1	Mesoscale assemblages of fish and megainvertebrates as evidence of
2	benthiscapes on continental shelves.
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19	Abstract
20	Despite the relatively small proportion of ocean surface they represent, continental
21	shelf ecosystems are among the most productive in the world. Located at the interface
22	between terrestrial and marine environments, these habitats are structured by strong

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environmental forcings, especially on the sea bottom. A clear understanding of the spatial 23 24 distribution of these habitats, along with knowledge on the composition and functioning of 25 their associated communities, is essential for fisheries management and ecosystem 26 conservation. Here, we used data from yearly EVHOE otter trawl surveys (2008-2020) to 27 characterize the spatial structuration of benthic communities of the entire continental shelf 28 of the Bay of Biscay (France), and to investigate the potential environmental drivers of 29 these patterns. Two separate biological components were studied: epibenthic 30 megainvertebrates and bentho-demersal fish. Clustering analyses identified seven 31 assemblages within each component. We detected a strong correlation between the 32 spatial organization of the different assemblages identified for fish and megainvertebrates, providing evidence for broad-scale spatial structuration of benthic habitats-33 34 benthiscapes—in this shelf ecosystem. The most influential environmental variables were 35 identified as bottom temperature, sediment type, and primary production. Patterns in 36 certain structural parameters, such as biomass, revealed possible spatial differences in 37 ecological functioning. For example, we observed a drop in biomass from the coast to the 38 central part of the shelf, followed by an increase in biomass near the edge of the Armorican 39 shelf. These patterns reflect major large-scale processes (river inputs versus shelf-break 40 upwelling) structuring the entire Bay of Biscay ecosystem. A comparative analysis 41 revealed that the biological features and functioning observed in this study are shared with other European continental shelves. In addition to improving our knowledge of benthic 42 43 environments, studies such as this one can promote improvements in ecosystem-based management and marine spatial planning of a fast-changing ecosystem under multiple 44 anthropogenic stresses. 45

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- 47 Keywords: Continental shelf, Bay of Biscay, Benthos, Demersal, Environmental
- 48 forcings, Spatial structure

49 **1** INTRODUCTION

Continental shelves occupy a small proportion of the total ocean surface (review in Hall, 50 51 2002) but are among the most productive ecosystems in the world (Costanza et al., 1997). 52 In addition, the ecosystem services they provide are among the most valuable (e.g., 53 carbon storage, fisheries, biogeochemical cycling) (Costanza et al., 1997; Longhurst et 54 al., 1995; Raffaelli et al., 2003). Within these environments, the benthic domain is of 55 considerable importance: it is responsible for a significant proportion of marine biological 56 production, especially the part exploited by humans (Hall, 2002), it makes an essential 57 contribution to bentho-pelagic coupling, and it represents a major reservoir for terrestrial 58 carbon storage (Zhang et al., 2021). The spatial location of these ecosystems-at the 59 interface between terrestrial and marine environments-places them under strong pressure from natural processes that vary in time and space, and exposes them to 60 extensive anthropogenic influences (Halpern et al., 2008; Worm et al., 2006). These 61 62 intense and dynamic environmental conditions structure the habitats and ecological 63 niches present and determine how species are brought together and interact with one 64 another (Guimerà et al., 2010).

65 Due to their long (multi-year) lifespans and wide spatial dispersion, assemblages of 66 epibenthic megainvertebrates (i.e., invertebrates larger than 1 cm living on the seabed) 67 and bentho-demersal fish (fish species living on or near the seabed) can provide useful 68 information on the structure of the benthic ecosystem and its response to environmental 69 drivers across large spatial and temporal scales. This approach—investigating the spatial 70 structure of ecosystems through the analysis of various benthic components—has been 71 previously applied to different European shelves (Bremner et al., 2006; Daan et al., 2005; 72 Neumann et al., 2016; Sánchez et al., 2008). While some recent studies have considered epibenthic megainvertebrates and bentho-demersal fish separately (e.g., Reiss et al.,
2010; Sánchez et al., 2008), many older studies did not (Colloca et al., 2004; Ellis et al.,
2011; Kaiser et al., 2004; Rees et al., 1999), relying instead on the hypothesis that these
two communities share common biological traits and could therefore be grouped together.

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Whether grouped together or separately, knowledge on the spatial structuration of these two benthic communities can provide a better understanding of the meso-scale (from 10 to 100 km) functioning of shelf ecosystems (Hooper et al., 2005; Shojaei et al., 2021) in the face of local and/or global environmental change. Moreover, this type of work can help to identify ecologically coherent spatial entities which can then be integrated into management efforts (e.g., Bell et al., 2021), with particular benefits for the ecosystembased management of fisheries (Garcia et al., 2003).

