

# Multi-benthic size approach to unveil different environmental conditions in a Mediterranean harbor area (Ancona, Adriatic Sea, Italy) (#81322)

1

First submission

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# Multi-benthic size approach to unveil different environmental conditions in a Mediterranean harbor area (Ancona, Adriatic Sea, Italy)

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Ports are hubs of human activity and are subject to the continuous discharge and release of industrial, agricultural, and municipal waste and contaminants. Benthic organisms are largely known to reflect environmental conditions they live in. The meio- and macrofauna are ecologically distinct components of the benthos and as such may not necessarily respond to environmental conditions and/or disturbances in the same way. However, in few field studies the spatial patterns of meio- and macrofauna have been simultaneously compared. In the present study, we hypothesized a different response (*i.e.* abundance, diversity, and distribution patterns) from the two benthic size classes to the different environmental conditions they live in (*i.e.* concentration of selected trace metals and Polycyclic aromatic hydrocarbons (PAHs); organic matter contents into the sediment; grain size and microbial abundance) characterizing the Ancona harbor (Adriatic sea). Meio- and macrofauna provided partially similar and complementary types of information depending on the indices used (univariate measures or community structure/ species composition) and the different 'response-to-stress'. The community composition of both benthic size components clearly showed differences among sampling stations located from inside to outside the harbor, reflecting the marked environmental heterogeneity and disturbance typically characterizing these systems. Notwithstanding, the univariate measures (*i.e.* meio- and macrofauna total abundance, diversity indices and equitability) didn't show similar spatial patterns. Meiofauna results were generally statistically more significant than those obtained for macrofauna. Meiofauna were likely to be more sensitive to the effects of

environmental features and contaminants than macrofauna. Overall, trace metals and PAHs affected the community composition of the two benthic components, but only the meiofauna abundance and diversity were related to the environmental variables considered (*i.e.* quantity and quality of food sources). Our results pinpoint the importance and the advantage of the complementary use of two sets of faunistic groups which could provide greater insight into the processes affecting the investigated area.

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26

27 **Abstract**

28 Ports are hubs of human activity and are subject to the continuous discharge and release of  
29 industrial, agricultural, and municipal waste and contaminants. Benthic organisms are largely  
30 known to reflect environmental conditions they live in. The meio- and macrofauna are  
31 ecologically distinct components of the benthos and as such may not necessarily respond to  
32 environmental conditions and/or disturbances in the same way. However, in few field studies the  
33 spatial patterns of meio- and macrofauna have been simultaneously compared. In the present  
34 study, we hypothesized a different response (*i.e.* abundance, diversity, and distribution patterns)  
35 from the two benthic size classes to the different environmental conditions they live in (*i.e.*  
36 concentration of selected trace metals and Polycyclic aromatic hydrocarbons (PAHs); organic  
37 matter contents into the sediment; grain size and microbial abundance) characterizing the  
38 Ancona harbor (Adriatic sea).

39 Meio- and macrofauna provided partially similar and complementary types of information  
40 depending on the indices used (univariate measures or community structure/ species

41 composition) and the different ‘response-to-stress’. The community composition of both benthic  
42 size components clearly showed differences among sampling stations located from inside to  
43 outside the harbor, reflecting the marked environmental heterogeneity and disturbance typically  
44 characterizing these systems. Notwithstanding, the univariate measures (*i.e.* meio- and  
45 macrofauna total abundance, diversity indices and equitability) didn’t show similar spatial  
46 patterns. Meiofauna results were generally statistically more significant than those obtained for  
47 macrofauna. Meiofauna were likely to be more sensitive to the effects of environmental features  
48 and contaminants than macrofauna. Overall, trace metals and PAHs affected the community  
49 composition of the two benthic components, but only the meiofauna abundance and diversity  
50 were related to the environmental variables considered (*i.e.* quantity and quality of food sources).  
51 Our results pinpoint the importance and the advantage of the complementary use of two sets of  
52 faunistic groups which could provide greater insight into the processes affecting the investigated  
53 area.

54

55

## 56 Introduction

57 Coastal waters are widely recognized as marine areas of high ecological and economic value, but  
58 also as highly threatened zone, exposed to multiple human activities and their negative impacts  
59 (Travizi *et al.*, 2019).

60 Harbors are enclosed areas and hubs of human activity. They are essential to the economic  
61 growth of coastal regions, where maritime traffic, shipping, international trade, and fishing are  
62 continuously increasing (Simonini *et al.*, 2005; Franzo *et al.*, 2022). Harbors are usually  
63 characterized by high sediment pollution levels due to heavy metals and hydrocarbons produced  
64 by intense maritime traffic and huge organic matter loads (Covazzi Harriague *et al.*, 2007;  
65 Baldrighi *et al.*, 2019). The high concentrations of contaminants and the relevant inputs of  
66 organic matter represent a persistent and ongoing threat, especially for the biota living in the  
67 sediment (Moreno *et al.*, 2008a; Veiga *et al.*, 2009). In addition, the exposure of the innermost  
68 part of harbors to both wind and waves is limited and may create conditions of reduced water  
69 renewal, favoring sedimentation processes, anoxia, and trapping pollutants (Guerra-García &  
70 García-Gómez, 2005; Spagnolo *et al.*, 2011). The EU Water Framework Directive (WFD;  
71 Directive 2000/60/EC) considers harbors as ‘heavily modified water bodies’, which cannot meet  
72 the common criteria of good ecological quality status. Therefore, their effective management is  
73 crucial for the sustainable use of these maritime spaces and for the protection of the adjacent  
74 coastal habitats (Chatzinikolaou *et al.*, 2018; Franzo *et al.*, 2022).

75 Benthic organisms are largely known to reflect environmental conditions they live in. Among  
76 benthic components, meio- and macrofauna are widely recognized as good ecological indicators  
77 (Schratzberger *et al.*, 2003; Patricio *et al.*, 2012). Benthic meiofaunal (<500  $\mu\text{m}$ ) and  
78 macrofaunal (>500  $\mu\text{m}$ ) communities are regularly utilized in impact assessment, but very few  
79 studies are carried out taking into account both communities (Whomersley *et al.*, 2009;  
80 Frontalini *et al.*, 2011). Meio- and macrofaunal assemblages do not exist in isolation and

81 therefore are part of an interacting system (Whomersley *et al.*, 2009). However, the meio- and  
82 macrofauna are ecologically distinct components of the benthos and as such may not necessarily  
83 respond to environmental conditions and/or disturbances in the same way (Patrício *et al.*, 2012;  
84 Covazzi Harriague *et al.*, 2013).

85 As well as being separated on the basis of size, meio- and macrobenthos each have a series of  
86 distinctive ecological and evolutionary characteristics. Meiofauna is characterized by small size,  
87 high abundance, ubiquitous distribution, rapid generation times, fast metabolic rates, and absence  
88 of a planktonic phase, resulting in a short response time and high sensitivity to different  
89 environmental conditions and certain types of disturbance. Meiofauna can also suitably reflect  
90 the ecological conditions present in a particular system. Due to their ecological characteristics,  
91 meiofaunal organisms can act as suitable indicators of changes in environmental conditions over  
92 small spatial scales (e.g., Schratzberger & Ingels, 2018; Ridall & Ingels, 2021). Consequently,  
93 meiofaunal communities have generated considerable interest as potential indicators of  
94 anthropogenic disturbances in aquatic ecosystems (e.g., Balsamo *et al.*, 2012; Semprucci *et al.*,  
95 2015; Semprucci *et al.*, 2022).

96 Benthic macrofauna is relatively sedentary and thus cannot easily escape natural and  
97 anthropogenic disturbances. Moreover, several species are very sensitive to environmental  
98 changes (Simboura & Zenetos, 2002). For these reasons, these organisms have been commonly  
99 used as bioindicators of local pressure, essential for obtaining information on the state of  
100 conservation of marine ecosystems (e.g., Grotti *et al.*, 2016). Macrobenthic invertebrates have  
101 been identified by the EU WFD as key biological components to assess the ecological status of  
102 aquatic ecosystems, due to their important role in ecosystem functioning and to their  
103 involvement in food-web nutrient recycling (Punzo *et al.*, 2017).

104 Our understanding of benthic systems and how they behave in response to disturbance events  
105 may therefore be improved if a more holistic ecosystem approach to disturbance impact studies  
106 was taken, simultaneously considering more than a single faunal group (Warwick *et al.*, 2006).  
107 The complementary use of two sets of faunistic groups with contrasting ecological characteristics  
108 could provide greater insight into the processes affecting such an area.

109 Distribution and composition of benthic communities is influenced by a wide and complex array  
110 of environmental factors (Covazzi Harriague *et al.*, 2013). The physical instability of some  
111 environments such as harbors, the morphodynamic features of their sediment texture, presence of  
112 pollutants and organic loads contribute to the structure and distribution of benthic assemblages  
113 (e.g., Spagnolo *et al.*, 2011; 2019; Losi *et al.*, 2013; Baldrighi *et al.*, 2019). Up to now, in a few  
114 field studies in which the spatial patterns of meio- and macrofauna have been simultaneously  
115 compared, changes in both assemblages as a response to natural gradients were found to be  
116 scattered across habitats (Patrício *et al.*, 2012). The investigations have demonstrated the  
117 fundamental advantage of a multi-species approach, with the inclusion of many taxonomic and  
118 functional groups that have a broad range of sensitivities to any given environmental regime  
119 (e.g., Frontalini *et al.*, 2011).



120 Adriatic Sea ecosystem is negatively affected by many kinds of biological and ecological threats  
121 *e.g.*, eutrophication, pollution, fragmentation of benthic habitats, invasion of alien species  
122 (Katsanevakis *et al.*, 2011; Pećarević *et al.*, 2013; Corriero *et al.*, 2016). In the regional  
123 perspective the basin is highly positioned on the list of ‘Priority issues in the Mediterranean  
124 environment’ drawn up by the European Environment Agency (EEA), with 20 (15%) out of the  
125 131 hotspot pollution sites identified along the Mediterranean coastline in the frame of the  
126 Strategic Action Programme (SAP) of the United Nations Environment Programme (UNEP;  
127 EEA, 2006), including many harbors such as the Ancona one (Travizi *et al.*, 2019).  
128 Since previous works published on Ancona harbor have always treated the different benthic  
129 components separately (Mirto & Danovaro 2004; Spagnolo *et al.*, 2011; 2019; Baldrighi *et al.*,  
130 2019; Travizi *et al.*, 2019; Franzo *et al.*, 2022), the aim of the present work is to test whether  
131 meio- and macrofaunal assemblages could provide a comparable and/or complementary  
132 assessment of its ecological conditions. Considering that meio- and macrofauna are ecologically  
133 distinct components of the benthos, **we hypothesized a different response** (*i.e.* abundance,  
134 diversity, and distribution patterns) from the two benthic size classes to the different  
135 environmental conditions they live in (*i.e.* concentration of selected **Trace Elements** (TEs) and  
136 Polycyclic aromatic hydrocarbons (PAHs); organic matter contents into the sediment; grain size  
137 and microbial abundance) characterizing the Ancona harbor.

138

## 139 **Materials & Methods**

### 140 **Sampling area and sampling strategy**

141

142 The Ancona harbor (water depth, 4-15 m) is located in the western coast of central Adriatic Sea  
143 (Fig. 1), it has a water sheet of 700,000 m<sup>2</sup> and 5,400 m of docks (Spagnolo *et al.*, 2011). The  
144 harbor is one of the most important of the Adriatic Sea, with intense ferryboat and merchant ship  
145 activity. More than one million passengers on ferries and cruise ships travel from Ancona to the  
146 Adriatic eastern coasts, and both container and oil traffic have also developed in recent years. In  
147 the shipyards, all kinds of ships are designed and built, and shipbuilding is the largest  
148 entrepreneurial reality in the harbor (Franzo *et al.*, 2022). Previous investigations reported that  
149 the area is subjected to organic waste dumping derived from fishing boats and is also affected by  
150 a strong industrial pollution due to the presence of shipyards. Consequently, a huge organic  
151 matter load and high heavy metal and hydrocarbon concentrations are present inside the harbor  
152 area (Mirto & Danovaro, 2004; Bianchelli *et al.*, 2016).

153 In the present study, sediment samples were collected in winter 2015 from five sampling stations  
154 located from inside to outside the Ancona harbor (Fig. 1). The five sampling stations were  
155 chosen according to the different environmental features and anthropogenic activities present in  
156 the area: **MAN station was located in the inner part of the harbor where small fishing boats dock;**  
157 **PORT station was located in a transition area where work ships are berthed; DS and LR stations**


158 were located in a more external position, nearby shipping facilities such as active berths; API  
159 station was located outside the harbor where no activity takes place.

160 At each station, sediment samples for **characterizing the benthic fauna (meio- and macrofauna)**  
161 **and environmental features were collected with a box-corer (40×30 cm wide and 50 cm high) in**  
162 **three independent replicates**, processed and preserved differently according to the analysis to be  
163 performed (see below). At all stations, the temperature and salinity at the sea bottom were  
164 measured using CTD (Conductivity, Temperature, and Depth) probe equipped with previously  
165 calibrated sensors.

