
33 most notable case represented by cnidarians, such as hermatypic corals (Sheppard et al., 2009),
34 several other animal taxa have such an ability, including sponges (Bell et al., 2013), molluscs
35 (Gutiérrez et al., 2003), polychaetes (Naylor and Viles, 2000) and bryozoans (Wood et al.,
36 2012). Gregarious tube-dwelling worms of the Sabellaridae family (Annelida), such as for
37 example *Phragmatopoma californica* in California and *Gunnarea capensis* in South Africa,
38 can form, provided suitable environmental conditions, extensive reefs (Achary, 1974).
39 According to the Habitats Directive definition, reefs can be defined as “*submarine, or exposed*
40 *at low tide, rocky substrates and biogenic concretions, which arise from the sea floor in the*
41 *sublittoral zone but may extend into the littoral zone where there is an uninterrupted zonation*
42 *of plant and animal communities. These reefs generally support a zonation of benthic*
43 *communities of algae and animal species including concretions, encrustations and*
44 *corallogenic concretions”* (European Commission DG Environment, 2003). Along the
45 European coasts, the honeycomb worms *Sabellaria alveolata* and *S. spinulosa* form the most
46 important sabellariid bio-constructors. These can develop in the intertidal and shallow
47 subtidal zone as encrusting formations adhering to rocks, ball-shaped (‘hummocks’) structures
48 or large banks (Wilson, 1971; Dubois et al., 2002, 2006; Desroy et al., 2011), whose ‘reefiness’
49 can be quantified based on a number of physical and biological characteristics (Hendrick and
50 Foster-Smith, 2006). Sabellariid constructions are not only found on natural substrates, but also
51 on artificial structures (Pearce et al., 2014; Firth et al., 2015).

52 The ecological importance of sabellariid bio-constructions is widely acknowledged and is
53 primarily due to their modulation of crucial abiotic and biotic processes, including substrate
54 stabilization, sediment trapping and the provision of structurally complex microhabitats used
55 for refuge, nursery and feeding by associated, mainly macrofaunal, organisms (Holt et al.,
56 1998; Dubois et al., 2002, 2006; Cocito, 2004; Plicanti et al., 2017). Sabellariid reef-associated
57 organisms span a wide range of taxa, although the largest proportion is typically represented by
58 other polychaetes (Dias and Paula, 2001; Dubois et al., 2006; Cole and Chapman, 2007). The
59 majority of species are common in other benthic habitats, but species using sabellariid reefs as
60 their exclusive or preferential habitat are also known (Killeen and Light, 2000; Schimmenti et
61 al., 2016) and contribute to make the *Sabellaria*-associated macrofauna generally more diverse
62 and abundant than that of surrounding areas (Desroy et al., 2011). This macrofauna is also
63 peculiar as it is represented, in the same habitat, by species typically found on hard, muddy or
64 sandy bottom (Dubois et al., 2002).

65 *Sabellaria* reefs, however, represent highly dynamic systems exposed to several threats,
66 including natural perturbations and increasing disturbance directly and/or indirectly related to
67 human activities, which can affect both the habitat-forming species and their associated
68 organisms. For instance, activities such as oyster farming and fishing may cause critical
69 damage to the structure of both mature and developing reefs (Dubois et al., 2002, 2006, 2007).
70 Even the physical disturbance produced by low intensities of trampling was experimentally
71 indicated as responsible for significant reductions in the cover of intact *S. alveolata* concretions
72 (Plicanti et al., 2016). Concomitantly, biogenic reefs are subject to physical damage by
73 meteorological events like extreme storms, precipitation, temperature and acidification. The
74 intensity, frequency and variance of such events was demonstrated by empirical data to
75 increase in the last decades (e.g. Muller and Stone, 2001; Wolff et al., 2016) and is modelled to
76 further increase in the near future due to climate change (Webster et al., 2005; Sriver and
77 Huber, 2006; Trapp et al., 2007). The tube-building activity of *Sabellaria* may effectively

78 repair small damaged areas of the reef within weeks or months (Cunningham et al., 1984;
79 Vorberg, 2000; Plicanti et al., 2016), but heavier impacts may take years up to decades (Firth et
80 al., 2015) to recover depending on temporally variable processes, such as larval supply (Gruet,
81 1986; Ayata et al., 2009). Permanent changes to the structure of the reef are also reported
82 (Gibb et al., 2014).

83 The functional role played and the diversity and severity of threats they are subject to
84 render sabellariid reefs a very valuable and vulnerable marine habitat, that must be considered
85 for protection by European Union (EU) legislation, including the Habitats Directive (Directive
86 92/43/EEC), the Water Framework Directive (Directive 2000/60/EC) and the Marine Strategy
87 Framework Directive (Directive 2008/56/EC). Moreover, *Sabellaria* reefs are included in the
88 European Red List of Habitats (Gubbay et al., 2016) that is intended to provide an overview of
89 the type, extent and status of benthic marine habitats across Europe. Unfortunately, these reefs,
90 especially in the Mediterranean Sea, are listed as ‘Data Deficient’, thus needing comprehensive
91 and specific investigation. In particular, *Sabellaria* reefs along the Italian coast are reported to
92 cover thousands km² (IUCN, 2016), but largely lack empirical and quantitative data. Previous
93 studies in the Mediterranean region focused on very local scales and a few morphological and
94 ecological traits (Porrás et al., 1996; La Porta and Nicoletti, 2009; Lezzi et al., 2015;
95 Schimmenti et al., 2016), providing insufficient and fragmentary information to identify the
96 current status, distribution, dynamics and threats of *Sabellaria* reefs. Such information, instead,
97 is essential to assess their conservation needs and to guide effective management and
98 protection strategies. This is especially true for the urgently needed knowledge on patterns of
99 distribution of reefs and their associated biodiversity (but see Cole and Chapman, 2007). In
100 fact, identifying the relevant scales of variation of organisms is an essential requirement to
101 design protection schemes able to preserve not only those populations directly, but also their
102 driving processes (García-Charton et al., 2000; Benedetti-Cecchi et al., 2003; Frascchetti et al.,
103 2005a; Micheli et al., 2005; Bertocci et al., 2012; Anderson et al., 2014).

104 The present study was aimed at contributing to fill the above illustrated gaps by estimating
105 and comparing spatial patterns of distribution, abundance and diversity of sabellariid bio-
106 constructors and their associated fauna from biogenic reefs occurring in the shallow subtidal
107 environment along the Italian coast of the Sicily Channel (central Mediterranean Sea).
108 Specifically, we tested the hypothesis that patterns of the abundance, size and spatial variance
109 of reef-forming polychaetes (*S. alveolata* and *S. spinulosa*), of the overall structure (identity
110 and relative abundance of constituting taxa), richness and Shannon's diversity index of the
111 associated faunal assemblage, and of the abundance and variance of the main faunal taxa
112 differed at each of three scales ranging from tens of centimetres to ~200 km.