85 Our study focuses on the Bay of Biscay (France) as a representative case study of a North Atlantic temperate continental shelf. This zone features an extensive shelf that stretches 86 87 far offshore until finally breaking towards the abyssal plain (review in Borja et al., 2019). 88 In its coastal section, it faces numerous anthropogenic pressures (eutrophication, 89 pollution) and is the site of major fishing activity, which targets bentho-demersal species 90 in particular (Druon et al., 2021). Additionally, this area is now of growing interest for the 91 development of technologies for renewable marine energy production (Fofack-Garcia et 92 al., 2023; Taormina et al., 2018).

93 The first overview of benthic communities in the Bay of Biscay and the Celtic Sea is 94 credited to Le Danois (1948); although descriptive and not quantitative, this study provided 95 an initial picture of the overall structure of communities from the shelf to the slope.

Macrobenthic communities of the Bay of Biscay were further characterized in the north by 96 97 Glémarec (1969) and in the south by Lagardère (1973), who both described spatial 98 structuration that was mainly determined by a thermal depth gradient and sedimentary 99 substrate characteristics. A partial revisitation in the north in 2001–2002 revealed notable 100 changes in macrobenthic fauna as the result of strong fishing pressure (Hily et al., 2008). 101 At the scale of the entire bay, work by Souissi et al. (2001) provided a characterization of 102 the main assemblages of bentho-demersal fish. To date, though, there has not been a 103 wide-scale study to generate a standardized picture of the larger benthic communities of 104 the entire Bay of Biscay. In particular, there is a need for the identification of the main 105 patterns and environmental factors that structure communities, especially with respect to 106 a joint analysis of fish and invertebrate fauna.

The aim of this study was to assess the spatial structuration of the benthic ecosystem of the continental shelf of the Bay of Biscay, specifically by analyzing communities of epibenthic megainvertebrates and bentho-demersal fish and their relationships with environmental forcings. For this, we made use of the large, homogenous dataset available thanks to the recurrent international bottom trawl survey "EVHOE" (Laffargue et al., 2020). Our work was carried out in two steps:

113 1. Identify and characterize assemblages of epibenthic megainvertebrates and bentho 114 demersal fish and their spatial distribution within the study area
 115 2. Analyze the links between the spatial patterns observed and the environmental forcings

115 2. Analyze the links between the spatial patterns observed and the environmental forcings116 that structure the identified assemblages.

117 **2** MATERIALS AND METHODS

118 **2.1 Study area**

This study covers the entire French part of the continental shelf of the Bay of Biscay, extending from the island of Sein to the canyon of Capbreton (Figure 1). The present work considers both coastal and deep circalittoral areas along the margin of the continental shelf, with bathymetry ranging from 16 to 200 m in depth. This continental shelf is composed of a broad northern shelf (Armorican shelf, ~180 km wide) and a narrower southern shelf (Aquitaine shelf, ~90 km wide) (K. Koutsikopoulos & Le Cann, 1996).

125 The shelf is characterized by numerous complex seasonal hydrographic processes: 126 slope-shelf exchanges, northern slope fronts, southern coastal upwellings, and high 127 terrigenous contributions, mainly from the Loire, Vilaine, and Gironde Rivers (Akpınar et 128 al., 2020; Borja et al., 2019; Charria et al., 2013; K. Koutsikopoulos & Le Cann, 1996; 129 Labry et al., 2001; Loyer et al., 2006).



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131Figure 1. Study area: continental shelf of the Bay of Biscay. Gray dots represent132sampling points from EVHOE survey used in the analysis. Red frame divides the shelf133into the northern Armorican shelf and the southern Aquitaine shelf.

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135 **2.2 Biological data**

136 The observations used in this study were recorded as part of the EVHOE (EValuation 137 Halieutique Ouest de l'Europe) survey undertaken within the framework of IBTS surveys of European fisheries (ICES, 2017). The EVHOE survey took place each year between 138 139 October and December. Samples were collected using an otter trawl with a large vertical 140 opening (GOV, Grande Ouverture Vertical) and a 20-mm mesh size cod-end (Laffargue 141 et al., 2020). Each haul lasted 30 minutes at a speed of about 4 knots and covered about 142 0.07 km². The dataset covers the period 2008–2020 (except 2017, which was incomplete 143 due to cruise damage), with a total of 685 stations sampled.

Whenever possible, sampled organisms were sorted to the lowest possible taxonomic
level on board, with verification in the laboratory for some specimens, if necessary.
Species names were assigned as per the World Register of Marine Species (WoRMS,
2022).