166

### 167 **Environmental variables and microbial component**

168

169 The content of each box-corer was sub-sampled with PVC corers (inner diameter, 4.5 cm) to  
170 assess the biochemical composition of the organic matter and grain size. The top 3 cm of  
171 sediment from three independent replicates for each parameter were frozen at -20°C, except for  
172 the grain size determination, for which samples were kept at *in situ* temperature in single  
173 replicate until brought to the laboratory. The biochemical composition of the organic matter  
174 **(protein, carbohydrate, and lipid concentration)** and chloroplastic pigments (chlorophyll-*a* (Chl-  
175 *a*) and phaeopigment (Phaeo) concentration) were determined by standard techniques (Danovaro,  
176 2010). Concentrations were calculated using standard curves and normalized to sediment dry  
177 weight after desiccation (60°C, 24 h). Biopolymeric organic carbon (BPC) was calculated as the  
178 sum of the carbon equivalents of carbohydrates, proteins, and lipids (Fabiano *et al.*, 1995) and  
179 was used as a proxy for the available trophic resources. The value of the protein to carbohydrate  
180 ratio (PRT/CHO) was utilized as descriptor of the nutritional quality of organic matter in the  
181 sediment, with a PRT/CHO ratio  indicating relatively high quality and high food  
182 availability (Pusceddu *et al.*, 2010).

183 For grain size determination, aliquots of fresh sediment were sieved over a 63 µm mesh. The two  
184 fractions (>63 µm, sand; <63 µm, silt/mud) were dried in an oven at 60°C and weighed. Data  
185 were expressed as a percentage of sediment total dry weight (Pusceddu *et al.*, 2010).

186 For **microbial abundance determination, samples of surface seawater** were collected at each  
187 sampling station using sterile containers (capacity, 1 L) and preserved in formaldehyde (final  
188 concentration, 2%). The Total Prokaryotic Number (TPN) was determined by acridine orange  
189 staining technique (Luna *et al.*, 2002), using an Axioskop 2 epifluorescence microscopy (Carl  
190 Zeiss AG, Oberkochen, Germany; magnification, 1,000x).

191

### 192 **Contaminants into the sediment**

193

194 Concentration of **selected TEs and PAHs** was determined in the surface sediment (0-3 cm)  
195 collected at each sampling station in triplicate and frozen at -20°C. Analyses were conducted  
196 following previous validated methods, fully described in Benedetti *et al.* (2014) and Etiope *et al.*  
197 (2014).

198 In brief, TEs were determined after digestion under pressure with nitric acid and hydrogen  
199 peroxide (7:1), using a Mars 6 Microwave Digestion System (CEM Corporation, Charlotte, NC,  
200 USA). As, Cd, Cr, Cu, Fe, Mn, Ni, Pb, V, and Zn were analyzed by Atomic Absorption  
201 Spectroscopy (AAS), with flame (SpectrAA 220FS Spectrometer, Varian Inc., Palo Alto, CA,  
202 USA) and flameless atomization (240Z AA Spectrometer, Agilent Technologies Inc., Santa  
203 Clara, CA, USA), while the Hg content was quantified by Cold Vapor Atomic Absorption  
204 Spectroscopy (CVAAS; QuickTrace M-6100 Mercury Analyzer, Teledyne CETAC  
205 Technologies Ltd., Omaha, NE, USA).

206 PAHs were determined after KOH-methanol extraction with a Mars 6 Microwave Digestion  
207 System (CEM Corporation). Extracts were concentrated using a RC 10.09 Vacuum  
208 Concentration System (Jouan SA, Saint-Herblain, France) and purified by J.T.Baker™  
209 BAKERBOND™ Octadecyl (C18) Solid Phase Extraction (SPE) cartridges (Avantor Inc.,  
210 Radnor, PA, USA). PAHs were analyzed by High-Performance Liquid Chromatography (HPLC)  
211 using both Fluorimetric (FLD) and (UV) Diode Array Detection (DAD) (Infinity 1260 Series,  
212 Agilent Technologies, Santa Clara, CA, USA).

213 For both TEs and PAHs, appropriated blank solutions and the Standard Reference Material  
214 (SRM) 1944: *New York/New Jersey Waterway Sediment* (NIST – National Institute of Standards  
215 and Technology), digested as samples, were used to check for accuracy, precision, and  
216 recoveries of the employed analytical methodologies; concentrations obtained from SRM  
217 analyses were always within the 95% confidence intervals of the NIST certified values.  
218 The results obtained in this study were compared with threshold values for chemicals specified in  
219 the Ministerial Decree 173/2016, the Italian normative that rules the management of dredged  
220 sediments and sets out their quality. Only those values exceeding the upper thresholds values  
221 (L2) are defined as alerting values (Table 2.5, Ministerial Decree 173/2016).

222

## 223 **Meiofauna and macrofauna**

224

225 For meiofauna samples, the content of each box-corer (three independent replicates) was sub-  
226 sampled with PVC corers (inner diameter, 4.5 cm). The top 3 cm of sediment, where meiofaunal  
227 organisms are typically more abundant, were preserved in 4% buffered formaldehyde. For  
228 meiofaunal extraction, sediment samples were sieved through a 1000 µm mesh; a 32 µm mesh  
229 was used to retain the smallest metazoan organisms. The latter fraction was centrifuged 3 times  
230 with LUDOX® HS-40 colloidal silica (diluted with water to a final density of 1.18 g cm<sup>-3</sup>) and  
231 stained with Rose Bengal (0.5 g L<sup>-1</sup>; Heip *et al.*, 1985). Meiofaunal organisms were counted (no.  
232 of individuals 10 cm<sup>-2</sup>) and identified to the higher taxonomic level under a stereomicroscope.  
233 For macrofauna samples, the first 20 cm of three independent box-corer deployments were  
234 sieved *in situ* using a 500 µm mesh and all organisms retained were preserved in 5% buffered  
235 formaldehyde. Macrofauna was sorted in laboratory using a stereomicroscope and a binocular  
236 microscope, identified and classified to the lowest possible taxonomic level using standard  
237 nomenclature, and quantified (no. of individuals m<sup>-2</sup>). The data collected was subjected to a

238 control and validation process and organized in a dedicated database. The nomenclature of the  
239 species was verified and validated using the web portal <https://www.marinespecies.org/> and the  
240 ‘WoRMS Taxon Match Tool’ (World Register of Marine Species Editorial Board, 2022).  
241 Taxon richness (*i.e.* higher taxa and species for meio- and macrofauna, respectively; S), total  
242 number of individuals per taxon/species (N), Shannon’s diversity index ( $H'$ , based on  $\log_2$ ;  
243 Shannon & Weaver, 1949), and Pielou’s equitability index ( $J'$ ; Pielou, 1969) of benthic  
244 communities were calculated.

## 245 Statistical analysis

247  
248 To assess differences in benthic communities among the sampling stations a Permutational  
249 Multivariate Analysis of Variance (PERMANOVA; 9999, number of random unrestricted  
250 permutations of raw data) was used (Anderson, 2001). The design included one factor: the  
251 sampling station (five levels, fixed). The analysis was based on Bray-Curtis’ similarity of  
252 previously fourth root transformed meio- and macrofaunal data. In case of significant differences  
253 obtained by the main test, the pairwise test was performed and, as there was a limited number of  
254 unique permutations, the  $p$  values were obtained from Monte Carlo tests (Anderson & Robinson,  
255 2003). Permutational Multivariate Analysis of Dispersion (PERMDISP) test was applied to  
256 assess if differences among the sampling stations (between-group) were due to real differences in  
257 benthic community composition and not to differences in the multivariate dispersion of replicates  
258 (within-group) among their respective centroids. A non-metric Multidimensional Scaling  
259 (nMDS) ordination was carried out to visualize similarities among the sampling stations.  
260 Similarity Percentages (SIMPER) analysis (cut-off, 90%) was used to identify the meio- and  
261 macrofaunal taxa that contributed to the dissimilarity among the sampling stations. These  
262 procedures were performed with PRIMER™ and PERMANOVA+ ecological software (Clarke  
263 & Gorley, 2006; Anderson *et al.*, 2008). For all tests the significance threshold was set to 0.05.  
264 One-way Analysis of Variance (ANOVA) was used to explore differences among the sampling  
265 stations for organic matter content and microbial component, total abundance, diversity and  
266 equitability indices in meio- and macrofaunal benthic communities. The ANOVA assumptions  
267 were tested graphically plotting residuals *vs.* fitted values, normality of residuals and residuals  
268 *vs.* covariate (factor station) to assess the variance homogeneity (Zuur *et al.*, 2016). When  
269 ANOVA showed significant differences, the Tukey’s Honestly Significant Difference (HSD) test  
270 was performed to find significant effects between different levels. ANOVA was performed using  
271 the free software R (R Core Team, 2018).

272 The number of environmental and pollutant-related variables were separated in two groups:  
273 environmental (comprising TEs, PAHs, and silt/mud percentage) and biotic (comprising bacteria  
274 and all the organic compounds of biogenic origin). Since many compounds were linked each  
275 other, multicollinearity (cut-off at Spearman’s correlation =  $|0.8|$ ) among variables was assessed  
276 to reduce the dataset and avoid problems in the analysis algorithms. In case of multicollinearity,  
277 the variables with a more stringent biological or environmental value were retained acting as

278 proxy for the omitted collinear ones. To further reduce the number of variables (as they were still  
279 more numerous than the meio- and macrofaunal ones), both environmental and biotic variables  
280 were analyzed (separately) by means of a Principal Component Analysis (PCA) to identify  
281 groups with similar variability. Only those principal components showing eigenvalues  $>1$   
282 (Kaiser-Guttman criterion; Zwick & Velicer, 1986) were considered. To obtain a better insight  
283 into the output loadings, the orthogonal varimax rotation of extracted PCA components was  
284 performed. After a varimax rotation, each original variable tends to be associated with one (or a  
285 small number) of PCA axis. By doing so, groups of variables were created each of which  
286 represented by a single PCA score, and all the scores obtained were used as covariates in the  
287 following analysis to assess the relations between faunal communities and the environmental  
288 variables. Spearman's rank correlation analysis was performed to test relationships between  
289 meio- and macrofauna total abundance, taxon richness, Shannon's diversity and Pieolu's  
290 equitability indices, and the environmental variables considered. Spearman's rank correlation  
291 coefficients ( $\rho$ ,  $\rho_h$ ) were considered significant at  $p$  values  $<0.05$ .  
292 In order to verify the existence of a significant relation between the benthos data matrix and the  
293 environmental data, the distance-based Linear Modeling (DistLM) procedure was utilized with a  
294 backward selection of the variables, and each model assessed by means of the Akaike's  
295 Information Criterion corrected for small samples (AICc; Anderson *et al.*, 2008). A distance-  
296 based Redundancy Analysis (dbRDA) was then applied to visually investigate the relationship  
297 between the community assemblages and the environmental variables (Anderson *et al.*, 2008).

298

## 299 Results

300

### 301 Environmental features and contamination levels in the Ancona harbor

302

303 Values of main environmental features and considered pollutant concentrations are reported in  
304 Table 1 and Table 2, respectively.

305 Bottom water temperature reported a mean value of  $10.3^{\circ}\text{C}$  and salinity ranged from a minimum  
306 of 31.3 (at LR station) to a maximum of 37.6 PSU (at API station). The sandy fraction ( $>63\ \mu\text{m}$ )  
307 characterized stations located outside the harbor (DS, LR, and API stations), while the silty  
308 muddy fraction ( $<63\ \mu\text{m}$ ) was predominant in the inner stations MAN and PORT (Table 1).

309 Analyses of Chl-*a* and organic matter contents into the sediments showed significantly ( $F_{4,10} =$   
310  $12.97$ ,  $p = 0.001$ ) higher values of 'fresh' material (*i.e.* Chl-*a*) at MAN and DS stations, while the  
311 BPC significantly ( $F = 22.21$ , 4 d.f.,  $p = 0.001$ ) decreased from inside to outside the harbor  
312 (Table 1). The innermost station MAN was also characterized by the highest abundance of  
313 prokaryotes (TPN) although not significantly different from the other stations. The significant ( $F$   
314  $= 12.61$ , 4 d.f.,  $p = 0.001$ ) highest value in the quality of organic matter (*i.e.* PRT/CHO) was  
315 reported at PORT station (Table 1), due to a particularly low concentration of CHO into the  
316 sediment of this station (Table S1).

317 Analyses aimed at determining TE concentrations in the sediment of the Ancona harbor have  
318 highlighted a general decrease of contamination values moving from the innermost sampling  
319 station (MAN) to those more external (Table 2). MAN station turned out to be affected by  
320 pollution levels 3-4 times higher than those detected at API station, reaching even Cu and Zn  
321 concentrations 13 and 6 times higher, respectively. Just these two elements, Cu and Zn, were the  
322 only exceeding the threshold limits imposed by the Ministerial Decree 173/2016 (46.31  $\mu\text{g g}^{-1}$  of  
323 Cu detected at MAN station; 297.2 and 114.9  $\mu\text{g g}^{-1}$  of Zn detected at MAN and PORT stations,  
324 respectively). In the remaining sampling stations there were no overruns (Table 2).