113

114 **2. Methods**

115 *2.1. Study system*

116 The Sicily Channel lies between the southern coast of Sicily (Italy) and the north-eastern
117 coast of Tunisia, separating the Western from the Eastern Mediterranean sub-basins. As such, it
118 is directly affected by the Atlantic Ionian Stream, which flows eastwards from the Atlantic
119 Ocean through the Strait of Gibraltar (Poulain and Zambianchi, 2007). The main
120 oceanographic conditions, coastal morphology and prevailing habitats in the study region are
121 described in detail elsewhere (Schimmenti et al., 2016). Briefly, the present study was carried
122 out at three locations (Triscina = TRI: 37°58' N, 12°80' E; Eraclea Minoa = ERA: 37°39' N,
123 13°28' E; Donnalucata = DON: 36°75' N, 14°64' E) distributed along about 190 km of coast in
124 southern Sicily, which is characterized by a prevailing sandy bottom interspersed with
125 meadows of the seagrass *Posidonia oceanica*, rocky shores and sabellariid reefs. A map of the
126 study area illustrating the sampling locations is given in Appendix A.

127 The sampled reefs had a pillow-like form with a brain-like structure, ranging between 0.5
128 and 2 m in diameter and between 0.3 and 1.5 m in height and occurred between 1.5 and 3 m in
129 depth depending on the location. All reefs occurred on a sandy bottom, interspersed with

130 pebbles and close to a *P. oceanica* meadow at Triscina, and located at the base of a rocky
131 breakwater at Donnalucata (for further details, see Schimmenti et al., 2016).

132

133 2.2. Sampling design and collection of data

134 Data were collected, by SCUBA divers, in June 2013 at two sabellariid reefs (hundreds of
135 metres apart, hereafter indicated as sites) chosen at random in each location. Within each site,
136 four replicate blocks (0.1 x 0.1 x 0.1 m, some metres apart) were collected at random using a
137 hammer and a putty knife and immediately preserved, in separate bags, in 70% ethanol for
138 subsequent analysis in the laboratory. In order to avoid edge effects due to the occurrence of
139 different habitats adjacent to the sampled reefs at each location, all blocks were collected from
140 the centre of each reef.

141 In the laboratory, all individuals of each *Sabellaria* species and of the associated fauna
142 were sorted under a stereomicroscope, identified to the finest possible taxonomic level (species
143 in most cases) and counted.

144 Moreover, the opercular length (maximum width of the operculum), considered as a proxy
145 of body size and, hence, of the home-tube size, was measured on 40 *S. alveolata* (the dominant
146 habitat-forming polychatete) individuals selected at random among those found in each of two
147 replicate blocks chosen at random among the four available in each site and location.

148

149 2.3. Analysis of data

150 A two- way permutational multivariate analysis of variance (PERMANOVA: Anderson,
151 2001) based on Bray-Curtis untransformed dissimilarities was used to examine differences in
152 the structure of the faunal assemblages associated with the *Sabellaria* habitat-forming
153 polychaetes among locations and between sites. The model for the analysis included the factors
154 ‘Location’ (random, three levels) and ‘Site’ (random, two levels, nested within Location), with
155 four replicates provided by the blocks collected in each site. Since Bray-Curtis dissimilarities

156 combine differences in both the identity and the relative abundance of taxa, the same analysis
157 was repeated using either the matrix of raw abundances, or that of presence/absence data as
158 input.

159 Multivariate patterns were visually displayed using Principal Component Ordination (PCO)
160 based on Bray-Curtis dissimilarities calculated on both the abundance and the
161 presence/absence data matrix. The SIMPER procedure (Clarke, 1993) was adopted to quantify
162 the absolute (δ_i) and the percent ($\delta_i \%$) contribution of each taxon to the total dissimilarity
163 between paired locations, using a cut-off of 75% of cumulative dissimilarity for excluding low
164 contributions. Data on the abundance (number of individuals) of each *Sabellaria* species, their
165 sum, the total number and the Shannon's diversity index (Shannon, 1948) of associated taxa,
166 and the abundance of each taxon identified as relevant by SIMPER were analysed with analysis
167 of variance (ANOVA) based on the same design as that used for PERMANOVA.

168 The opercular length of *S. alveolata* was analysed with a three-way ANOVA model
169 including the same two factors of the previously described analyses, plus a third factor
170 'Replicate' (random, two levels represented by two replicate blocks in each site). The 40
171 individuals selected in each replicate block provided the replicates for this analysis.

172 Before each ANOVA, the assumption of homogeneity of variances was assessed with
173 Cochran's *C* test, and data were log-transformed when the assumption was not met. When
174 heterogeneous variances could not be stabilised by transformation, untransformed data were
175 analysed and results were considered robust if not significant ($p > 0.05$) or significant at $p < 0.01$.
176 These options were considered appropriate since the probability of Type II error is not affected
177 by heterogeneous variances, and to compensate for increased probability of Type I error,
178 respectively (Underwood, 1997).

179 Univariate variance components and their multivariate analogous at each scale were
180 calculated, respectively, from ANOVA and PERMANOVA by equating empirical and
181 expected mean squares (Winer et al., 1991; Underwood, 1997). Negative estimates of variance

182 were interpreted as sample underestimates of very small to null variances and set to zero
183 (Searle et al., 1992; Underwood, 1997). All variance components were calculated from
184 untransformed data (e.g. Terlizzi et al., 2005).

185

186 **3. Results**

187 *3.1. Habitat-forming species, richness and structure of associated faunal assemblages*

188 *Sabellaria alveolata* and *S. spinulosa* coexisted in the sampled reefs, but the first species
189 represented about 99% of all habitat-forming polychaetes and thus drove the overall patterns of
190 abundance of total *Sabellaria* specimens counted throughout the study (Fig. 1 A-C). The
191 abundance of the two *Sabellaria* species together and that of *S. alveolata* showed large and
192 significant variation between sites, but not among locations (Table 1 and Table 2). Both
193 species, however, differed in abundance between sites at Triscina more than at Eraclea Minoa
194 and Donnalucata, with the first location showing a non-significant trend of larger abundances
195 compared to the other two locations (Fig. 1 A and B). The range of *S. alveolata* opercular
196 length was between 0.5 and 5 mm, with a mean (± 1 SE) of 2.04 (± 0.04) mm calculated over all
197 sampled individuals ($n = 480$). Significant variation in this variable was found only between
198 replicate blocks, although more than 83% of the total variance was provided by the among-
199 individuals scale (Table 1 and Table 2 in Appendix B). In spite of graphically evident
200 differences between sites within each location, the abundance of *S. spinulosa* also did not vary
201 significantly, neither between sites nor among locations (Table 1 and Fig. 1 C), showing most
202 variation at the among-replicates scale (Table 2).