148 Only epibenthic megainvertebrates and bentho-demersal fish were retained for these 149 analyses. "Megainvertebrates" refers here to organisms with a size greater than 1 150 centimeter that can live on or in the substrate and that need to move on the substrate to 151 perform a vital and/or reproductive function. This excluded from the analysis large 152 swimming cephalopods (e.g., Loligo sp., Illex sp.) and gelatinous taxa. Colonial organisms 153 (from the groups Bryozoa, Hydrozoa, and Ascidiacea) were uncountable on-board, and 154 thus were also removed from the analyses. The epibenthic megainvertebrates component 155 will be further referred to as "benthos".

As we were particularly interested in bentho-demersal fish, we excluded species whose lifecycles and activities are mainly related to pelagos. The bentho-demersal ichtyofauna component will be further referred to as "fish".

159 **2.3 Environmental variables and data**

We selected a total of seven major environmental variables that are assumed to have astructuring effect on benthic communities.

162 The coupled hydrodynamic-ecosystem 3D model NEMO-NEMOVAR-ERSEM was used 163 to obtain data for a large number of environmental variables at each haul station (EU-164 Copernicus, 2020b, 2020a). Those variables were: chlorophyll concentration, mixed layer 165 depth, nitrate concentration, dissolved oxygen concentration, pH, phytoplanktonic carbon, 166 phosphate concentration, primary production, salinity, and temperature. Monthly

averaged environmental data were gridded with a resolution of 0.111° x 0.067° (cells of 167 168 about 7 km of horizontal and vertical resolution). Considering the distribution of stations 169 and the distance sampled per haul (about 3.5 km), we judged that this resolution was 170 appropriate for the spatial scale of the study. With the exception of primary production, we 171 selected the surface and near-bottom values for each of these variables. For net primary 172 production, we considered the maximum value in the water column for each station and 173 each month. Data were integrated over the three years preceding each sampling event, 174 since the life expectancy for the majority of taxa considered here is expected to be at least 175 three years. For each available environmental variable and each year of the dataset, we 176 calculated the minimum, maximum, standard deviation, and mean values. The values 177 assigned to each station were then averaged over the three years prior to the observation 178 date. In this way, we obtained an initial set of 72 environmental variables and derived 179 calculations for each sampling station.

180 To complete the set of environmental variables, additional data were selected regarding 181 bottom current and substrate. Bottom stress, which represents the velocity of the water 182 movements experienced by benthic organisms due to currents and waves, has been 183 found to be an important variable structuring the diversity of benthic communities (Reiss 184 et al., 2010). Here, this was quantified as the 90th percentile of the kinetic energy at 1 m 185 from the seabed, with a horizontal resolution of 1/36° (about 3 km). Since the NEMO-186 NEMOVAR-ERSEM model does not include wave components, we obtained data on the 187 kinetic energy due to currents from the MANGA500 model, available on the EMODnet 188 platform (Accensi & Maisondieu, 2015), and data on the kinetic energy due to waves from 189 the WAVEMATCH III model (Caillaud et al., 2016). Data were averaged between 2010 190 and 2015. We assumed that there were no strong temporal variations in these two variables (energy due to currents and energy due to waves). These two types of energy
were summed into a single quantitative "bottom energy" variable.

We performed a preliminary analysis to avoid over-correlation among variables. From the initial set of variables, we selected six quantitative descriptors corresponding to key ecological processes:

- Standard deviation of surface salinity ('sd_Sal'), assumed to reflect the
 proximity to river plumes with variable seasonal inputs.
- Standard deviation of bottom seafloor temperature ('sd_Temp'), reflecting the
 climatic gradation of the benthic domain (infralittoral, coastal circalittoral, and deep
 circalittoral) (Chassé & Glémarec, 1976).
- Bottom dissolved oxygen minimum ('min_Oxy'), reflecting the input at the
 air/water interface and the use by living organisms (respiration, bacterial organic
 mineralization).
- Kinetic energy at the seabed ('q90_Energy'). This variable has a direct influence,
 through the physical pressure exerted, on the ability of a species to colonize or live
 in a given habitat, as well as an indirect influence on the trophic capacity of an area
 through organic matter resuspension and export for example.
- Average depth of the mixed layer ('avg_MLD'). This variable was assumed to
 increase close to frontal zones, where strong mixing occurs that is thought to
 support primary production.
- Maximum net primary production in the water column ('max_NetPP'),
 averaged over the year. This highlighted areas with high net primary production
 rates, notably below the water surface in offshore areas.

214 Calculation of these six quantitative descriptors is summarized in Appendix Table A.1.

To this subset of environmental descriptors, we added a qualitative variable representing substrate type, following the categories defined by the French Marine Hydrographic and Oceanographic Service (SHOM, 2021): mud, silt, fine sand, sand, gravel, and rock. Substrate categories were viewed as indicative of overall environmental conditions, with low current and high deposition rates in muddy areas, and strong hydrological conditions in coarser areas.