325 The concentration levels of  $\Sigma_{19}$  PAHs ranged from 73.38 to 213.4  $\mu\text{g g}^{-1}$  following, although not  
326 linearly, the same pattern highlighted for TEs, with higher values in the innermost sampling  
327 station (MAN), decreasing towards outer ones (Table 2). All the sediment samples showed a  
328 distinct predominance of Low Molecular Weight PAHs, mainly driven by the Naphthalene  
329 concentration (exceeding the legislative thresholds at MAN, DS, and LR stations), which  
330 averagely accounted for 37% of the  $\Sigma_{19}$  PAHs, followed by its methylated isomers: 1- and 2-  
331 Methylnaphthalene (24 and 22%, respectively). This pronounced prevalence of volatile, easily  
332 transportable PAHs, with 2-3 aromatic rings (Fig. 2a) suggest that fuel combustion linked to  
333 harbor traffic may be the major source of these organic contaminants in the area (for wind  
334 direction and speed on Ancona harbor area during February 2015, please see Fig. S1). Excluding  
335 Naphthalene and its related compounds, the PAH residual contamination was mainly ascribable  
336 to Phenanthrene and Fluoranthene (Table 2), and the principal PAH diagnostic ratios  
337 (Tobiszewski & Namieśnik, 2012), commonly used as a tool to discriminate the analyte origin  
338 and sources (Giuliani *et al.*, 2019; Pizzini *et al.*, 2021), would seem to suggest a clear petrogenic  
339 origin (Fig. 2b).

340 The results from PCA plots considering 'biotic' and 'abiotic' environmental variables separately,  
341 clearly summarized the differences among the sampling stations. In details, considering the  
342 biotic variables (Fig. 3a) MAN and, for a lesser extent, LR were grouped together being  
343 characterized by higher values of potential food sources and prokaryotes the former, and by the  
344 lowest salinity value the latter. This explained the variability along the first axis, while variability  
345 along the second axis was mainly explained by the contrast between the innermost stations  
346 (MAN and PORT), characterized by higher quantity and quality of organic matter, and the outer  
347 sampling stations. The first two components accounted for 75.7% of the total variability. In  
348 regard to abiotic variables (Fig. 3b), MAN station separated from all the other sampling stations  
349 due to highest values of pollution (both by inorganic and organic contaminants) and in this case  
350 the first two components accounted for 87.3% of the total variability.

351

## 352 **Meiofauna assemblages**

353

354 A total of 12 taxa were identified with Nematoda representing the dominant taxon at all sampling  
355 stations, with a percentage ranging from 75 to 95% (Fig. 4a). The second most represented taxon  
356 was Copepoda with their *nauplii*; among less represented taxa (*i.e. others*) Bivalvia, Ciliata,

357 **Foraminifera**, Kinorhyncha, Oligochaeta, Ostracoda, Platyhelminthes, Polychaeta, Sipuncula,  
358 and Tardigrada constituted from 3 to 15% of the meiobenthic community (Fig. 4b). Meiofauna  
359 abundance and values of its diversity indices are reported in Table S2 and represented in Figure  
360 5. The ANOVA tests detected significant differences among all the sampling stations for N  
361 (Table 3), with the highest value reported at DS station (Fig. 5a). The ANOVA tests reported  
362 significant differences among the sampling stations also for all diversity and equitability indices  
363 (Table 3 and Table S2). Regarding the number of taxa, PORT station showed a significant lower  
364 value on average compared to all the other sampling stations (Fig. 5b). Shannon's diversity and  
365 Pielou's equitability indices showed both similar patterns, with MAN station presenting the  
366 highest value (Fig. 5c and Fig. 5d).  
367 PERMANOVA analysis with pairwise test reported significant differences (Table 4) in  
368 meiofaunal community composition among all the sampling stations; PERMDISP test did not  
369 show any significant dispersion around centroids, confirming that the differences among the  
370 sampling stations were due to a real difference in meiobenthic composition ( $F_{2,4} = 5.38$ ,  $P(\text{perm})$   
371  $= 0.088$ ).  
372 The nMDS plot (Fig. 6a) clearly shows the separation among the sampling stations, as well as a  
373 low inter-replica variability. In particular, the innermost MAN and PORT stations were separated  
374 from to the outermost ones. SIMPER analysis detected a dissimilarity percentage from 15 (LR  
375 vs. DS stations) to 54% (PORT vs. LR stations). The highest values of dissimilarity percentage  
376 were always associated with PORT station and mainly due to very low abundances in some of  
377 the most represented taxa such as: Copepoda, Foraminifera, Nematoda, and Polychaeta (Table  
378 S3). For the other sampling stations, the dissimilarity was mainly due to the presence/absence or  
379 differences in the abundances of the taxa *others*: Bivalvia, Ciliata, Kinorhyncha, Oligochaeta,  
380 Ostracoda, Platyhelminthes, and Sipuncula (Table S3). Several positive correlations emerged  
381 between meiofauna descriptors and quantity of organic matter and TPN; moreover, meiofauna  
382 abundance and its diversity were positively correlated to some TEs and to silt/mud content into  
383 the sediment (Table S4a). The best model selected by the DistLM analysis, reported in Table 5,  
384 comprised only the pollutant compounds (ARC1, 2, 3, and 4). The resulting dbRDA showed that  
385 the first two axes explained 91.8% of the fitted model variation and the 83.4% of the variance of  
386 meiofaunal community composition, corresponding to the combination of the following  
387 environmental factors: **V/Ni ratio**, Naphthalene, and Benzo[*a*]pyrene (ARC1) and Zn, Fluorene,  
388 and percentage of silt/mud (ARC4; Fig. S2a).

### 389 **Macrofauna assemblages**

390  
391  
392 A total of 93 taxa were identified, these included: 43 Anellida (42 Polychaeta and 1  
393 Oligochaeta), 29 Mollusca (22 Bivalvia, 6 Gastropoda, and 1 Scaphopoda), 10 Crustacea (6  
394 Amphipoda, 2 Cumacea, 1 Isopoda, and 1 Tanaidacea), 5 Nematoda, 2 Bryozoa, 2 Cnidaria (1  
395 Anthozoa and 1 Hydrozoa), 1 Nemertea, and 1 Ophiuroidea (Table S5). Anellida was the most  
396 represented group (from 68 to 90% at LR and MAN stations, respectively), followed by

397 Mollusca (from 4 to 24% at MAN and LR stations, respectively) at all the sampling stations.  
398 Other less represented groups such as Cnidaria, Isopoda, Ophiuroidea, and Tanaidacea were  
399 found only at one or two sampling stations (Fig. 7). Macrofauna abundance ranged from  $610 \pm$   
400  $241$  to  $3455 \pm 425$  individuals  $m^{-2}$  at MAN and LR stations, respectively; values of its diversity  
401 indices are reported in Table S2 and represented in Figure 8. The ANOVA tests reported  
402 significant higher abundance value at LR station (Fig. 8a) compared to all the other stations  
403 (Table 3). LR station was also characterized by the highest number of taxa (Fig. 8b) but also by  
404 lowest values (Fig. 8c and Fig. 8d) of Shannon's diversity and Pielou's equitability indices  
405 (Table 3 and Table S2). PERMANOVA analysis with pairwise test reported significant  
406 differences (Table 6) in macrofaunal community composition among the majority of the  
407 sampling stations; PERMDISP test did not show any significant dispersion around centroids,  
408 confirming that the differences among the sampling stations were due to a real difference in  
409 macrobenthic composition ( $F_{2,4} = 0.99$ ,  $P(\text{perm}) = 0.767$ ).

410 The nMDS plot (Fig. 6b) shows the separation among the sampling stations. In detail, the  
411 innermost MAN station and the outermost API station were particularly distinguished from the  
412 others. SIMPER analysis reported high percentages of dissimilarity ranging from 63.4 (LR vs.  
413 DS stations) to 88.3% (MAN vs. API stations) between all pairs of sampling stations. The great  
414 dissimilarity is mainly due to the presence/absence of species or to a particularly abundant  
415 presence of them in one station compared to the others (Table S6). Species like *Spiophanes*  
416 *bombyx*, *Kurtiella bidentata*, and *Euclymene oerstedii* characterized mainly the outermost API  
417 station, while some other species such as *Tubificoides swirencoides*, *Streblospio* sp.,  
418 *Heteromastus filiformis*, and *Chaetozone caputesocis* were found inhabiting the innermost  
419 stations (Table S6). Four significant correlations (three out of four were negative) were detected  
420 between macrofauna descriptors and environmental variables; only macrofauna species richness  
421 was (negatively) correlated to three TEs and to the percentage of finest sediment fraction (Table  
422 S4b).

423 The best model selected by the DistLM analysis, reported in Table 7, comprised only ARC2  
424 (Benzo[ghi]perylene, and Anthracene/(Anthracene + Phenanthrene) and  
425 Fluoranthene/(Fluoranthene + Pyrene) diagnostic ratios) and ARC4 (Zn, Fluorene, and  
426 percentage of silt/mud). The resulting dbRDA showed that the first two axes explained 100% of  
427 fitted model variation and the 42.9% of the variance in the macrofaunal community composition  
428 (Fig. S2b). This latter low percentage indicates that the residual variance associated to the  
429 community was not captured by the graph, and it is likely that an unobserved variable should  
430 have had improved the general plot. In any case, along the first axis of the dbRDA there was a  
431 clear separation between the innermost stations MAN and PORT, characterized by high values of  
432 Zn and percentage of silt/mud compared to the outermost stations LR, DS, and API; while along  
433 the second axis LR and DS stations were separated from API, MAN, and PORT stations.

## 434 Discussion

435  bors are deeply modified coastal areas to meet human requirements. Pressures associated



437 with harbor ecosystems are becoming increasingly important, causing significant damage to  
438 water and sediment quality and, subsequently, to marine life and ecosystems, as well as to human  
439 health (Mestres *et al.*, 2010).  
440 Ancona harbor has been the subject of some previous investigations to evaluate the presence and  
441 impact of several contaminants on macrobenthic community (Spagnolo *et al.*, 2011) and  
442 meiobenthic nematodes (Franzo *et al.*, 2022); to assess the presence of alien species (Non-  
443 indigenous species, NIS) introduced by ballast waters (BW) (Spagnolo *et al.*, 2019; Travizi *et*  
444 *al.*, 2019); to investigate the response of meiobenthic communities under BW impact (Baldrighi  
445 *et al.*, 2019). All these previous studies, conducted on different benthic size components, have  
446 reported that Ancona harbor is a receptor of multiple contaminants, such as toxic compounds,  
447 heavy metals, hydrocarbons, and organic matter, as well as a habitat for NIS introduced by BW  
448 due to intense maritime traffic and which, in turn, exert consequences on the composition and  
449 structure of the benthic communities.

450

#### 451 Environmental features of Ancona harbor

452



453 In Ancona harbor, the innermost stations were characterized by muddy sediments, while sand  
454 dominated the outermost stations. This grain size distribution was clearly due to a reduced  
455 exposure to hydrological factors (wind, waves, and currents) which create conditions of poor  
456 water renewal inside the harbor, favoring the presence of fine sediments (Spagnolo *et al.*, 2011).  
457 The sediment deposition rate can influence sediment organic matter load and pollutant content,  
458 with fine-grained components commonly showing a high content in organic matter and  
459 pollutants (Papageorgiou *et al.*, 2010). This might have facilitated an overall accumulation of  
460 TEs and PAHs inside the harbor. In particular, the high values of Cu and Zn detected at MAN  
461 and PORT stations were likely correlated to the shipyard activities present within the harbor, in  
462 particular with the use of new generation antifouling paints (Costa *et al.*, 2016; Pereira *et al.*,  
463 2018). Furthermore, the pronounced prevalence of volatile, easily transportable PAHs (*e.g.*,  
464 Naphthalene; Fig. 2a) pointed out that fuel combustion linked to maritime traffic was the major  
465 source of these organic contaminants in the harbor basin. The petrogenic origin of the PAH  
466 residual contamination (*e.g.*, Phenanthrene and Fluoranthene; Fig. 2b) was supported by the  
467 detection in the sediment samples of V, Ni, and Pb, commonly considered as tracers of  
468 accidental oil spills and/or marine fuels (El Nemr *et al.*, 2006), as well as by the values of the  
469 V/Ni ratio, marker of an intense maritime traffic (Viana *et al.*, 2014).  
470 Considering the threshold values for chemicals specified in the Ministerial Decree 173/2016, few  
471 values were reported exceeding the established alerting thresholds and always from the  
472 innermost stations. It is also true that the Decree does not report threshold values for all TEs and  
473 PAHs considered in our study, thus making the sediment quality evaluation based only on  
474 Ministerial Decree 173/2016 somewhat weak.  
475 Not only contaminants, but also natural and/or anthropogenic changes in the benthic trophic  
476 status (*i.e.* organic matter quantity and sediment biochemical composition) may affect the

477 benthic communities (Pusceddu *et al.*, 2011; Foti *et al.*, 2014). The protein, carbohydrate, lipid,  
478 and BPC content in the sediments have been proposed and utilized to assess the benthic trophic  
479 status of marine coastal environments, including the Adriatic Sea (Vezzulli & Fabiano, 2006). In  
480 Dell'Anno *et al.* (2002) PRT and CHO sedimentary contents were suggested as proxies and  
481 threshold values for ranking the trophic status and the environmental quality of coastal marine  
482 ecosystems along the Apulia Region. Applying those thresholds to the investigated sediments,  
483 the trophic status of Ancona harbor could be ranked as hyper-trophic (PRT >4 mg g<sup>-1</sup>) with the  
484 exception of the API station, ranked as eutrophic (PRT ≈ 4 mg g<sup>-1</sup>). However, in terms of CHO  
485 content, Ancona harbor should be ranked as meso-oligotrophic system (CHO <5 mg g<sup>-1</sup>).  
486 Pusceddu *et al.* (2009; 2011) identified as eutrophic systems those characterized by BPC  
487 concentration >3 mgC g<sup>-1</sup>, as found at MAN, PORT, LR, and DS stations, and as mesotrophic  
488 systems those characterized by BPC concentration in the range 1-3 mgC g<sup>-1</sup>, as in the case of the  
489 API station. In Ancona harbor PRT/CHO ratio resulted always >1, indicating a great input of  
490 recent production's material and highlighting the good trophic quality of the organic matter  
491 (Pusceddu *et al.*, 2009). Concentrations of Chl-*a* here reported were extremely high if compared  
492 to those reported in February in a previous study conducted along the Adriatic coasts (0.11-0.23  
493 μg g<sup>-1</sup>; Bianchelli *et al.*, 2016), indicating the presence of 'fresh' primary organic matter. TPN  
494 did not change significantly among the sampling stations, however the trend in the abundance  
495 values appeared similar to that of organic matter content.