203 In total, 86 faunal taxa (*Sabellaria* spp. excluded) associated with biogenic reefs, including
204 43 Annelida, 42 Arthropoda (27 Amphipoda and 14 Decapoda among Crustacea, and 1
205 Arachnida) and 1 Echinodermata (Ophiuroidea), were identified in this study (Appendix C)
206 and used to calculate the total richness of taxa (a proxy for species richness) and in the
207 multivariate analysis. Most taxa ($n=67$) were identified to species level, while the remaining 19

208 were identified to genus (n=12), family (n=5) or higher taxa (n=2). The total number of taxa
209 was comparable across sites and locations (Table 1 and Fig. 1 D), with about 98% of the total
210 variance provided almost equally by the replicate and the location scale (Table 2). The
211 Shannon's diversity index varied significantly among locations, with a decreasing trend from
212 Triscina to Donnalucata to Eraclea Minoa (Table 1 and Fig. 1 E) and most variance provided
213 by the location scale (Table 2).

214 The structure of the reef-associated assemblages varied at each examined scale
215 independently of analysing abundance or presence/absence data (Table 3). In both cases, most
216 multivariate variation occurred at the among-replicates scale, followed by the among-locations
217 and the between-sites scales (Table 2). The PCO clearly separated the three locations and the
218 two sites within each location in terms of dissimilarities calculated from both abundance and
219 presence/absence data (Fig. 2 A and B).

220

221 3.2. Faunal taxa associated with sabellarid reefs

222 The SIMPER procedure identified 11, 13 and 7 taxa as collectively contributing with over
223 75% to the total dissimilarity of Triscina vs. Eraclea Minoa, Triscina vs. Donnalucata, and
224 Eraclea Minoa vs. Donnalucata, respectively (Table 1 in Appendix D). Out of the total 17 taxa
225 identified as either exclusive of one of the three contrasts or shared by multiple contrasts, all
226 but Sipunculidae were analysed individually.

227 Significant differences among locations were found for 8 species, namely the polychaetes
228 *Eulalia ornata*, *Syllis pulvinata*, *Syllis garciai*, *Nereis splendida* and *Arabella iricolor*, and the
229 amphipods *Apolochus neapolitanus*, *Tethylembos viguieri* and *Caprella acanthifera* (Tables 1-
230 3 in Appendix E). *Eulalia ornata* was, on average, the most abundant species and was
231 commonly found at all sites and locations, although in higher numbers at Eraclea Minoa than at
232 both Triscina and Donnalucata (Fig. 3 A). Differently, *S. pulvinata* (Fig. 3 B), *N. splendida*
233 (Fig. 3 E), *A. iricolor* (Fig. 3 H), *A. neapolitanus* (Fig. 4 E) and *C. acanthifera* (Fig. 4H) were

234 relatively more abundant at Donnalucata and very rare, if not completely absent (i.e., *A.*
235 *iricolor* and *C. acanthifera* at Eraclea Minoa), at the other locations. All these species tended to
236 vary in abundance also between sites, but significant differences at such scale were found for *S.*
237 *pulvinata* only (Table 1 in Appendix E and Fig. 3 B). *Syllis garciai* (Fig. 3 I) and *T. viguieri*
238 (Fig. 4 G), instead, were mostly found at Triscina, less abundant at Donnalucata and absent at
239 Eraclea Minoa.

240 Adding to *S. pulvinata*, four more species showed significant variation between sites
241 (Tables 1-3 in Appendix E), including the polychaetes *Syllis armillaris* (Fig. 3 C), *Syllis*
242 *gracilis* (Fig. 3 D) and *Nereis rava* (Fig. 3 G), and the amphipod *Gammaropsis ulrici* (Fig. 4
243 B). In some cases, however, between-sites variability might have masked some visually
244 evident differences among locations, such as in *S. armillaris* and *G. ulrici*, which tended to be
245 relatively more abundant at Triscina, and in *S. gracilis*, which was represented in higher
246 numbers at Eraclea Minoa (Fig. 3 C, Fig. 4 B and Fig. 3 D, respectively).

247 Four taxa did not show significant results (Tables 1-3 in Appendix E), including the
248 polychaete *Brania arminii* (Fig. 3 F) and the amphipods *Jassa ocia* (Fig. 4 C), *Caprella rapax*
249 (Fig. 4 D) and *Quadrimeaera* sp. (Fig. 4 F). All of them were generally represented by low
250 numbers, with *B. arminii* and *C. rapax* being absent at Eraclea Minoa, and *Quadrimeaera* sp.
251 being absent at Donnalucata (Fig. 3 F, Fig. 4 D and Fig. 4 F, respectively).

252 Sipunculidae were only present at Triscina and were not analysed statistically (Fig. 4 A).

253 Finally, the comparison of variance components of *Sabellaria*-associated taxa analysed
254 individually indicated the smallest examined scale, i.e. among-replicates, as that providing the
255 largest contribution in the majority (56%) of cases. These included, *N. splendida*, *B. arminii*, *A.*
256 *iricolor*, *J. ocia*, *C. rapax*, *A. neapolitanus*, *Quadrimeaera* sp., *T. viguieri* and *C. acanthifera*
257 (Table 2). Only four (25%, represented by *S. armillaris*, *S. gracilis*, *N. rava* and *G. ulrici*) and
258 three (19%, represented by *E. ornata*, *S. pulvinata* and *S. garciai*) species showed the largest
259 variance at the site and the location scale, respectively (Table 2).

260

261 **4. Discussion**

262 This study tested the main hypothesis that patterns of distribution of habitat-forming
263 sabellariid polychaetes (the dominant *S. alveolata* and the much less numerous *S. spinulosa*)
264 and of the overall structure and individual abundance of associated fauna in the Sicily Channel
265 varied at spatial scales ranging from metres (among replicates) to ~50-100 km (among
266 locations). Results indicated that relevant scales of variation were not consistent across all
267 examined organisms. While the abundance of total *Sabellaria* and of the dominant *S. alveolata*
268 varied significantly at the site, but not the location, scale, the structure of the associated
269 assemblages and the abundance of many of the most common taxa showed marked differences
270 at the largest scale. This observation is rather surprising since it could be expected that the
271 patterns of distribution of bio-constructors are matched by those of the faunal taxa using
272 biogenic reefs as habitat. Moreover, obvious macroscopic differences in the general reef
273 structure, size, surrounding habitats and depth among sampled locations (as illustrated in detail
274 by Schimmenti et al., 2016) did not correspond to significant differences in patterns of
275 distribution and abundance of sabellariid worms at the same scale. The present findings
276 suggest that the structure of the habitat and the surrounding environment, although not
277 quantified in this study, may actually shape the structure and distribution of associated
278 assemblages, but that this effect can be decoupled from the mere abundance of the habitat-
279 forming species. An analogous observation was made by Schlund et al. (2016), who reported
280 that different structures (platforms vs. 'reefs', i.e., more massive formations) of *S. alveolata*
281 bio-constructions in the Bay of Mont-Saint-Michel were characterized by different
282 macrofaunal assemblages, yet comparable density of *S. alveolata*. Such differences are often
283 explained with critical changes in the structural complexity of the bio-construction, which
284 would provide different ranges, types and extent of microhabitats to diverse organisms (Gruet,
285 1986; Porras et al., 1996; Dubois et al., 2002). All present reefs, however, were characterized

286 by a comparable convoluted tridimensional structure (detailed pictures are shown in
287 Schimmenti et al., 2016), suggesting that such mechanisms were relatively less important in
288 the studied system.