This set of seven descriptors (six quantitative and one qualitative) was found to strongly
reflect the variability in the original data (Gower's dissimilarity, Spearman's correlation:
85.6%, Mantel permutation test significance: p-value <0.001).

224 2.4 Biological data analysis

225 Each analysis was conducted separately on benthos and fish. Analyses were 226 performed using the vegan, stats, and cluster packages in R 4.3.1 software (Oksanen et 227 al., 2022; R core team, 2023). Only species with an occurrence equal to or greater than 228 5% were retained, and abundance matrices were double square-root transformed to avoid 229 the distorting effect of heavily abundant species. Bray-Curtis dissimilarity matrices were 230 generated for each biological component (fish or benthos), and the dissimilarity matrices 231 were used to conduct ascendant hierarchical clustering using Ward linkage to aggregate 232 hauls into several clusters, the number of which was determined using visual and inertia 233 criteria (Borcard et al., 2018b). Clusters that were faunistically distinct from others were 234 denoted "assemblages". The significance of the clustering pattern was assessed through 235 an analysis of similarity using the "ANOSIM" function of the vegan package.

Spatial overlap between benthos and fish assemblages was first assessed visually and
then quantified using a Spearman's correlation between the two dissimilarity matrices.
This correlation was tested with a Mantel permutation test.

239 Each cluster was taxonomically characterized using the IndVal method from the labdsv package. Indices reflecting structural parameters—biomass $(kg.km^{-2})$, abundance 240 (thousands of individuals per km²) (10^3 ind.km⁻²), and Shannon diversity—were used to 241 242 identify spatial patterns in both benthos and fish clusters. Calculations of observed 243 taxonomic richness (per haul) and Pielou's evenness are available in the Appendix 244 (Tables A.2 and A.3). Pairwise Wilcoxon tests were performed to assess differences 245 between clusters with respect to the structural indices. We also compared the overall 246 distribution of the structural parameters of each assemblage per station by performing 247 Pearson correlations between invertebrates and fish for each parameter (biomass, 248 abundance, and diversity). In order to balance the weight of the different assemblages 249 depending on the number of stations represented, we calculated median correlations 250 based on 1000 random selections of 26 stations per cluster. For this analysis, we only 251 kept the stations (444 stations out of a total of 685) with a similar cluster attribution for the 252 two biological components.

We performed distance-based redundancy analyses (db-RDAs) on Bray-Curtis dissimilarity matrices to explicitly explore the relationships between species distributions and environmental variables (Borcard et al., 2018a). Variance partitioning was performed to assess the importance of each environmental variable in determining the structure of benthic assemblages. By combining the results of the variance partitioning with canonical ordination, we were able to identify relationships among the identified assemblages as well as relationships between the assemblages and environmental variables.

260 **3 Results**

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261 **3.1 Spatial organization of the clusters**

In total, 141 fish and 342 benthos taxa were recorded during the sampling period. Of these, 47 fish and 77 benthos taxa were considered to be non-rare species (occurrence > 5%) and were retained for the hierarchical clustering. This procedure identified seven clusters within each of the two communities (corresponding dendrograms are in Appendix Figures A.3 and A.4). For both fish and benthos, clustering patterns were spatially coherent (Figure 2) and significant (ANOSIM, R = 0.555 (p-value < 0.001) and 0.542 (p-value < 0.001) for benthos and fish, respectively).



Figure 2. Clustering of megainvertebrates and bentho-demersal fish. Each color
 represents one cluster. Triangular points represent invertebrate samples; square points
 represent fish.

273 The north coast cluster was mostly located in shallow depths (median depth of 32 m for 274 benthos and 26 m for fish) and close to estuarine areas (Loire, Vilaine, and Gironde) on 275 the Armorican shelf and the northern part of the Aquitaine shelf (Figure 2). The south 276 coast cluster was found mainly in shallow waters in the southern portion of the Aguitaine 277 shelf at a median depth of 32 and 35 m for benthos and fish, respectively. Moving offshore, 278 the shallow north coast cluster was followed by the inner shelf cluster (median depth of 279 89 and 64 m for benthos and fish, respectively) and then by a cluster associated with the 280 Grande Vasière (median depth of 102 and 100 m, respectively, for benthos and fish), a 281 well-characterized muddy area that hosted distinct assemblages of both fish and benthos. 282 Further offshore, the southern outer shelf, located on the Aquitaine shelf, was also 283 characterized by unique assemblages of both benthos and fish (median depth of 146 and 284 127 m, respectively), as were the northern middle shelf (respective median depths of 128 285 and 123 m) and the northern outer shelf (respective median depths of 146 and 143 m). 286 The southern portion of the Aquitaine shelf appeared to be predominantly structured into 287 two clusters: the south coast and the southern outer shelf.