496 The PCA on measured environmental variables indicated the presence of a clear spatial  
497 heterogeneity among the sampling stations and a separation between innermost stations and  
498 outermost ones due to higher organic matter, prokaryote and contaminant loads inside the harbor  
499 basin, as previously reported in the same study area (Spagnolo *et al.*, 2011; Baldrighi *et al.*,  
500 2019) and in other enclosed systems (Vezzulli *et al.*, 2003; Losi *et al.*, 2013). This marked  
501 environmental variability in harbors is a common feature. Indeed, environmental disturbance  
502 within harbors may change rapidly over spatial scales of a few meters, depending on various  
503 factors like the localization and magnitude of pollution sources, allochthonous inputs of different  
504 nature, tidal regime, water circulation, harbor position, shape, and size (Vassallo *et al.*, 2006).

505

### 506 **Meiofaunal response to harbor environmental conditions**

507

508 Harbor communities are subject to broad spatial and temporal variability of physical-chemical  
509 conditions and environmental disturbances, and meiofauna is usually able to respond rapidly to  
510 such changes (Vezzulli *et al.*, 2003).

511 In Ancona harbor, the meiofaunal total abundance, community structure and, for a lesser extent,  
512 univariate measures (*i.e.* diversity and equitability indices) reflected the marked spatial  
513 heterogeneity showed by the PCA and the clear separation both between inner and outer  
514 sampling stations and among the sampling stations themselves (MAN vs. PORT vs. DS + LR vs.  
515 API). Meiofaunal abundance was in the range described by Baldrighi *et al.* (2019) for Ancona  
516 harbor and for other harbors and coastal areas affected by pollution and/or high organic matter

517 loads (Vezzulli *et al.*, 2003; Veiga *et al.*, 2009; Pusceddu *et al.*, 2011; Dal Zotto *et al.*, 2016;  
518 Semprucci *et al.*, 2016). The only exception was represented by the paucity of meiofaunal  
519 organisms found at PORT station. Considering that total meiofaunal abundance was positively  
520 linked to products derived from primary production (Chl-*a*, Phaeo, and chloroplast pigment  
521 equivalents - CPE), its low abundance at PORT station could be partially justified by the lowest  
522 detected value of 'fresh' material (Chl-*a*) and/or by a recent physical disturbance due to the  
523 position of this sampling station in an area defined of 'transit' within the harbor. Apart from Zn,  
524 all TE and PAH values detected at PORT station appeared overall lower if compared to those of  
525 the other sampling stations, suggesting a recent resuspension or removal of sediment. Given its  
526 small size, low mobility, and lack of dispersive life stages (Giere, 2009), meiofauna is more  
527 susceptible to within-habitat physical variability and environmental disturbances than larger,  
528 more mobile, and potentially more highly dispersed members of the macrofauna (Schratzberger  
529 *et al.*, 2008). This would explain the drop in meiofauna abundance, not reported for the  
530 macrofauna, at that sampling station. The meiofauna showed an overall good number of taxa  
531 (12) and in four sampling stations out of five the majority of taxa were represented. The  
532 measures of diversity (*i.e.* S, H', and J') were comparable to the values reported in harbor areas  
533 (*e.g.*, Moreno *et al.*, 2008a; Moreno *et al.*, 2008b) and in enclosed/transitional systems in the  
534 Adriatic Sea (*e.g.*, Pusceddu *et al.*, 2007; Pusceddu *et al.*, 2011; Frontalini *et al.*, 2014).  
535 However, the strong dominance of Nematoda justified the low values reported for the Pielou's  
536 equitability index, particularly at LR and DS stations. The dominance of the most resistant and  
537 adaptable group is a peculiarity of more stressed and less stable environments, such as harbors  
538 (Semprucci *et al.*, 2015). Abundance and diversity indices were correlated to different proxies of  
539 food sources (quantity and quality) into the sediment and to its grain size, confirming the effect  
540 of these environmental variables on meiofaunal populations (Balsamo *et al.*, 2010).  
541 As previously mentioned, the structure of the meiofaunal assemblages exhibited a clear spatial  
542 variability between inside vs. outside the harbor but also among the five sampling stations,  
543 consistent with changes detected for environmental features. The dissimilarity among the  
544 sampling stations was mainly due to some less abundant and more sensitive taxa, not present in  
545 the innermost stations. Usually, organisms that can cope with unfavorable conditions take over  
546 (*e.g.*, Nematoda), whereas more sensitive taxa disappear or become rare (Mirto *et al.*, 2014;  
547 Zeppilli *et al.*, 2015). In the case of Ancona harbor, Bivalvia, Kinorhyncha, Platyhelminthes,  
548 Sipuncula, and Tardigrada were found only at the outermost stations (LR, DS, and API) being  
549 identified as less tolerant taxa (Baldrighi *et al.*, 2019 and literature therein). Conversely, the more  
550 tolerant and widespread groups of Ciliata, Oligochaeta, and Polychaeta (Pusceddu *et al.*, 2007;  
551 Moreno *et al.*, 2008a; Moreno *et al.*, 2008b; Semprucci *et al.*, 2015) characterized the innermost  
552 stations, along with an important presence of soft-shelled Foraminifera inhabiting the sediment at  
553 all the investigated sampling stations. Soft-shelled monothalamous Foraminifera are an  
554 important component living in the sediment and populating the Adriatic Sea (Sabbatini *et al.*,  
555 2013), but most of the time this component is overlooked in meiofauna studies. The high  
556 presence of this group has been found to be associate to high values of Chl-*a*, eutrophic

557 conditions, and high variability of environmental parameters (*e.g.*, organic matter loads, salinity,  
558 temperature, oxygen content; Sabbatini *et al.*, 2013). Their strong tolerance and positive response  
559 to environmental stress (Sabbatini *et al.*, 2010) can perfectly justified their presence in the  
560 Ancona harbor. Changes in the community structure were supported by DistLM analysis, which  
561 revealed that pollutants and, secondly, the grain size could explain the variability in the  
562 meiofaunal composition.

563 Food sources did not have any effect on the meiobenthic community. According to Dell'Anno *et al.*  
564 *al.* (2002) and Pusceddu *et al.* (2009), the system of Ancona harbor can be ranked as eutrophic  
565 (inside) - mesotrophic (outside) with high quality of organic matter. Thus, food sources did not  
566 constitute a limiting factor for the meiofaunal community, as reported instead for oligotrophic  
567 systems (*e.g.*, Covazzi Harriague *et al.*, 2013) or in estuary's waters (Patricio *et al.*, 2012). Same  
568 results were reported in Franzo *et al.* (2022) analyzing the nematode communities inhabiting  
569 different Adriatic harbors, including that of Ancona. Authors showed that the main  
570 environmental factor that shaped the nematode assemblages in all harbors were the PAH  
571 concentration levels, while food sources and the grain size were much less relevant.  
572 Interestingly, some positive correlations between TEs and meiofaunal abundance and its related  
573 univariate measures were reported also in this study. In the study conducted by Cibic *et al.*  
574 (2017), authors pinpointed as heavy metal content may influence meiofaunal abundance and its  
575 composition. The positive nature of the correlation could be the result of a meiobenthic  
576 community well adapted to permanent stress conditions (Cibic *et al.*, 2017).

577

### 578 **Macrofaunal response to harbor environmental conditions**

579

580 Benthic macroinvertebrates are traditionally used as biological indicators of ecosystem health in  
581 the marine environment, especially infaunal assemblages associated with soft-bottom habitats  
582 (Borja *et al.*, 2003), and they act as integrators of stress over months to years (Weisberg *et al.*,  
583 1997; Paul *et al.*, 2001). Thus, the structure and spatial distribution of benthic communities can  
584 be directly linked with pollutant or disturbance exposure (Borja *et al.*, 2003). The macrobenthos  
585 in the internal parts of a harbor usually exhibits a low abundance and a numerical importance of  
586 pollution-tolerant species (Pearson & Rosenberg, 1978; Callier *et al.*, 2009). Species richness,  
587 abundance, biomass, diversity and equitability indices generally show a gradual increase from  
588 the interior to the exterior of an harbor, following an internal-external gradient (Callier *et al.*,  
589 2009).

590 In the present study, total macrofaunal abundance and its measures of diversity (*i.e.* S, H', and  
591 J'), fell within the range of values reported by Spagnolo *et al.* (2011) and Travizi *et al.* (2019) for  
592 Ancona harbor. Results here reported showed an overall increasing trend in macrofaunal  
593 abundance and species richness from inside to outside the study area; however LR station  
594 significantly differed from the other sampling stations when univariate measures were  
595 considered. Regarding the community structure, species composition was different enough to  
596 differentiate between the internal and the external sampling stations.

597 The macrobenthic community was mostly composed by the dominant groups of Anellida,  
598 Mollusca, and Crustacea, as usually reported from enclosed environments impacted by pollutants  
599 and characterized by high organic matter loads (Guerra-García & García-Gómez, 2004b;  
600 Spagnolo *et al.*, 2011; Travizi *et al.*, 2019). Among the group of Anellida, the Oligochaeta  
601 species *T. swirencoides* was identified for the first time in Ancona harbor and it was found to be  
602 particularly abundant in all the sampling stations and even dominant at LR station. Only at API  
603 station the species was absent. Tubificid oligochaetes, also called sludge worms, are very  
604 common in high polluted areas (Brusca & Brusca, 2003) and they are recognized as a pollution-  
605 tolerant taxon (Pelletier *et al.*, 2010). Thus, *T. swirencoides* was the species mostly responsible  
606 for the difference between the innermost sampling stations and the outermost one. Species  
607 composition may be affected by pollutant concentrations and high levels of organic matter,  
608 through a decrease in diversity and abundance of sensitive species (Callier *et al.*, 2009). The  
609 majority of the species found inhabiting the Ancona harbor sediments belonged to the ecological  
610 groups of disturbance-tolerant, second- and first-order opportunistic species (Borja *et al.*, 2000).  
611 All these species were typically soft-bottom species with the only exceptions of *Mytilus*  
612 *galloprovincialis* and *Hiatella arctica* found at LR station, as also reported in Spagnolo *et al.*  
613 (2011). These two species are commonly from hard-substrates and their origin may have been  
614 from the pier close to the LR station. Polychaeta species, along with the tubificid oligochaeta *T.*  
615 *swirencoides*, were the species most represented and diversified. Many Polychaeta species have a  
616 high level of tolerance to adverse effects such as pollution and natural perturbations (Borja *et al.*,  
617 2000), and for this reason they usually constitute the majority of benthic organisms living in  
618 harbor systems (Guerra-García & García-Gómez, 2004b). Usually, Polychaeta species richness  
619 and their diversity inside of harbor areas are low because of high pollution levels and the lack of  
620 oxygen in the water column (Estacio *et al.*, 1997; Dhainaut-Courtois *et al.*, 2000). The same  
621 trend of increasing Polychaeta diversity moving outside the harbor area was also reported in the  
622 present study, with tolerant and opportunistic species such as *Capitella capitata*, *C. capitata*,  
623 *H. filiformis*, *Sternaspis scutata* and *Streblospio* sp. particularly abundant in the innermost  
624 sampling stations. The species *Prionospio cirrifera*, mainly recorded outside the harbor (API  
625 station), is traditionally identified as an opportunistic spionid living in silty-clay sediments with  
626 high organic content (Borja *et al.*, 2000; Simonini *et al.*, 2004). Spagnolo *et al.* (2011) reported  
627 the same finding and authors explained this as a result of a scarce tolerance of *P. cirrifera* to  
628 copper, detected at a concentration 9 times higher at the innermost MAN station compared to the  
629 average concentration detected at all the other sampling stations. A similar consideration could  
630 arise for the high presence of the opportunistic species *S. bombyx* at API station. On the other  
631 hand, Polychaeta species ranked as disturbance-sensitive (Borja *et al.*, 2000), such as *Aricidea*  
632 *fragilis*, *Glycera capitata*, *Jasmineira elegans*, *Paradoneis armata*, and *Paraonis fulgens* have  
633 also been found inhabiting the most impacted sampling stations inside the harbor. Due to their  
634 economic and ecological importance, as well as their sedentary life, Mollusca have assumed a  
635 major role in monitoring contaminants worldwide (Pizzini *et al.*, 2015; Pizzini *et al.*, 2017;  
636 Grotti *et al.*, 2016). *Anadara transversa*, *Kurtiella bidentata*, *M. galloprovincialis*, and *Nucula*

637 *nitidosa* are defined as disturbance-tolerant species and they tended to dominate the innermost  
638 sampling stations in Ancona harbor, with the only exception for *K. bidentata*, particularly  
639 abundant at API station. *Abra alba* characterized LR and DS stations, confirming its preference  
640 for sandy sediments with medium-high levels of organic matter quantity and quality (Guerra-  
641 García & García-Gómez, 2004b). This species has been reported abundant in harbors affected by  
642 heavy metal pollution (Dhainaut-Courtois *et al.*, 2000).