289 Local environmental conditions, instead, may be more relevant for determining location-
290 scale differences in the structure, diversity and abundance of the reef-associated fauna. A main
291 driving factor could involve the colonization of different organisms from surrounding habitats
292 (Dubois et al., 2006). All the sampled locations, in particular, presented sabellariid reefs
293 interspersed within a predominant sandy bottom, but these were close to a *P. oceanica* meadow
294 at Triscina and to a breakwater rocky barrier at Donnalucata, i.e., markedly different habitats
295 that are likely to provide different pools of colonizers to adjacent *Sabellaria*-associated
296 assemblages. For example, shallow hard substrates in the Mediterranean Sea are reported to
297 host diverse and abundant assemblages of syllid polychaetes (Chatzigeorgiou et al., 2012;
298 Musco, 2012; Dorgham et al., 2014), a feature that could have contributed to the larger
299 abundance of *S. pulvinata*, the most numerous syllid in this study, at Donnalucata compared to
300 Triscina. Concomitantly, significant among-locations variation in the distribution of reef-
301 associated fauna could also be driven by patterns of dispersal of organisms varying over the
302 same scale (e.g., Johnson et al., 2001; Frascetti et al., 2005b). This explanation is made
303 plausible by the potential effect of the Atlantic Ionian Stream originating from the Atlantic
304 Ocean, entering the Mediterranean Sea through the Strait of Gibraltar and reaching the surface
305 of the Sicily Channel in its eastward path. Such an oceanographic feature is associated with
306 environmental heterogeneity in terms of salinity, temperature (Lermusiaux and Robinson,
307 2001) and productivity (Ciappa, 2009) along the southern coast of Sicily. In addition,
308 transportation through the Atlantic Ionian Stream is likely to produce a gradient of distribution
309 of drifting colonizers which could have resulted in the observed variation among locations
310 tens-100 km apart along about 200 km of coast. On the contrary, other factors potentially
311 relevant for directly and indirectly driving differences in reef-associated assemblages, such as

312 the tidal height and wave exposure (e.g. Schlund et al., 2016, and references therein), were less
313 likely in the present system given the comparable depth and exposure to south-western winds
314 of all sampled reefs.

315 The peculiar position of the Sicily Channel at the separation between the Western and the
316 Eastern Mediterranean sub-basins may also be responsible for the local coexistence of two
317 *Sabellaria* species, although with a clear dominance of *S. alveolata*. So far, the studies
318 available for the Mediterranean Sea suggested that only *S. alveolata* and only *S. spinulosa*
319 build-up reefs in the Western and the Eastern (Adriatic Sea included) sub-basins, respectively
320 (La Porta and Nicoletti, 2009; Lezzi et al., 2015; Gubbay et al., 2016). Moreover, the present
321 locations are the southernmost records of the overall distribution of *S. alveolata* and *S.*
322 *spinulosa* reefs. Upwelling events occurring along the southern coast of Sicily (e.g. Piccioni et
323 al., 1988) may be responsible, at least in part, for such biogeographic patterns. In non-
324 Mediterranean regions, the majority of studies reported the exclusive occurrence of *S. alveolata*
325 (e.g., Wilson, 1976; Dubois et al., 2002; Plicanti et al., 2016) or *S. spinulosa* (e.g., Hendrick
326 and Foster-Smith, 2006), with the first species normally being more common and abundant in
327 the intertidal environment (e.g., Firth et al., 2015), while the second in the subtidal
328 environment (e.g., Pearce et al., 2014). In some cases, coexisting *S. alveolata* and *S. spinulosa*
329 have been recorded (e.g., Wilson, 1970a, 1970b), but not at the small scale documented in the
330 Sicily Channel.

331 Despite specific patterns of significant or non-significant differences at each examined
332 scale, the structure of the reef-associated assemblages and the abundance of the majority of
333 their constituting species showed the highest variance at the smallest (i.e., among replicates)
334 scale. Recurrent analogous findings suggest that relevant variation at spatial scales of tens-
335 hundreds of centimetres up to tens-hundreds of metres is a common feature of a wide range of
336 benthic organisms, habitats and geographic areas (Underwood and Chapman, 1996; Kendall
337 and Widdicombe, 1999; Benedetti-Cecchi, 2001; Frascchetti et al., 2005b; Terlizzi et al., 2007;

338 Bertocci et al. 2012), often explained primarily with the effects of the topography of the
339 substratum (e.g. Denny et al., 2004). In sabellariid reefs, small-scale heterogeneity could be
340 due to the availability of different microhabitats formed by worms, hence tubes, of a range of
341 sizes. The present data did not show significant differences in the opercular size of *S.*
342 *alveolata*; nevertheless, most variance in this trait occurred at small scales (among individuals
343 within the same block and between replicate blocks), suggesting that the heterogeneous spatial
344 pattern of available microhabitats could have maintained an analogously heterogeneous
345 distribution of the associated fauna. However, a previous comparison of patterns of distribution
346 of annelids associated with *S. alveolata* reefs from Wales and Portugal indicated that the
347 identification of small-scale variability can also depend on the level of taxonomic resolution,
348 being less likely for fine taxonomic groups, such as species and families, and more evident for
349 broad taxonomic groups, such as classes and phyla (Cole and Chapman, 2007). Instead, other
350 processes operating at a larger scale (hundreds of metres, i.e., between sites) may be invoked to
351 explain the large variance of *S. alveolata*. For instance, a potentially relevant factor which
352 could vary at this scale is the grain size of the sand used by the worms to build the tubes.
353 Changes in this variable between reefs separated by a distance comparable with that of present
354 sites were reported in Wales (UK) and related with changes in growth, morphology and
355 composition of *S. alveolata* aggregations (Naylor and Viles, 2000). Indeed, the degree of
356 development or degradation of sabellariid reefs is typically associated with considerable
357 changes in their hosted biodiversity, with the taxonomic richness of associated benthic
358 macrofauna tending to decrease within ‘healthy’ areas with large densities of honeycomb
359 worms, and to decrease in degraded areas (Gruet and Bodeur, 1997), often characterized by an
360 intense colonization by epibionts (Dubois et al., 2006). Quantitative data on the healthy vs.
361 degraded status of our sampled reefs were not available, but some considerations may lead to
362 exclude such differences as the most likely explanation for between-sites variation. In fact, the
363 reefs sampled at each location were all characterized by the virtual lack of conspicuous

364 epibionts (detailed pictures are reported by Schimmenti et al., 2016). At the same time, no
365 significant differences between sites were found for the opercular length of *S. alveolata*.
366 Assuming that the proportion of small *S. alveolata* individuals, being directly related with the
367 number of young recruits, should differ depending on the development stage, and possibly the
368 level of degradation, of reefs, it could be hypothesized that the sampled reefs were comparable
369 in terms of such variables.