It was evident from this analysis that there are strong spatial convergences between the benthos and fish communities. Indeed, in a comparison of the fish and benthos components (Bray-Curtis dissimilarity matrices), we detected a significant overall Spearman correlation of 68% (Mantel statistic R: 0.6763, p < 0.001, 10000 permutations).</p>

292 **3.2 Description of the assemblages**

The main groups of benthos were Malacostraca (93 taxa), Gastropoda (45 taxa), Bivalvia (37 taxa), Polychaeta (30 taxa), Anthozoa (26 taxa), Asteroidea (19 taxa), Echinoidea (17 taxa), Ophiuroidea (16 taxa), Cephalopoda (10 taxa), Holothuroidea (10 taxa), and Ascidiacea (9 taxa). 297 We used an indicator species analysis (IndVal) to identify the most representative species 298 within each cluster (Figure 3). The north coast cluster was represented by coastal species 299 such as the hermit crab Pagurus bernhardus, the crab Liocarcinus holsatus, and the 300 polychaete Aphrodita aculeata. Although the south coast cluster shared many species 301 with the north coast, but was less faunistically typed with the sea anemone Calliactis 302 parasitica and the crab Polybius henslowii as the most characteristic species. Species in 303 the inner shelf cluster were less indicative than those of the other clusters; here, the 304 highest indicator values were assigned to the pennatulid *Pteroeides griseum*, the sea 305 anemone Adamsia palliata associated with the hermit crab Pagurus prideaux, but these 306 species were all also observed in several other clusters. The Grande Vasière cluster 307 appeared to be highly characterized by the presence of Norway lobster 308 (*Nephrops norvegicus*), the shrimp Crangon allmanni, and the squat lobster 309 Munida rugosa. Instead, the northern middle shelf was mostly defined by the deep-water 310 crinoid Leptometra celtica, and the northern outer shelf by the starfish Porania (Porania) 311 *pulvillus* and the polychaete *Hyalinoecia tubicola* (Figure 3). Finally, the southern outer 312 shelf cluster was characterized by the brittlestar Ophiura ophiura and the anemone 313 Actinauge sp., even those these taxa were also shared with other clusters.

With respect to fish, the taxa represented were mainly Gadiformes (22 taxa), Perciformes (20 taxa), Pleuronectiformes (18 taxa), Sparidae (12 taxa), Gobiiformes (11 taxa), Rajiformes (8 taxa), Labridae (7 taxa), and Syngnathiformes (7 taxa).

Several species were identified in multiple clusters: the dogfish *Scyliorhinus canicula*, the European hake (*Merluccius merluccius*), the red gurnard (*Chelidonichtys cuculus*), and the thickback sole (*Microchirus variegatus*). The coastal, Grande Vasière, and outer shelf clusters appeared to be more faunistically defined than the northern middle shelf or inner shelf clusters, with more-characteristic species: the bogue (*Boops boops*), greater weever fish (*Trachinus draco*), and the striped red mullet (*Mullus surmuletus*) for the southern coast; the whiting (*Merlangius merlangus*) and the common sole (*Solea solea*) for the northern coast; the goby *Lesueurigobius friesii* for the Grande Vasière; the lesser silver smelt (*Argentina sphyraena*) and the ray *Leucoraja naevus* for the northern outer shelf; and the megrim *Lepidorhombus boscii* and two gurnards, *Eutrigla gurnardus* and *Trigla lyra*, for the southern outer shelf.



Figure 3. The five most indicative taxa of each of the seven clusters, as determined from indicator values (IndVal analysis). The warmer the color of the cell, the higher the indicator value (blue = low; red = high).

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For both benthos and fish, the north coast clusters had on average the highest biomass
and abundance (Figure 4, summarized values given in Appendix Tables A.2 and A.3) over
the study period (2008–2020). Within this cluster, benthos had an average biomass of

336 249.75 kg/km^2 and an average abundance of 20.29 10^3 ind.km^{-2} , and fish had an 337 average of 1302.27 kg/km^2 of biomass and an abundance of 20.27 10^3 ind.km^{-2} .