643 A large number of crustaceans (Amphipoda, Isopoda, Tanaidacea) have been categorized as  
644 pollution-sensitive taxa, especially compared to Polychaeta (Pelletier *et al.*, 2010). Crustacean  
645 communities have been considered to be among the most sensitive to changes in environmental  
646 variables (Gómez-Gesteira & Dauvin, 2000), and for this reason crustacean species richness and  
647 diversity inside harbors are generally considerably low (Estacio *et al.*, 1997; Dhainaut-Courtois  
648 *et al.*, 2000). Three abundant species characterized the innermost sampling stations in Ancona  
649 harbor: the Amphipoda *Leptocheirus pilosus*, the Caprellida *Phtisica marina*, and the Tanaidacea  
650 *Apseudopsis latreillii*. *P. marina* and *A. latreillii* have been reported in high number in sediments  
651 containing less sand and high concentrations of N, P, Cu, and organic matter (Guerra-García &  
652 García-Gómez, 2004a). *A. latreillii* belongs to the group of species that may occur under normal  
653 conditions, but whose populations are stimulated by organic enrichment; while *P. marina*  
654 belongs to the group of species very sensitive to organic enrichment. According to the present  
655 study and the results of other previous investigations (Conradi *et al.*, 2000; Guerra-García &  
656 García-Gómez, 2001; Guerra-García & García-Gómez, 2004a) this species is able to live even in  
657 impacted habitats with moderate-high levels of heavy metals and PAHs. Conversely, the more  
658 sensitive Amphipoda *Ampelisca diadema* dominated the crustacean assemblages at API station.

659 As for the meiofauna, pollutant content and the different sediment texture inside and outside the  
660 Ancona harbor affected the macrofaunal composition, as previously reported (Guerra-García and  
661 García-Gómez, 2001; Guerra-García & García-Gómez, 2004a; Guerra-García & García-Gómez,  
662 2004b; Spagnolo *et al.*, 2011; Travizi *et al.*, 2019). Univariate descriptors as well as the analysis  
663 of species characterizing the benthic communities indicated the presence of modified, but quite  
664 diverse and presumably well-established soft-bottom communities in all the investigated  
665 sampling stations. This might reflect the successful adaptation of many pollution-tolerant species  
666 to the long-term pollution and unstable environmental conditions of Ancona harbor (Travizi *et*  
667 *al.*, 2019). The dominance of opportunistic and tolerant species confirmed the results reported  
668 from DistLM analysis, which identified pollution as important driver of macrobenthic  
669 assemblage structure.



670

## 671 Conclusions

672 The benthic community represents a source of information at different food-web levels and can  
673 be utilized to investigate and characterize the habitat where the community exists. As meiofauna  
674 and macrofauna have different ecological roles in marine ecosystems, they may respond to  
675 environmental features and stress conditions at different spatial and temporal scales (Frontalini *et*  
676 *al.*, 2011).

677 We initially hypothesized a different response from the two benthic size classes to the different  
678 environmental conditions characterizing the Ancona harbor. Meio- and macrofauna provided  
679 partially similar and complementary types of information depending on the indices used  
680 (univariate measures or community structure/ species composition) and the different ‘response-  
681 to-stress’. The following considerations emerged:

- 682 - In Ancona harbor the presence of pollutants and the sediment type determined the meio- and  
683 macrofaunal community structure. Moreover, the meiofauna was affected by the quality and  
684 quantity of organic matter, suggesting that meiobenthic assemblages were more receptive to  
685 within-habitat food variability than macrofauna. Meiofauna results were generally  
686 statistically more significant than those obtained for macrofauna. Meiofauna were likely to  
687 be more sensitive to the effects of environmental features and contaminants than  
688 macrofauna.
- 689 - Both invertebrate groups were characterized by distinctive assemblages across the harbor,  
690 consistent with changes detected for environmental features. However, for the macrofaunal  
691 communities a gradient rather than a clear separation among the sampling stations was  
692 reported, while a clear separation among them and moving from inside to outside the harbor  
693 area was detected for the meiofaunal communities.
- 694 - This investigation confirmed the fundamental advantage of a multi-benthic size approach,  
695 with the inclusion of many taxonomic and functional groups. Optimally, both groups should  
696 be used in marine pollution monitoring programs included in the EU Marine Strategy  
697 Framework Directive (MSFD; Directive 2008/56/EC) in the context of its Descriptor 1  
698 ‘maintenance of biodiversity’ and Descriptor 6 ‘sea floor integrity’.

699

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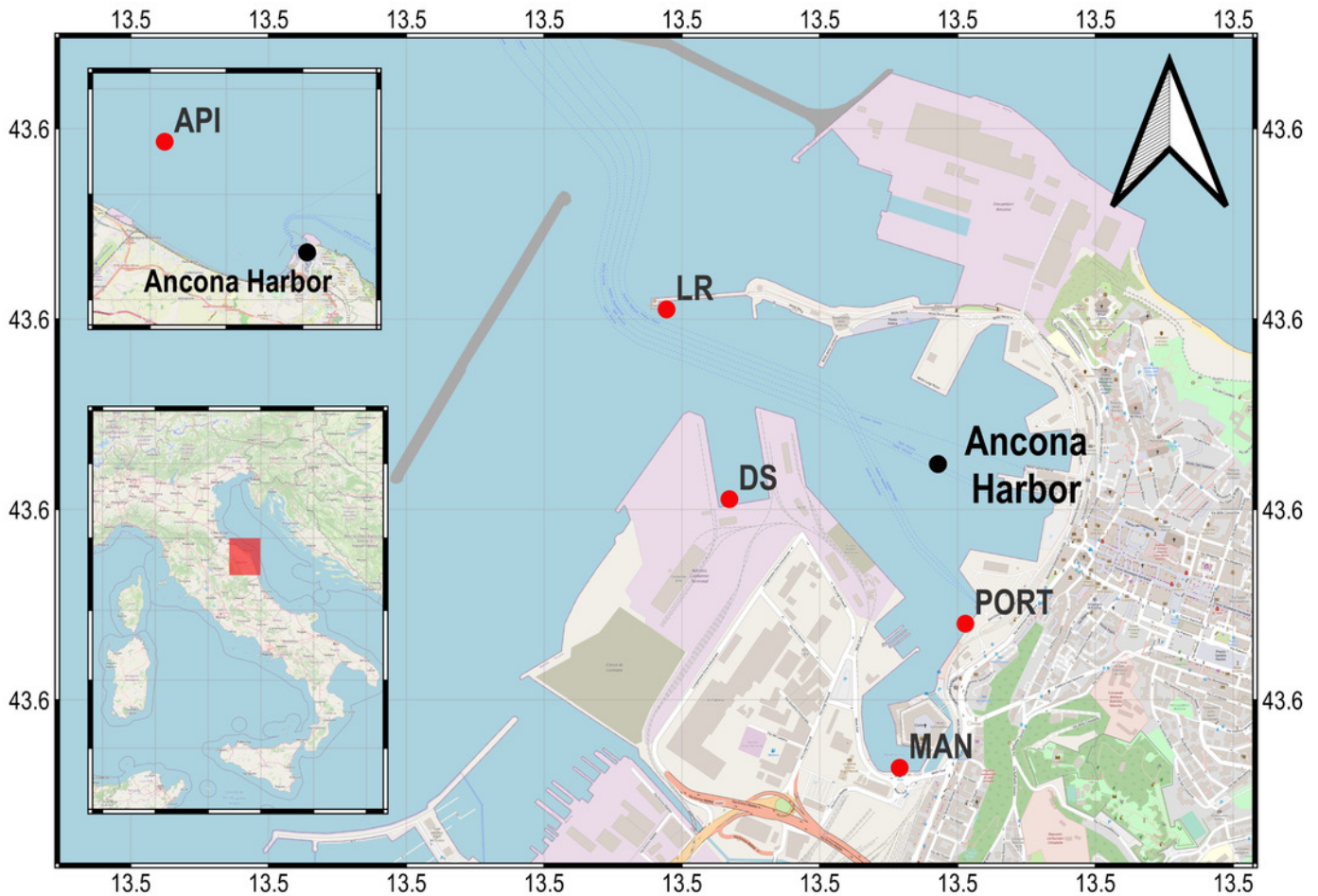
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# Figure 1

Map of the sampling area (Ancona harbor) and location of the five sampling stations.

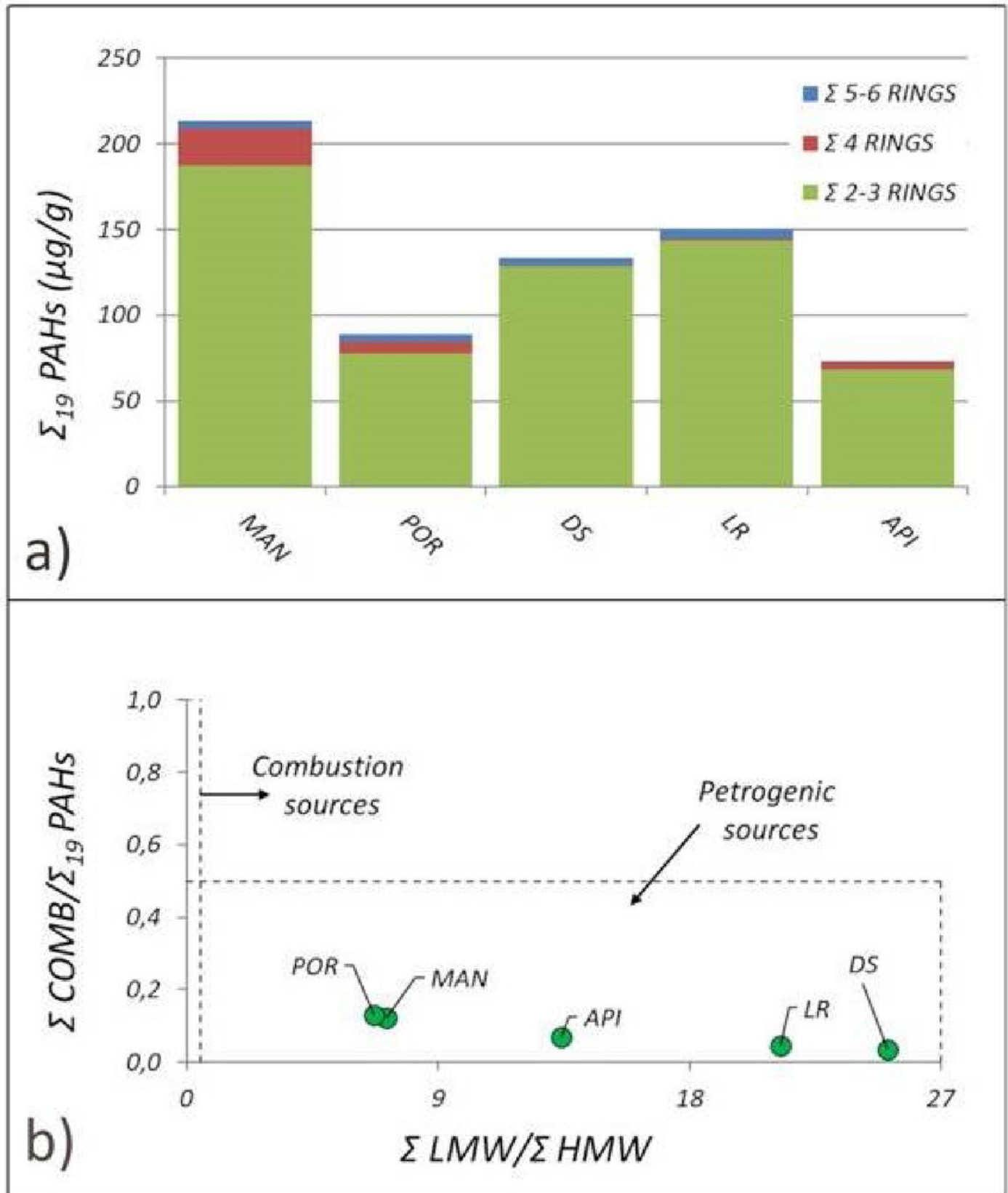
The pink rectangle indicates the geographical position of the Ancona harbor, Italy.



## Figure 2

(a) Distribution pattern and (b) origin of Polycyclic aromatic hydrocarbons (PAHs) characterizing the sediment at the investigated sampling stations.