370 Within the fauna associated with present biogenic reefs, the most abundant species was *E.*
371 *ornata*. This phyllodocid polychaete uses *S. alveolata* bio-constructions as a preferential
372 habitat, as reported along European east Atlantic and Mediterranean coasts (Pleijel and Dales,
373 1991; Schimmenti et al., 2016). Actually, the Atlantic-Mediterranean distribution of *E. ornata*
374 mirrors that of *S. alveolata* reefs, suggesting the fidelity of the phyllodocid to such biogenic
375 habitat (Schimmenti et al., 2016). Finding *E. ornata* as the most abundant species in *S.*
376 *alveolata* reefs opens ecologically relevant questions, including whether Mediterranean *Eulalia*
377 specimens previously named as *E. clavigera* or *E. viridis* were correctly identified, and what
378 functional role *E. ornata* plays in the functioning and dynamics of biogenic reefs. If
379 carnivorous like most phyllodocids, such as *E. viridis* (Jumars et al., 2015), large numbers of *E.*
380 *ornata* should rely on the availability of abundant prey in the reef. Since most biomass in this
381 system is provided by the bio-constructing worm itself, intense predation by *E. ornata* on *S.*
382 *alveolata* could critically affect the structure and dynamics of the entire reef (Schimmenti et al.,
383 2016).

384 Following *E. ornata*, the syllid *S. pulvinata* was common and numerous in the examined
385 samples, providing interesting data on its occurrence in the Mediterranean Sea. Specifically, *S.*
386 *pulvinata* was not listed in the Italian fauna (Castelli et al., 1995) until recently (Musco et al.,
387 2004), possibly due to synonymy (Licher, 1999). Concomitantly, the peculiarity of the faunal
388 assemblages of sabellariid reefs in the Sicily Channel is reinforced by the occurrence of some
389 numerically scarce species, such as the amphipods *G. ulrici* and *J. ocia*. The former species is

390 considered endemic to the Mediterranean Sea, but records along the Italian coasts are very
391 sparse and rare (but see Lo Brutto and Sparla, 1993; Bedini et al., 2011; Iacifano et al., 2015).
392 The latter was previously found in large abundances within *S. alveolata* reefs located along the
393 central Tyrrhenian coast, about 500 km north of present locations (La Porta et al., 2009), and,
394 therefore, could be a species using the sabellariid bio-constructions as a preferential habitat.

395 In conclusion, identifying and testing the discussed processes as responsible for patterns of
396 distribution, abundance and diversity of sabellariid reefs and associated fauna were beyond the
397 goals of this study. Being based on a descriptive approach, present findings contribute to
398 propose plausible drivers of such patterns, but future, specifically designed, experiments are
399 needed to elucidate the actual causal role of each process and of their possible interactions.
400 Nevertheless, if including biogenic reefs in European directives and in the Red List of
401 endangered habitats has an obvious goal in protecting them as ecologically important, but
402 threatened, systems, the present findings have several implications for establishing the
403 corresponding priorities. First, identifying the relevant scales of variation of habitat-forming
404 species and of their associated biodiversity, as well as their faunal peculiarities, is crucial to
405 design protection schemes suitable to guarantee a representative set of target populations and
406 assemblages and, likely, of the processes driving their spatial variation. To this end, effective
407 strategies of protection of sabellariid reefs in the Sicily Channel would require to include all
408 locations examined here as collectively providing a representative sample of the biota
409 associated with this habitat. Any future decision in this direction, however, will need to be
410 based on the confirmation of present patterns through temporally replicated studies. Indeed, the
411 present findings necessarily apply only to the particular time at which this study was carried
412 out, and it cannot be assumed that spatial patterns are consistent over time (e.g. Underwood
413 and Petraitis, 1993; Benedetti-Cecchi et al., 2001). Second, recent experimental and theoretical
414 investigations started to clarify how natural and anthropogenic perturbations, especially those
415 due to climate change, ultimately modulate morphological, physiological and distributional

416 responses of natural populations and assemblages and the ecosystem services they provide
417 (reviewed by Torossian et al., 2016). In this context, this study may provide reference data
418 useful to relate climatic variation with ongoing changes in patterns of an important biogenic
419 habitat and its supporting species at the edge of their distribution limits. Marginal populations
420 are assumed to deal with sub-optimal environmental conditions (Bridle and Vines, 2007) which
421 could reduce their ability to resist to and/or recover from further disturbances (Guo et al., 2005;
422 Hampe and Petit, 2005). In addition, they are, like in the present case, often characterized by
423 distinctive ecological characteristics that need to be preserved against the negative impact of
424 environmental modifications, such as those observed and predicted as a consequence of climate
425 change, especially in the Mediterranean Sea (e.g. Marbà et al., 2015). Finally, sabellariid reefs
426 are the focus of recent attempts to restore their habitats where degraded (e.g. Callaway et al.,
427 2014), as required by the Habitats Directive. Present information on ecological and biological
428 patterns of sabellariid reefs in the Sicily Channel may contribute to assess their status, establish
429 terms of comparison for similar reefs elsewhere and evaluate the effectiveness of possible
430 restoration interventions of this habitat and its hosted biota.

431

432 **Acknowledgements**

433 This study was partially supported by the Norwegian Taxonomy Initiative (knr. 49-13, per.
434 70184228), MIUR - Italian Ministry of Education, University and Research through the
435 projects “Biocostruzioni costiere: struttura, funzione e gestione” (PRIN 2010-2011, grant F81J
436 120003 80001), and the University of Palermo (FFR 2012/2013). The authors are grateful to C.
437 Calvaruso and E. Keci for technical assistance, and to D. Iacofano for helping with amphipod
438 identification at the University of Palermo.

439

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Table 1. Results of analysis of variance (ANOVA) on the abundance (number of individuals) of *Sabellaria* spp., the total richness (number) of taxa and the Shannon's diversity index (H) of assemblages associated with sabellarid reefs in the Sicily Channel. * p<0.05, ** p<0.01, *** p<0.001.