338 When considering the distribution of biomass from the coast to offshore areas, we 339 detected an increase in benthos biomass from the inner shelf to the northern outer shelf cluster $(47.06 \text{ kg/km}^2 \text{ to } 132.56 \text{ kg/km}^2, \text{ respectively})$ (Figure 4). A similar pattern was 340 observed for abundance (ranges from 4.18 10^3 ind.km⁻² to 17.41 10^3 ind.km⁻²), with a 341 342 peak on the northern middle shelf (20.68 10^3 ind.km⁻², Figure 4). The northern coastal, 343 middle, and outer shelf clusters of benthos had significantly higher biomass than the 344 others (pairwise Wilcoxon test). High values for biomass in the northern middle shelf 345 cluster were mainly due to the presence of Leptometra celtica and the octopus Eledone 346 *cirrhosa*, while biomass patterns in the northern outer shelf appeared to be strongly 347 influenced by the crab Cancer pagurus and the holothurian Parastichopus regalis. Regarding diversity, values of Shannon's index were constant on the shelf and 348 349 significantly higher close to the shelf edge (northern outer shelf). Benthos diversity was 350 significantly higher in the northern clusters than in the southern clusters (south coast and 351 southern outer shelf).

352 For fish, the same global patterns in biomass and abundance were observed, but with 353 peak abundance occurring in the Grande Vasière cluster. On the inner shelf, average fish biomass and abundance were 693.75 kg/km^2 and $11.17 \ 10^3 \text{ ind.km}^{-2}$, respectively. 354 355 These values increased in the direction of the northern outer shelf cluster, with the highest values found for the Grande Vasière cluster (962.49 kg/km² and 22.86 10^3 ind.km⁻², 356 357 respectively). This cluster, along with the north coast cluster, had significantly higher fish 358 abundance than the others. The main species responsible for the high abundances and 359 biomass observed in these areas were the pouting (Trisopterus luscus) for the north coast and the hake (*Merluccius merluccius*) for the Grande Vasière. Values of Shannon diversity
and Pielou's evenness were more stable across the shelf for fish than for benthos (Figure
4, and Appendix Tables A.2 and A.3), and larger-scale diversity patterns from coast-tooffshore were less clear. However, we did observe that the Grande Vasière was home to
a significantly less diverse assemblage of fish than the outer shelf clusters.

For both benthos and fish, biomass and abundance were always significantly higher in the Grande Vasière cluster than in the inner shelf cluster. However, the benthos in the northern outer shelf cluster had significantly higher biomass and abundance than those in the Grande Vasière cluster (Figure 4). Among all the clusters, patterns of biomass and abundance were 56% and 60% correlated, respectively, between benthos and fish communities.

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Figure 4. Distribution of structural indices for each spatial cluster of benthos (left) and
fish (right). Biomass (in kg.km⁻²) and abundance (in thousands of individuals.km⁻²) are
log-transformed. Within each panel, boxplots with the same letters were not significantly
different following a pairwise Wilcoxon test. For visual convenience, outliers are
removed from the figure.

378 **3.3** Role played by environmental forcings



379 3.3.1 Influence of environmental forcings on benthos

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381 Figure 5. Distance-based redundancy analysis for benthos abundance at EVHOE 382 sampled stations. Ellipses represent the 95% confidence interval. Together with 383 substrate type, six environmental drivers were considered: annual average mixed layer depth (avg MLD), annual minimum dissolved oxygen at the seafloor (min Oxyg), annual 384 standard deviation of sea surface salinity (sd Sal), annual standard deviation of the 385 temperature at the seafloor (sd_Temp), annual maximum net primary production in the 386 water column (max NetPP), and 90th percentile of average bottom kinetic energy 387 388 (q90 Energy).

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With respect to the benthos abundance data, the distance-based redundancy analysis (db-RDA) explained 19.47% of the total variance. The first two axes of the analysis explained 73.99% of the fitted variation and 14.41% of the total variation in benthos clusters (Figure 5). The results highlighted a coast-to-offshore gradient, with a higher 394 range in seafloor temperature and net primary production in coastal areas. The southern 395 coast cluster had higher bottom energy than the north coast cluster, where bottom energy 396 was more variable. The Grande Vasière cluster, which featured the muddiest conditions 397 in the entire area, stood out from the others along axis 2. The inner shelf and southern 398 outer shelf were associated with the widest ellipses, which was likely a function of their 399 lower faunistic specificity (see above) and greater environmental variability. The inner 400 shelf cluster seemed to represent a transitional zone between the coastal and offshore 401 clusters, and did not appear to feature any clearly discriminating environmental 402 characteristics of its own. Offshore areas of the northern shelf were mainly characterized 403 by fine sand with a deep mixed layer, and low variation in bottom temperature. Finally, the 404 substrate of the northern middle shelf cluster was coarser than that of the northern outer 405 shelf.

406 Variance partitioning revealed that the seven environmental descriptors explained 18% of 407 the total variability (Appendix Figure A.5). The most influential factors were, in order: 408 mixed layer depth (average), substrate type, seafloor temperature (standard deviation), 409 maximum net primary production (average), and bottom energy (90th percentile). The 410 mixed layer was deeper, on average, for the northern middle and outer shelves than 411 southern ones, and for inner shelf clusters compared to coastal ones. Coastal clusters 412 were also distinguished from the others based on differences in seafloor bottom 413 temperature (standard deviation), maximum net primary production (average), and bottom 414 energy. Surface salinity (standard deviation) was higher only for the northern coast cluster, 415 reflecting the terrigenous influence in that location.