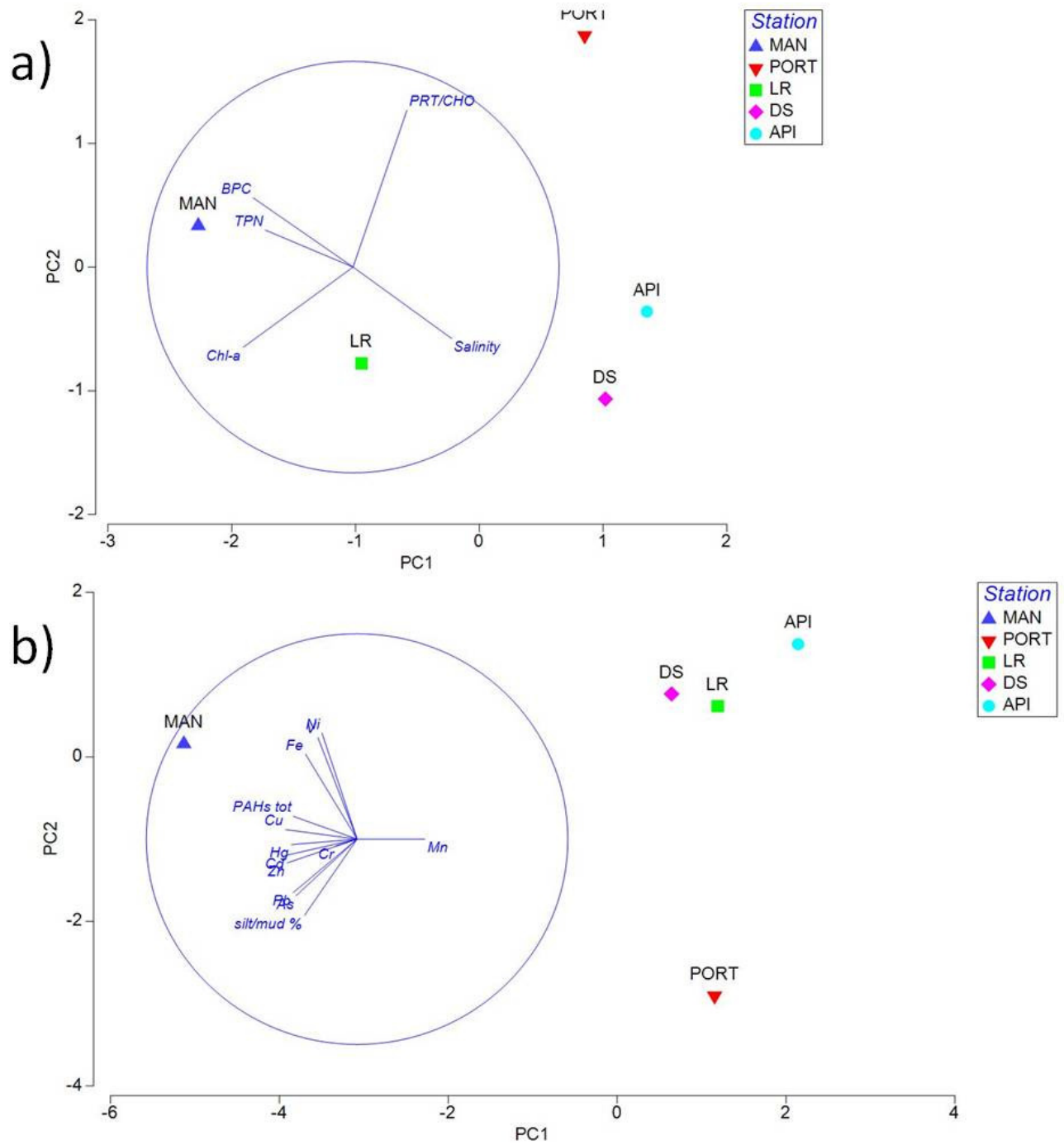
$\Sigma$  LMW = sum of Low Molecular Weight compounds (Naphthalene, 1-Methylnaphtalene, 2-Methylnaphtalene, Acenaphthylene, Acenaphthene, Fluorene, Phenanthrene, Anthracene);  $\Sigma$  HMW = sum of High Molecular Weight compounds (Fluoranthene, Pyrene, Benz[*a*]anthracene, Chrysene, Benzo[*b*]fluoranthene, Benzo[*k*]fluoranthene, Benzo[*a*]pyrene, 7,12-Dimethylbenz[*a*]anthracene, Benzo[*ghi*]perylene, Indeno[1,2,3-*cd*]pyrene, Dibenz[*a,h*]anthracene);  $\Sigma$  COMB = sum of 9 non-alkylated PAHs (Fluoranthene, Pyrene, Benz[*a*]anthracene, Chrysene, Benzo[*b*]fluoranthene, Benzo[*k*]fluoranthene, Benzo[*a*]pyrene, Benzo[*ghi*]perylene, Indeno[1,2,3-*cd*]pyrene).



## Figure 3

Principal Component Analysis (PCA) with (a) 'biotic' and (b) 'abiotic' environmental variables.

BPC = Biopolymeric organic carbon; Chl-*a* = Chlorophyll-*a* concentration; PAHs = Polycyclic aromatic hydrocarbons; PRT/CHO = Protein to carbohydrate ratio; TPN = Total Prokaryotic Number.

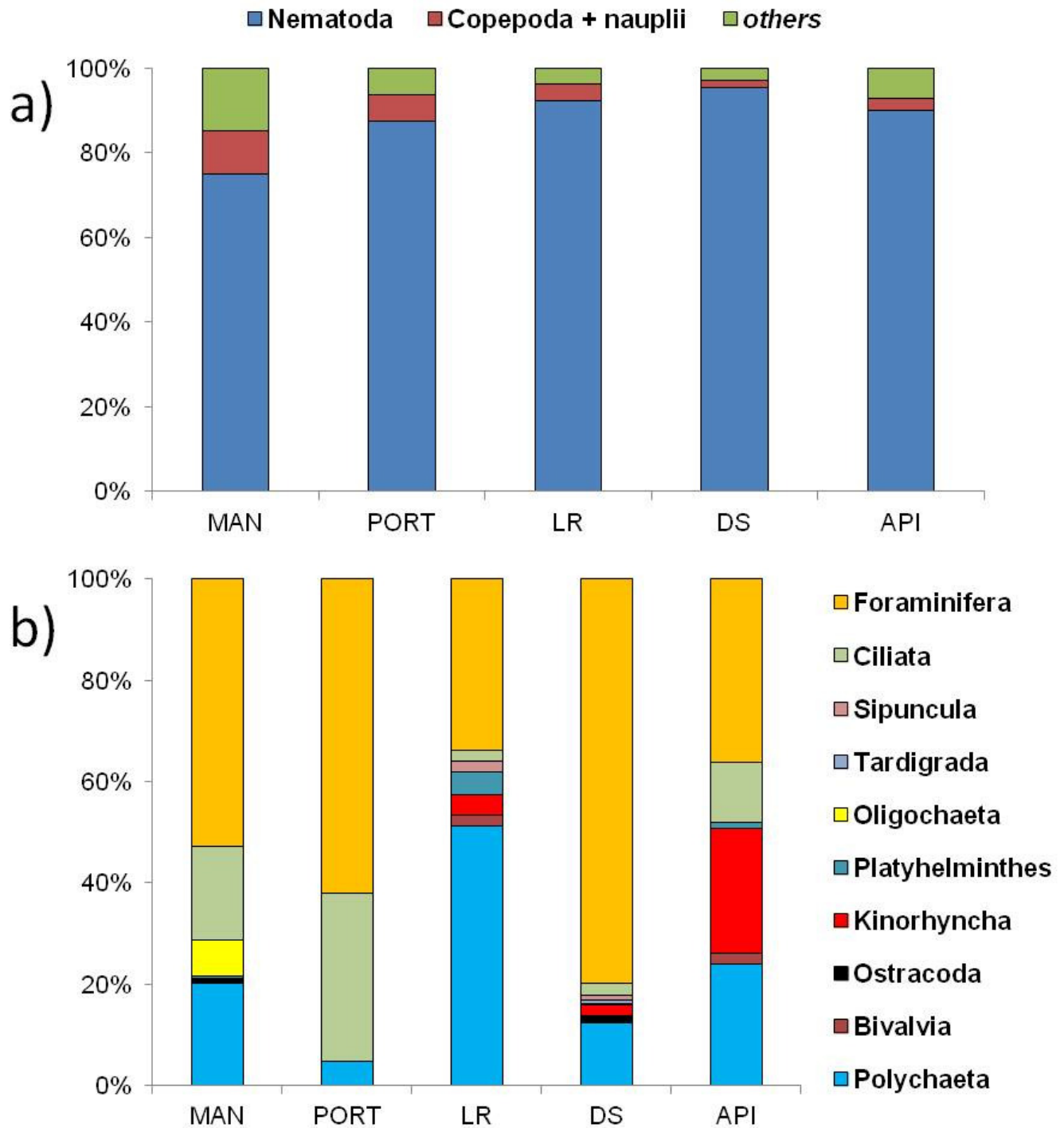


## Figure 4

(a) Meiofaunal community structure and (b) contribution of taxa *others* at each sampling station.

Mean values are shown.

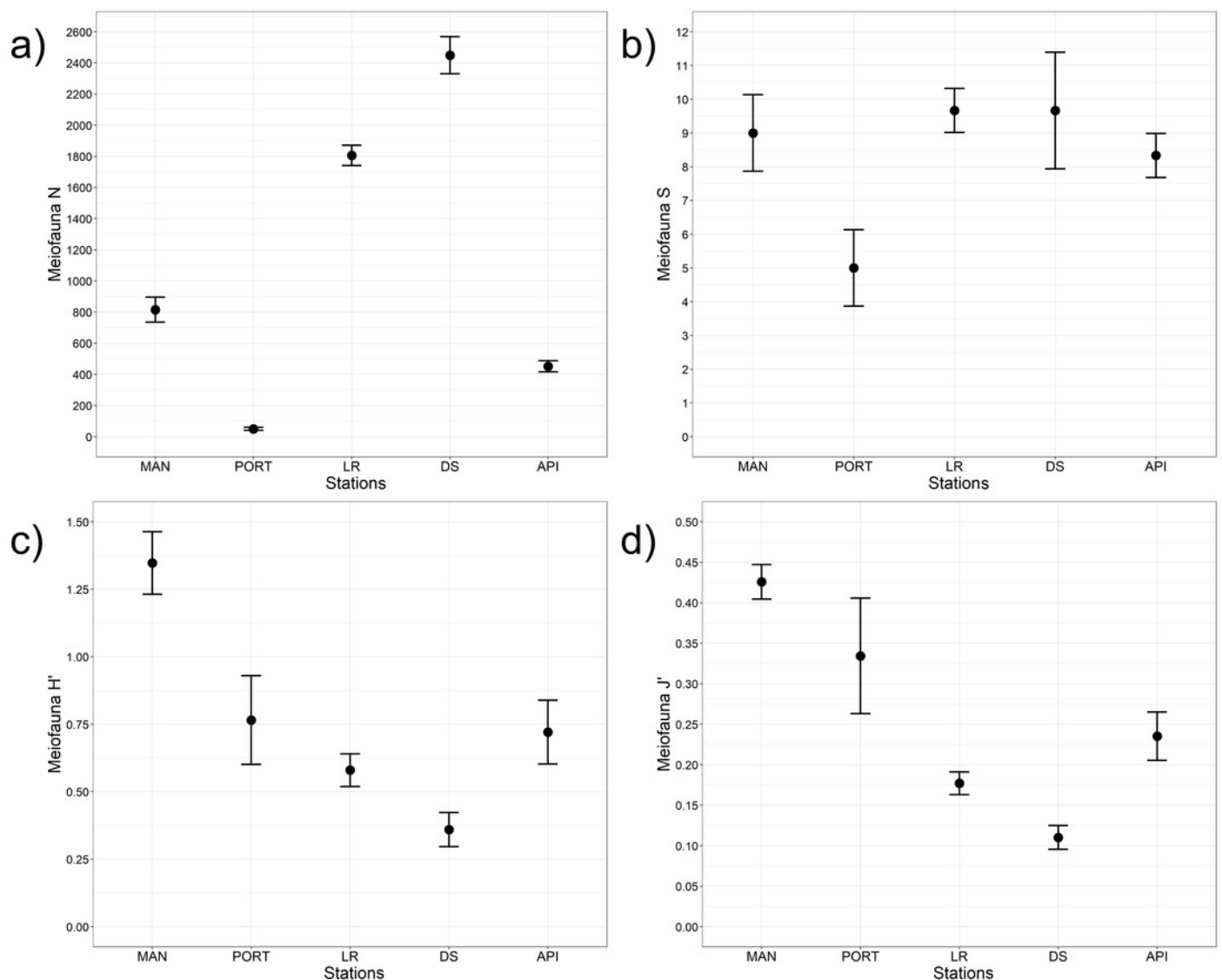




## Figure 5

Meiofaunal univariate measures.

(a) Meiofauna abundance ( $N$  = no. of individuals  $10\text{ cm}^{-2}$ ) and its diversity indices: (b) meiofauna taxa richness ( $S$ ), (c) Shannon's diversity index ( $H'$ , based on  $\log_2$ ), and (d) Pielou's equitability index ( $J'$ ). Bars represent 95% confidence intervals.