Source of variation	df	<i>Sabellaria total</i>		<i>Sabellaria alveolata</i>		<i>Sabellaria spinulosa</i>		Richness		Shannon's H		Denominator
		MS	F	MS	F	MS	F	MS	F	MS	F	
Location	2	1.43	0.79	1.41	0.77	2.88	1.34	215.54	8.07	1.80	10.51	Site(Location)
Site(Location)	3	1.80	40.70***	1.83	42.25***	2.15	1.45	26.71	1.19	0.17	1.36	Residual
Residual	18	0.04		0.04	1.49		22.43			0.13		
Cochran's test		C = 0.375		C = 0.374		C = 0.306		C = 0.518		C = 0.496		
Transformation		Ln(x+1)		Ln(x+1)		Ln(x+1)		None		None		

Table 2. Components of variation, expressed as untransformed variance values (σ^2) and percentage of total variance (%), at each of three spatial scales (among replicates, between sites and among locations) of the structure of whole faunal assemblages (*Sabellaria* spp. excluded), individual abundance of *Sabellaria* spp., richness (total number of taxa), Shannon's diversity index (H) and individual abundance of taxa associated with sabellarid reefs in the Sicily Channel.

Variable	Replicate scale		Site scale		Location scale	
	σ^2	%	σ^2	%	σ^2	%
<i>Sabellaria</i> tot.	104151	17.69	484621	82.31	0	0
<i>Sabellaria alveolata</i>	99452	17.15	480533	82.85	0	0
<i>Sabellaria spinulosa</i>	111.72	90.09	0	0	12.29	9.91
Richness	22.43	47.62	1.07	2.27	23.60	50.11
Shannon's H	0.13	38.24	0.01	2.94	0.20	58.82
Assemblage (abundances)	888.20	44.73	479.14	24.13	618.36	31.14
Assemblage (P/A)	1222.40	62.05	197.50	10.03	550.16	27.98
<i>Eulalia ornata</i>	521.94	44.32	130.70	11.12	522.95	44.48
<i>Syllis pulvinata</i>	368.22	29.40	144.53	11.54	739.62	59.06
<i>Syllis armillaris</i>	20.72	19.44	85.88	80.56	0	0
<i>Syllis gracilis</i>	14.18	29.67	25.17	52.67	8.44	17.66
<i>Nereis splendida</i>	67.63	79.79	0	0	17.13	20.21
<i>Brania arminii</i>	32.76	47.62	16.86	24.51	19.18	27.88
<i>Nereis rava</i>	1.17	37.86	1.92	62.14	0	0
<i>Arabella iricolor</i>	5.15	73.05	0	0	1.90	26.95
<i>Syllis garciai</i>	0.97	40.76	0	0	1.41	59.24
<i>Gammaropsis ulrici</i>	106.36	24.77	313.51	73.00	9.59	2.23

<i>Jassa ocia</i>	159.28	67.59	76.39	32.41	0	0
<i>Caprella rapax</i>	12.44	65.20	6.64	34.80	0	0
<i>Apolochus neapolitanus</i>	5.69	61.71	0	0	3.53	38.29
<i>Quadrimaera</i> sp.	6.81	92.15	0.31	4.19	0.27	3.65
<i>Tethylembos viguieri</i>	7.06	72.26	0	0	2.71	27.74
<i>Caprella acanthifera</i>	7.43	80.41	0	0	1.81	19.59

Table 3. Results of permutational multivariate analysis of variance (PERMANOVA) on the structure of sabellarid reef assemblages (*Sabellaria* spp. excluded), examined as abundance or presence/absence data, in the Sicily Channel. Significant effects are indicated in bold.

Source of variation	df	MS	pseudo- <i>F</i>	p	Unique perm.	Denominator
<u>Abundance</u>						
Location	2	7751.7	2.8	0.016	999	Site(Location)
Site(Location)	3	2804.8	3.2	0.001	998	Residual
Residual	18	888.2				
<u>Presence/Absence</u>						
Location	2	6413.5	3.2	0.005	998	Site(Location)
Site(Location)	3	2012.2	1.6	0.014	997	Residual
Residual	18	1222.4				

FIGURE CAPTIONS

Fig. 1. Mean (+SE, n = 4) abundance of total *Sabellaria*, *S. alveolata*, *S. spinulosa*, richness of taxa and Shannon's diversity index from three locations (TRI = Triscina, ERA = Eraclea Minoa, DON = Donnalucata) and two sites (paired bars of the same colour) within each location in the Sicily Channel.

Fig. 2. Principal Component Ordination (PCO) of assemblages (*Sabellaria* spp. excluded) from sabellarid reefs between three locations and two sites within each location in the Sicily Channel.

Fig. 3. Mean (+SE, n = 4) abundance of nine polychaete species associated with sabellarid reefs from three locations and two sites (paired bars of the same colour) within each location in the Sicily Channel. Abbreviations as in Fig. 2.

Fig. 4. Mean (+SE, n = 4) abundance of Sipunculidae and seven amphipod taxa associated with sabellarid reefs from three locations and two sites (paired bars of the same colour) within each location in the Sicily Channel. Abbreviations as in Fig. 2.