3.3.2 Influences of environmental forcings on fish 416



418 Figure 6. Distance-based redundancy analysis for fish abundance at EVHOE sampled stations. Ellipses represent the 95% confidence intervals. In addition to substrate type. 419 six environmental drivers were considered: annual average mixed layer depth 420 421 (avg_MLD), annual minimum dissolved oxygen at the seafloor (min_Oxyg), annual standard deviation of sea surface salinity (sd Sal), annual standard deviation of the 422 temperature at the seafloor (sd_Temp), annual maximum net primary production in the 423 424 water column (max NetPP), and 90th percentile of average bottom kinetic energy 425 (q90 Energy).

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427 For the fish abundance dataset, the redundancy analysis constrained 26.55% of the total 428 variance. The first two axes explained 78% of the fitted variation and 20.7% of the total 429 variance (Figure 6). Overall, redundancy analysis was able to explain more of the variation in fish distributions (26.55%) than it was for benthos (19.47%). As for benthos, the first 430 431 axis corresponded to a bathymetric gradient. The north coast cluster was associated with 432 an increasing range (standard deviation) of salinity, while the south coast cluster was 433 linked with stronger bottom energy. Instead, both coastal clusters were associated with 434 stronger net primary production. However, the overall separation of the coastal clusters 435 from the other assemblages was less clear for benthos than for fish. As observed for 436 benthos, the inner shelf cluster seemed to represent a transitional state between the 437 coastal and offshore clusters. Likewise, the northern middle shelf cluster (characterized 438 by a coarse substrate) was located in an intermediate position between the Grande 439 Vasière, the northern outer shelf, and the southern outer shelf (Figures 5 and 6). Variance 440 partitioning revealed that the seven variables explained 25% of the total variability 441 (Appendix Figure A.6). The forcings influencing fish clusters were mixed layer depth 442 (average), seafloor temperature (standard deviation), and maximum net primary 443 production (average). Compared to benthos communities, fish were less influenced by 444 substrate but more influenced by surface salinity variation (standard deviation).

To sum up, the coastal clusters corresponded to areas with a higher range of seafloor temperatures and higher primary production. The south coast cluster was located in a sandier and more energetic area than the north coast cluster. The cluster associated with the Grande Vasière was in a muddy area with low bottom energy. The inner shelf represented a transitional zone between the coastal and offshore areas, while the northern middle shelf bridged the gap between the Grande Vasière and the northern outer shelf.

452 With respect to substrate, that of the northern middle shelf was coarser than that of the 453 Grande Vasière and the northern outer shelf. The north and south outer shelf clusters 454 were distinguished from each other by substrate and mixed layer depth, which were finer 455 and deeper, respectively, for the northern shelf.

456 **4 Discussion**

457 4.1 Definition of assemblages as a tool for understanding the structure and 458 functioning of continental shelf ecosystems

459 There is a pressing need for improved characterizations of the spatial structure and 460 functioning of marine environments (Pittman et al., 2021), especially in the context of 461 marine ecosystem quality assessment and management (Lavialle et al., 2023). The 462 present work—a meso-scale (10–100 km) study of the spatial distribution of epibenthic 463 megainvertebrates and bentho-demersal fish on a continental shelf-represents an 464 important step forward in this regard. In particular, it provides complementary perspectives 465 to the previously published EUNIS typology of the Bay of Biscay, which was mainly based on macrofauna features (Bajjouk et al., 2015). 466

467 Our study identified seven clusters of epibenthic megainvertebrates and bentho-demersal 468 fish that were biologically coherent and spatially well-delineated and, thus, represented 469 distinct assemblages distributed within the larger study site. A limited number of 470 environmental forcings (e.g., hydrological features, estuarine influence, sediment type) 471 were identified as the main factors responsible for structuring the identified patterns within 472 each biological community (benthos or fish). Despite the biological differences between 473 bentho-demersal fish and epibenthic invertebrates, we found a high degree of similarity in 474 the spatial distribution of the different assemblages within the two communities 475 (correlation 68%), which exhibited similar coast-to-offshore patterns.