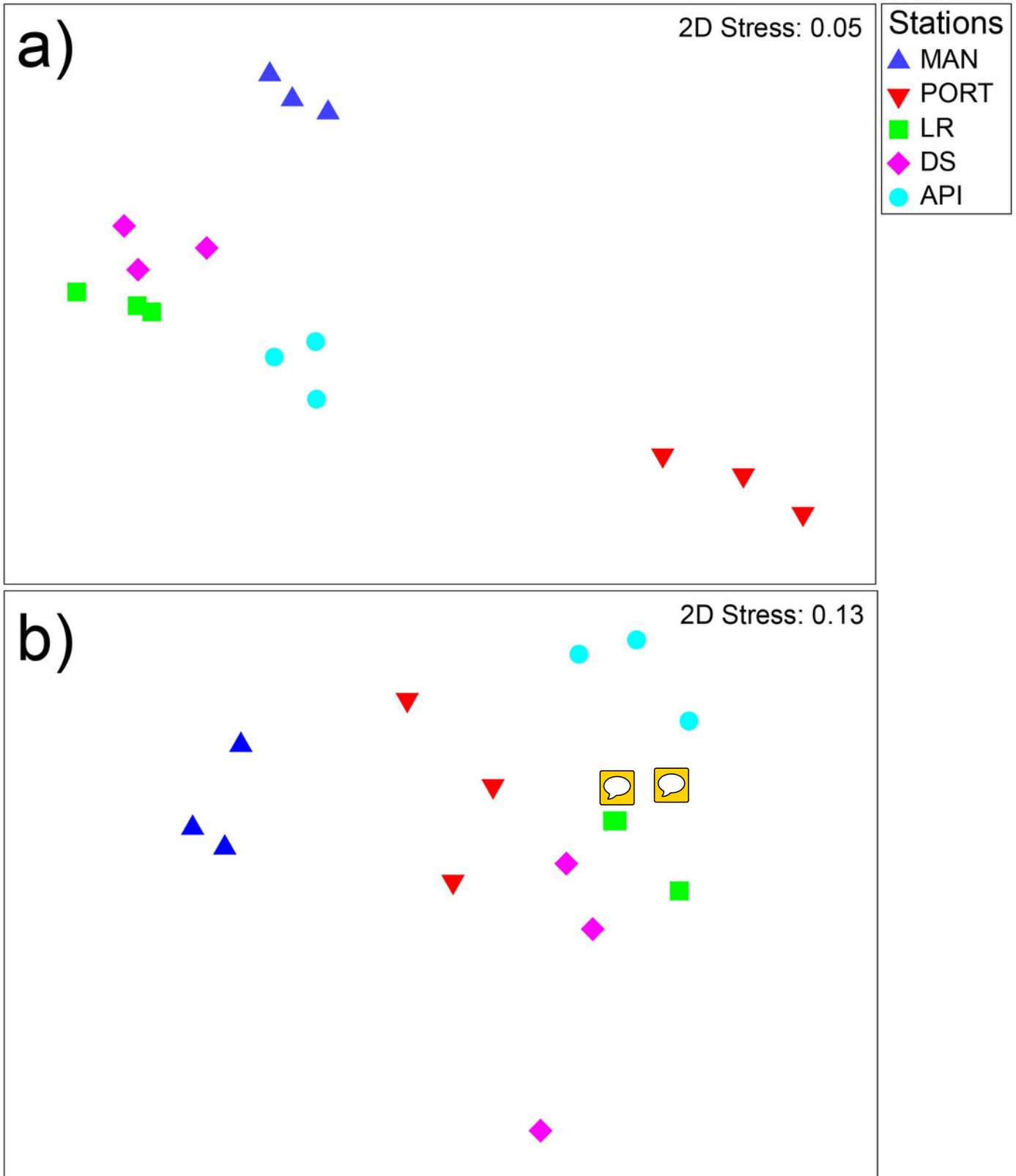


## Figure 6

Non-metric Multidimensional Scaling (nMDS) of benthic communities.

Non-metric Multidimensional Scaling (nMDS) plots on (a) meiobenthic and (b) macrobenthic community structures characterizing the sediment at the investigated sampling stations.

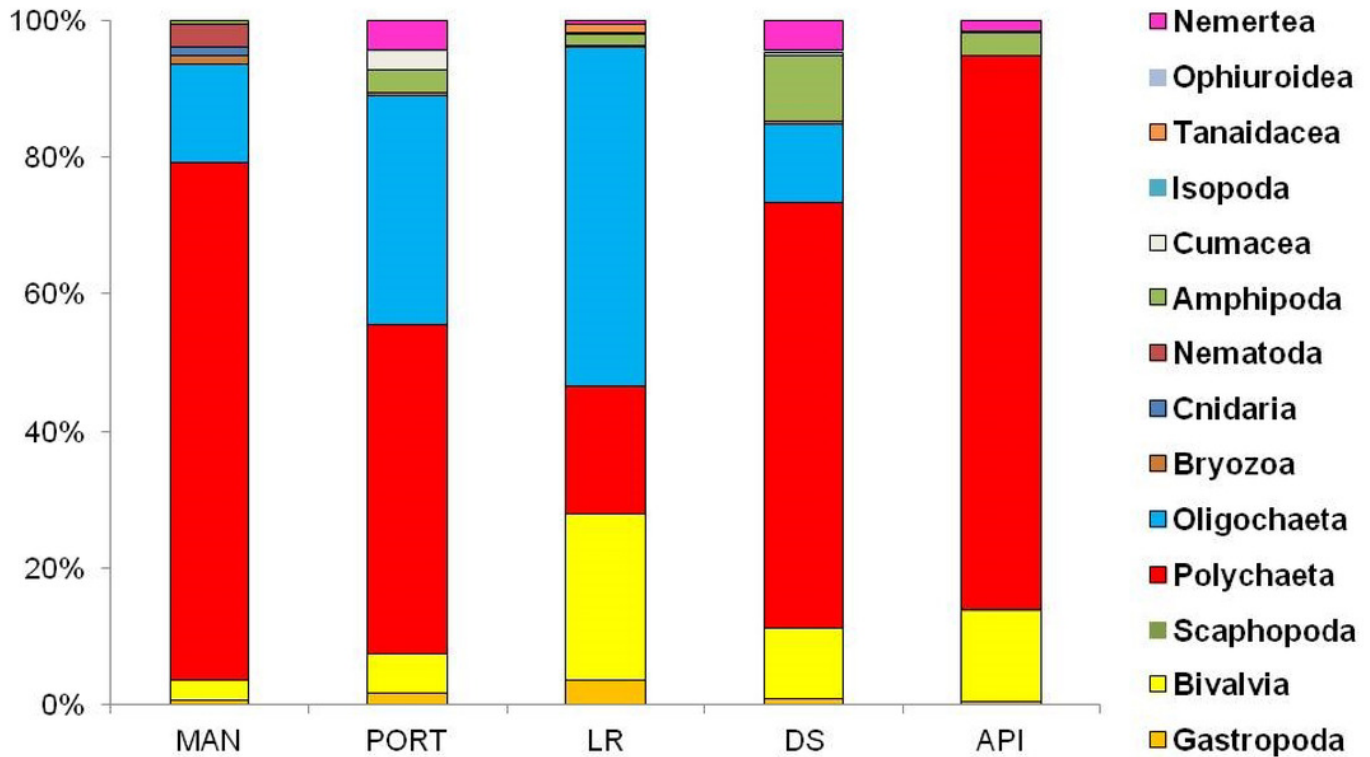
Green circles represent 40%  ilarity grouping.



## Figure 7

Macrofaunal community structure at each sampling station.

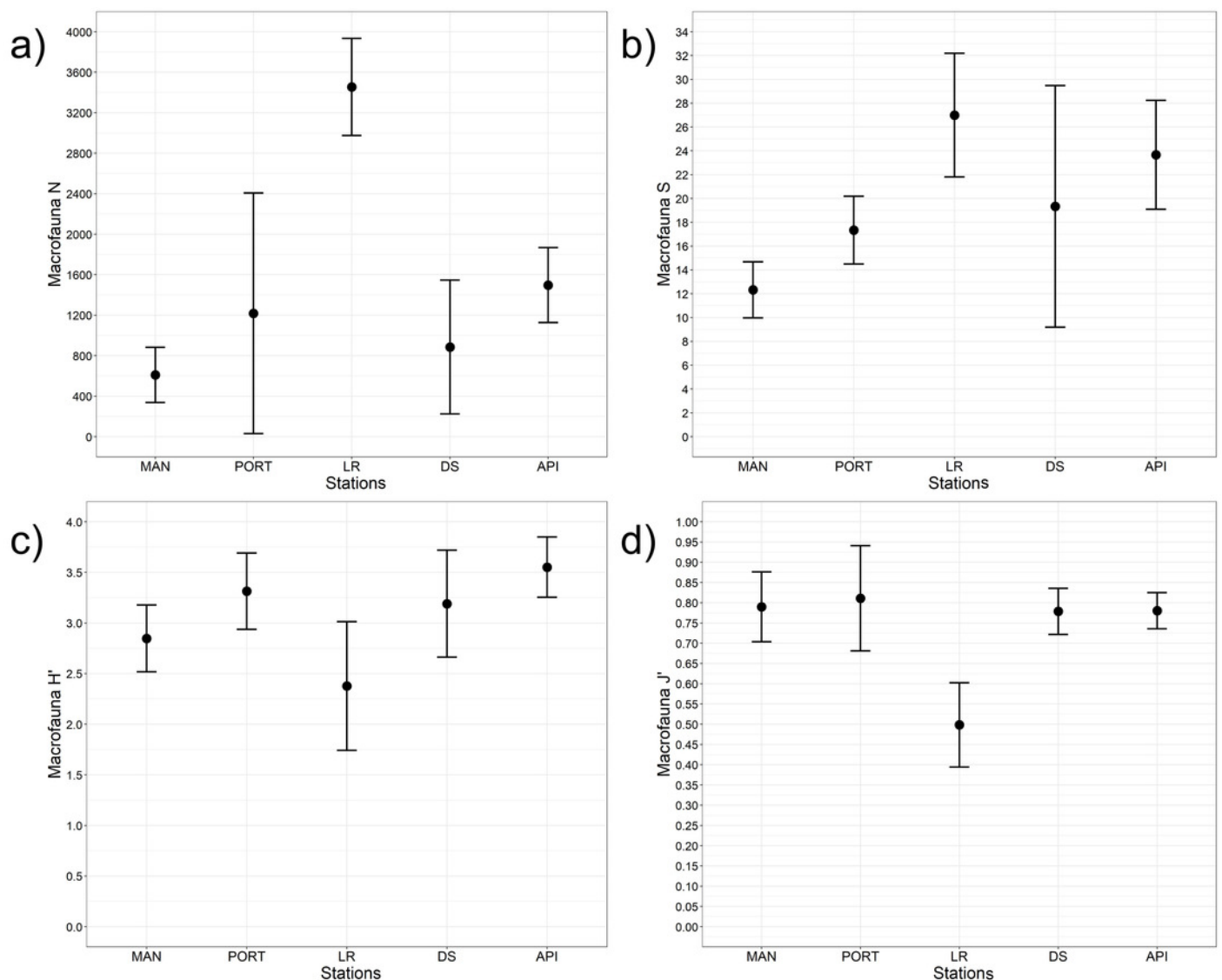
Mean values are shown.



## Figure 8

Macrofaunal univariate measures.

(a) Macrofauna abundance (N = no. of individuals  $m^{-2}$ ) and its diversity indices: (b) macrofauna taxa richness (S), (c) Shannon's diversity index ( $H'$ , based on  $\log_2$ ), and (d) Pielou's equitability index ( $J'$ ). Bars represent 95% confidence intervals.



**Table 1** (on next page)

Table 1. Location of the five sampling stations, environmental variables measured during the sampling, grain size, biochemical composition, and microbial component of the sediment samples.

Chl-*a* = Chlorophyll-*a* concentration; BPC = Biopolymeric organic carbon; PRT/CHO = Protein to carbohydrate ratio; TPN = Total Prokaryotic Number. Mean values  $\pm$  standard deviation are reported.

- 1 **Table 1. Location of the five sampling stations, environmental variables measured during the**  
 2 **sampling, grain size, biochemical composition, and microbial component of the sediment samples.**  
 3 Chl-*a* = Chlorophyll-*a* concentration; BPC = Biopolymeric organic carbon; PRT/CHO = Protein to  
 4 carbohydrate ratio; TPN = Total Prokaryotic Number. Mean values  $\pm$  standard deviation are reported.