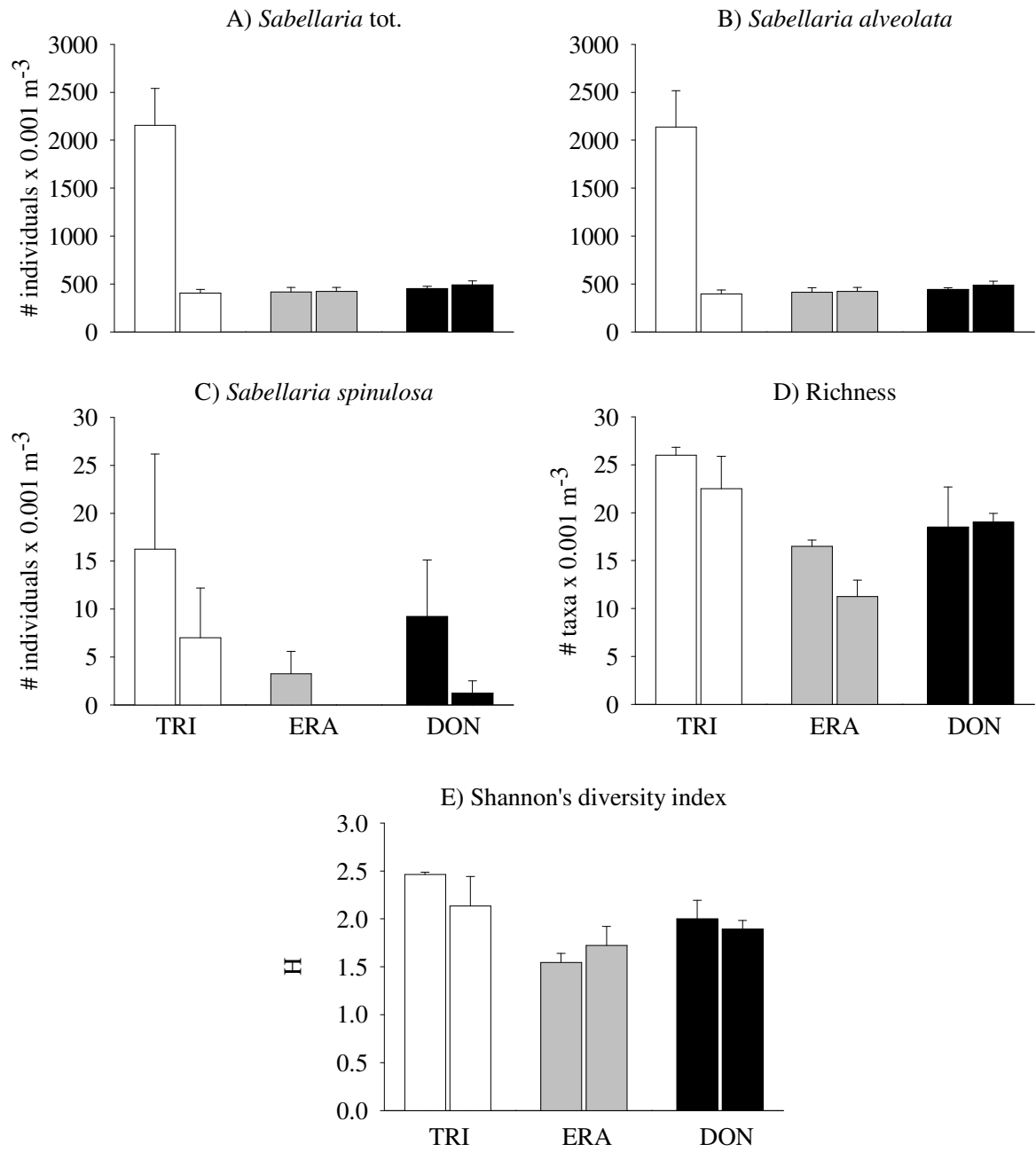


Fig. 1 Bertocci et al.

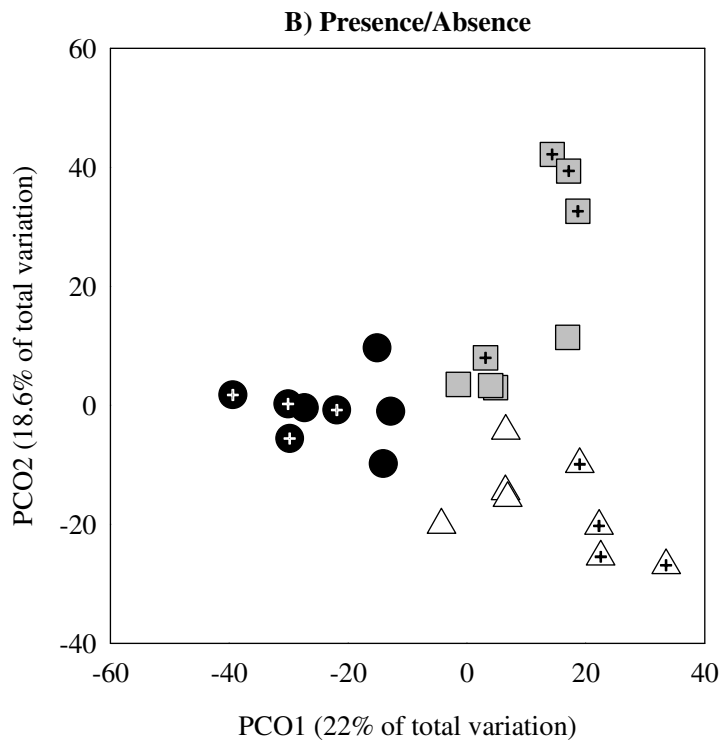
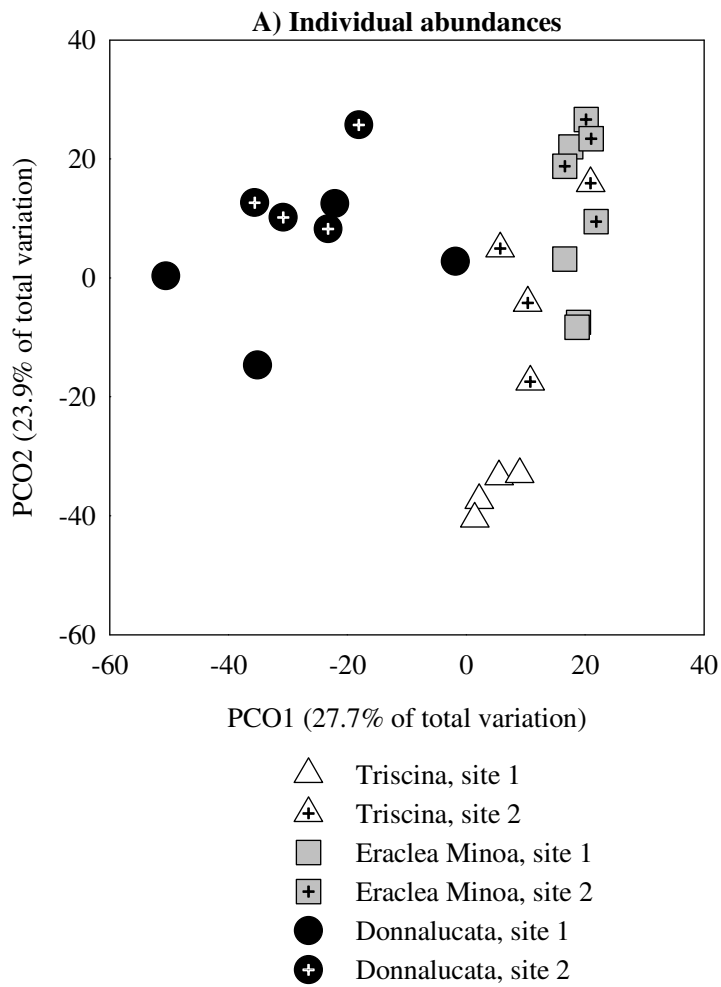


Fig. 2 Bertocci et al.