Sampling station		MAN	PORT	DS	LR	API
Latitude N		43°36.793	43°37.020	43°37.130	43°37.309	43°40.297
Longitude E		13°30.174	13°30.317	13°29.482	13°29.400	13°24.344
Water temperature (°C)		10.4	10.2	10.2	10.1	10.7
Salinity (PSU)		32.0	32.5	36.7	31.3	37.6
Depth (m)		3.1	6.2	11.0	5.0	14.0
Grain size	Sand%	2	20	84	84	79
	Silt/mud%	98	80	16	16	21
Chl- <i>a</i> ( $\mu\text{g g}^{-1}$ )		13.9 $\pm$ 5.2	2.3 $\pm$ 0.4	17.0 $\pm$ 4.6	8.1 $\pm$ 1.2	2.9 $\pm$ 0.6
BPC (mgC g <sup>-1</sup> )		10.2 $\pm$ 1.8	5.7 $\pm$ 1.5	3.7 $\pm$ 0.3	4.5 $\pm$ 0.5	2.3 $\pm$ 0.5
PRT/CHO		5.5 $\pm$ 1.5	40 $\pm$ 16.1	5.2 $\pm$ 1.4	4.2 $\pm$ 1.5	12.9 $\pm$ 2.7
TPN (no. of cells mL <sup>-1</sup> )		1.8E+06 $\pm$ 4.0E+05	8.4E+05 $\pm$ 7.4E+05	1.0E+06 $\pm$ 6.1E+05	3.8E+05 $\pm$ 3.7E+05	1.3E+06 $\pm$ 1.2E+05

5

6 **Table 2.** Concentration (in  $\mu\text{g g}^{-1}$ ) of selected Trace Elements (TEs) and Polycyclic aromatic  
 7 **hydrocarbons (PAHs)** detected in the sediment samples. In bold values exceeding the lower threshold  
 8 level (L1) specified in the Ministerial Decree 173/2016, in bold and underlined those values exceeding

Sampling station		MAN	PORT	DS	LR	API
TEs	Arsenic	3.628	3.193	2.941	2.882	2.386
	Cadmium	0.1915	0.0773	0.0533	0.0523	0.0493
	Chromium	23.07	22.14	26.54	21.17	19.28
	Copper	<b>46.313</b>	3.974	7.866	5.331	3.393
	<b>Iron</b>	19119	12332	16709	15666	15937
	Mercury	0.06835	0.02630	0.01865	0.01117	0.02761
	<b>Manganese</b>	355.7	508.0	449.2	507.0	558.4
	Nickel	3.911	2.786	3.555	3.539	3.703
	Lead	19.912	14.835	10.569	11.799	8.348
	Vanadium	28.31	18.99	24.90	24.83	26.09
	Zinc	<b><u>297.15</u></b>	<b>114.94</b>	58.78	67.70	47.03
PAHs	Naphthalene	<b>64.483</b>	21.611	<b>60.266</b>	<b>63.414</b>	30.370
	1-Methylnaphtalene	51.412	21.632	29.975	36.011	17.875
	2-Methylnaphtalene	50.917	24.928	27.438	31.036	13.150
	Acenaphthylene	8.267	<0.5	<0.5	<0.5	<0.5
	Acenaphthene	<0.5	<0.5	<0.5	<0.5	<0.5
	Fluorene	<0.5	1.578	1.280	1.881	1.428
	Phenanthrene	12.184	7.579	9.029	10.800	5.302
	Anthracene	<0.05	0.166	<0.05	0.123	0.163
	Fluoranthene	17.404	6.305	<0.05	0.861	4.022
	Pyrene	4.039	0.787	0.782	0.556	0.172
	Benz[ <i>a</i> ]anthracene	<0.05	<0.05	<0.05	<0.05	<0.05
	Chrysene	<0.05	<0.05	<0.05	<0.05	<0.05
	Benzo[ <i>b</i> ]fluoranthene	3.027	3.363	1.814	3.004	0.475
	Benzo[ <i>k</i> ]fluoranthene	0.308	0.373	1.220	1.114	0.322
	Benzo[ <i>a</i> ]pyrene	1.145	0.332	0.281	0.882	0.099
	7,12-Dimethylbenz[ <i>a</i> ]anthracene	<0.5	<0.5	<0.5	<0.5	<0.5
	Benzo[ <i>ghi</i> ]perylene	0.257	0.376	0.595	0.326	<0.05
	Indeno[1,2,3- <i>cd</i> ]pyrene	<0.05	<0.05	<0.05	<0.05	<0.05
	Dibenz[ <i>a,h</i> ]anthracene	<0.05	<0.05	0.411	<0.05	<0.05
	$\Sigma_8$ LMW PAHs	187.26	77.49	127.99	143.27	68.29
$\Sigma_{11}$ HMW PAHs	26.18	11.54	5.10	6.74	5.09	
$\Sigma_{19}$ PAHs	213.44	89.03	133.09	150.01	73.38	

9 also the upper threshold level (L2).

10 LMW = Low Molecular Weight compounds (Naphthalene, 1-Methylnaphtalene, 2-Methylnaphtalene,  
 11 Acenaphthylene, Acenaphthene, Fluorene, Phenanthrene, Anthracene); HMW = High Molecular Weight compounds

12 (Fluoranthene, Pyrene, Benz[*a*]anthracene, Chrysene, Benzo[*b*]fluoranthene, Benzo[*k*]fluoranthene,  
13 Benzo[*a*]pyrene, 7,12-Dimethylbenz[*a*]anthracene, Benzo[*ghi*]perylene, Indeno[1,2,3-*cd*]pyrene,  
14 Dibenz[*a,h*]anthracene).



15 **Table 3. One-way Analysis of Variance (ANOVA) results on total meiofauna and macrofauna**  
 16 **abundance (N) and their diversity indices.** Number of taxa (S), Shannon's diversity index (H', based on  
 17  $\log_2$ ), and Pielou's equitability index (J'), characterizing the sediment at the investigated sampling

<b>Meiofauna</b>					
<b>Index</b>		<b>d.f.</b>	<b>F value</b>	<b>p value</b>	<b>Tukey's HSD <i>post hoc</i> test</b>
<b>N</b>	Stations	4	714.3	<0.001	DS>LR>MAN>API>PORT
	Residuals	10			
<b>S</b>	Stations	4	11.3	<0.001	PORT<API=DS=LR=MAN
	Residuals	10			
<b>H'</b>	Stations	4	44.1	<0.001	MAN>PORT>API=LR(=DS)>DS
	Residuals	10			
<b>J'</b>	Stations	4	41.6	<0.001	MAN>PORT=API=LR(=DS)>DS
	Residuals	10			
<b>Macrofauna</b>					
<b>Index</b>		<b>d.f.</b>	<b>F value</b>	<b>p value</b>	<b>Tukey's HSD <i>post hoc</i> test</b>
<b>N</b>	Stations	4	10.6	0.002	LR>API=DS=MAN=PORT
	Residuals	10			
<b>S</b>	Stations	4	3.8	0.041	MAN<LR
	Residuals	10			
<b>H'</b>	Stations	4	8.2	0.003	LR<API=DS=MAN=PORT
	Residuals	10			
<b>J'</b>	Stations	4	3.9	0.036	LR<API
	Residuals	10			

18 stations.

19

20 d.f. = degrees of freedom; HSD = Honestly Significant Difference.

- 21 **Table 4. Permutational Multivariate Analysis of Variance (PERMANOVA) and pairwise test and**  
 22 **results on total meiofaunal community composition. In bold significant values are reported.**

<b>PERMANOVA</b>						
<b>Source</b>	<b>d.f.</b>	<b>SS</b>	<b>MS</b>	<b>Pseudo-F value</b>	<b>P(perm)</b>	<b>Unique permutations</b>
Stations	4	7315.8	1829	38.037	<b>0.0001</b>	9915
Residuals	10	480.84	48.084			
Total	14	7796.7				
<b>PAIRWISE TEST</b>						
<b>Groups</b>	<b>t value</b>	<b>P(perm)</b>	<b>P(MC)</b>			
MAN, PORT	7.39	0.103	<b>0.0006</b>			
MAN, LR	8.2125	0.0993	<b>0.0006</b>			
MAN, DS	4.8631	0.0997	<b>0.0043</b>			
MAN, API	6.1192	0.0992	<b>0.0015</b>			
PORT, LR	8.233	0.0953	<b>0.0002</b>			
PORT, DS	6.4351	0.1013	<b>0.0004</b>			
PORT, API	5.3464	0.0985	<b>0.0019</b>			
LR, DS	2.9523	0.1014	<b>0.0163</b>			
LR, API	5.2028	0.1014	<b>0.0018</b>			
DS, API	3.9778	0.0985	<b>0.0054</b>			

- 23 d.f. = degrees of freedom; SS = Sum of Squares; MS = Mean Square; P(perm) = Permutation  $p$ -value; P(MC) =  
 24 Monte Carlo  $p$ -value.

25 **Table 5. Distance-based Linear Modeling (DistLM) analysis on meiofaunal community composition**  
 26 **characterizing the sediment at the investigated sampling stations.** Variables are coded after Varimax  
 27 Rotated PCA Axis as follow: ARCx = Abiotic variables associated to the rotated axis (x indicates the  
 28 number of the axis); BRCx = Biotic variables associated to the rotated axis. For a complete list of the  
 29 variables refers to the text.

<b>START SOLUTION</b>							
<b>AICc</b>	<b>R<sup>2</sup></b>	<b>RSS</b>	<b>No. of variables</b>	<b>Selections</b>			
82.237	0.9374	488.09	6	All			
<b>SEQUENTIAL TESTS</b>							
<b>Variable</b>	<b>AICc</b>	<b>SS</b>	<b>Pseudo-F value</b>	<b>p value</b>	<b>Prop.</b>	<b>Cumul.</b>	<b>Res. d.f.</b>
BRC1	7.390	146.94	2.4083	0.0916	1.88E-02	0.91855	9
BRC2	8.213	78.41	1.1113	0.4061	1.01E-02	0.90849	10
<b>BEST SOLUTION</b>							
<b>AICc</b>	<b>R<sup>2</sup></b>	<b>RSS</b>	<b>No. of variables</b>	<b>Selections</b>			
74.597	0.9085	713.44	4	ARC1- ARC4			

30 AICc = Akaike's Information Criterion corrected for small samples; R<sup>2</sup> = Coefficient of determination; RSS =  
 31 Residuals Sums of Squares; SS = Sums of Squares; Prop. = Proportion of variation explained by the variable;  
 32 Cumul. = Cumulative total of Prop; Res. d.f. = Residual degrees of freedom.

33

34 **Table 6. Permutational Multivariate Analysis of Variance (PERMANOVA) and pairwise test and**  
 35 **results on total macrofaunal community composition.** In bold significant values are reported.

<b>PERMANOVA</b>						
<b>Source</b>	<b>d.f.</b>	<b>SS</b>	<b>MS</b>	<b>Pseudo-F value</b>	<b>P(perm)</b>	<b>Unique permutations</b>
Stations	4	23534	5883.5	4.5481	<b>0.0001</b>	9915
Residuals	10	12936	1293.6			
Total	14	36470				
<b>PAIRWISE TEST</b>						
<b>Groups</b>	<b>t value</b>	<b>P(perm)</b>	<b>P(MC)</b>			
MAN, PORT	2.0289	0.1017	<b>0.028</b>			
MAN, LR	2.6607	0.0979	<b>0.011</b>			
MAN, DS	2.4096	0.0933	<b>0.015</b>			
MAN, API	3.0865	0.1047	<b>0.007</b>			
PORT, LR	1.5617	0.1021	0.100			
PORT, DS	1.708	0.1013	0.069			
PORT, API	2.0282	0.1012	<b>0.031</b>			
LR, DS	1.5713	0.1040	0.106			
LR, API	2.2314	0.1007	<b>0.024</b>			
DS, API	2.034	0.0997	<b>0.036</b>			

36 d.f. = degrees of freedom; SS = Sum of Squares; MS = Mean Square; P(perm) = Permutation  $p$ -value; P(MC) =  
 37 Monte Carlo  $p$ -value.

38 **Table 7. Distance-based Linear Modeling (DistLM) analysis on macrofaunal community**  
 39 **composition characterizing the sediment at the investigated sampling stations.** Variables are coded  
 40 after Varimax Rotated PCA Axis as follow: ARC<sub>x</sub> = Abiotic variables associated to the rotated axis (x  
 41 indicates the number of the axis); BRC<sub>x</sub> = Biotic variables associated to the rotated axis. For a complete  
 42 list of the variables refers to the text.

<b>START SOLUTION</b>							
<b>AICc</b>	<b>R<sup>2</sup></b>	<b>RSS</b>	<b>No. of variables</b>	<b>Selections</b>			
130.09	0.6749	11856	6	All			
<b>SEQUENTIAL TESTS</b>							
<b>Variable</b>	<b>AICc</b>	<b>SS</b>	<b>Pseudo-F value</b>	<b>p value</b>	<b>Prop.</b>	<b>Cumul.</b>	<b>Res. d.f.</b>
BRC1	124.52	1625.6	1.0969	0.3775	4.46E-02	0.63033	9
ARC3	121.01	2259.5	1.5084	0.1502	6.20E-02	0.56838	10
BRC2	118.28	2173.9	1.3810	0.1845	5.96E-02	0.50877	11
ARC1	116.72	2906.3	1.7845	0.0484	7.97E-02	0.42908	12
<b>BEST SOLUTION</b>							
<b>AICc</b>	<b>R<sup>2</sup></b>	<b>RSS</b>	<b>No. of variables</b>	<b>Selections</b>			
116.72	0.42908	20821	2	ARC2;ARC4			

43 AICc = Akaike's Information Criterion corrected for small samples; R<sup>2</sup> = Coefficient of determination; RSS =  
 44 Residuals Sums of Squares; SS = Sums of Squares; Prop. = Proportion of variation explained by the variable;  
 45 Cumul. = Cumulative total of Prop; Res. d.f. = Residual degrees of freedom.