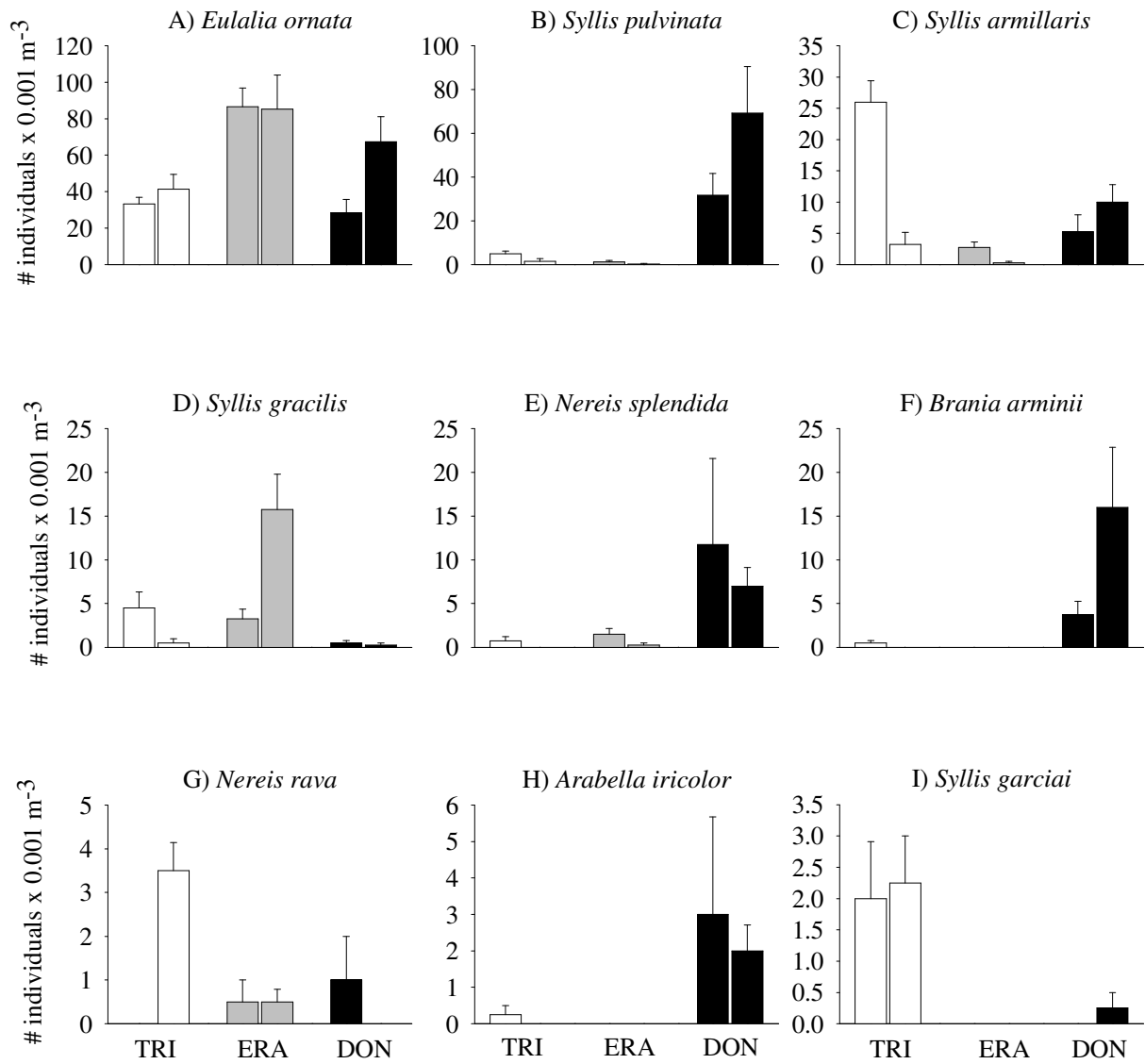


Fig. 3 Bertocci et al.

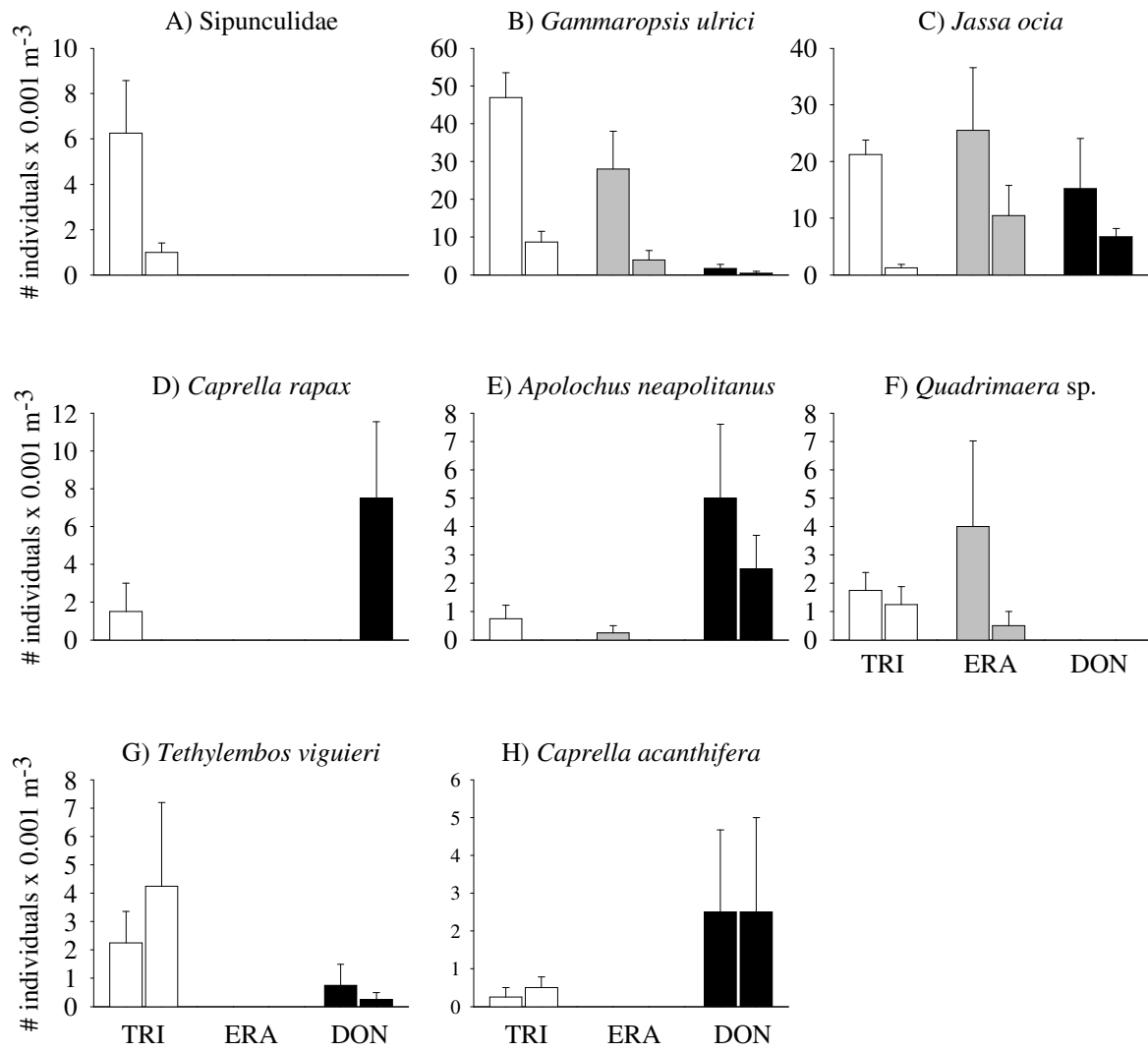


Fig. 4 Bertocci et al.