Any attempt to understand the ecosystem functioning of continental shelves based on the definition of biological assemblages is likely to be influenced by the choice of methodology. The spatial structures resulting from a given analysis can be highly dependent on the sampling strategy and the nature of the data recorded (e.g., sampling gear, spatial

resolution, sampling period). In this case, the equipment used was more appropriate for 480 481 sampling bentho-demersal fish than benthic invertebrates. Because the sampling gear 482 poorly penetrates the soft bottom, only larger epibenthic organisms can be properly 483 sampled. In addition, although our study relies on data from several years (2008–2020), 484 sampling was conducted in only a single month of the year (October). However, the 485 absence of any clear seasonal fluctuation in epifaunal megabenthic communities in 486 comparable environments (e.g., Hinz et al., 2004) gave us confidence that the overall 487 structure described in our study was very likely stable over an annual scale. Moreover, 488 we checked that each of the identified clusters remained distinct from the others over the 489 12 years of our study, and detected no strong inter-annual variations in the observed 490 spatial distributions. This hypothesis was also supported by a recent report that spatial 491 patterns were stable and strongly dominant compared to temporal variations in shaping 492 the beta-diversity of bentho-demersal fish in the Bay of Biscay and the Celtic Sea (Eme 493 et al., 2022).

494

495 4.2. Spatial distribution of assemblages and environmental drivers on the 496 continental shelf of the Bay of Biscay

The spatial structures identified here followed a pattern of coast-to-offshore succession, and were marked by a clear difference between the Armorican shelf in the north and the Aquitaine shelf in the south.

500 The northern coast assemblages are situated in a shallow circalittoral area (around 30 m 501 deep) and are influenced by river (Loire and Gironde) plumes, as evident from their higher 502 variability in surface salinity in the canonical ordination. Communities of both fish and 503 invertebrates featured species that are characteristic of coastal muddy areas, such as the 504 crab Liocarcinus holsatus and the flatfish Solea solea (C. Koutsikopoulos et al., 1989). 505 With respect to coastal fish, the main indicator species observed in our study (e.g., 506 Pomatoschistus minutus and Merlangius merlangus) were the same as those described 507 by Souissi et al. (2001). River plumes are a major structuring factor of the Bay of Biscay 508 as they are significant drivers of primary production (Lunven et al., 2005), leading to high 509 species abundance, biomass, and secondary production in coastal areas (Guillaud et al., 510 2008; Saulnier et al., 2019). Along the northern coast of the Bay of Biscay, this terrestrial 511 influence is also retrieved in coastal macrofaunal assemblages (Dutertre et al., 2013). 512 Another explanation for the greater taxonomic diversity (Stein et al., 2014) observed in 513 this area could be the high number and variety of small-scale habitats utilized by 514 epibenthic communities (Vasquez et al., 2015).

515 Southern coastal assemblages are also located in shallow circalittoral areas (around 30 516 m deep). They do not benefit from the influence of plumes but are subjected to high-517 velocity bottom currents, which is reflected in the mud-free nature of the substrate. Thus, 518 both currents and substrate type were found to have a significant structuring effect on the 519 fish and benthos assemblages in this region; as an example, the fish community was 520 characterized by the presence of species such as the lesser weever fish (Echiichthys 521 vipera), which is known to burrow in clear-sandy substrate (Wheeler, 1969). Similar 522 bentho-demersal fish species were identified by Souissi et al. (2001) within this area in 523 October 1990. Such "sandy habitat" was observed by Lagardère (1973) and Monbet (1972) 524 for macrobenthos communities along the Aquitaine shelf, which were also structured 525 along a bathymetric gradient. Compared to the northern coast, the abundance and 526 biomass of both fish and benthos were 10 times lower in the south, possibly the result of

a lower nutrient supply due to a weaker river influence. To our knowledge, our study
represents the first large-scale study on epibenthic megainvertebrates (with the exception
of decapods) along the coast of the Bay of Biscay, and sheds new light on the benthic
communities of this area.

531 Further offshore, in the central part of the continental shelf, the inner shelf assemblages 532 appeared to represent a transitional zone between shallow and deeper areas. This area 533 was characterized by a high degree of environmental variability, which likely promoted the 534 development of assemblages composed of habitat generalists rather than specialists.

535 The north central area of the shelf—the Grande Vasière—hosted a unique assemblage of 536 benthos, which appeared to be associated with the characteristic muddy substrate of this 537 zone. Instead, the fish community of this area was less distinctive, probably reflecting the 538 fact that these assemblages are less dependent on substrate type (Appendix Figures A.5 539 and A.6). Among the species observed here were typical burrowers such as the 540 crustacean decapod N. norvegicus or the goby L. friesii (Ellis et al., 2013). The high 541 amounts of benthos and fish biomass we found were expected, as this area is of great 542 importance to the fishing industry; it benefits from high spring bloom production (Guillaud 543 et al., 2008) and high quantities of detrital matter (Le Loc'h et al., 2008).

Offshore on the Armorican shelf, the northern middle shelf hosted assemblages of fish and benthos that were less well defined faunistically and environmentally than their neighbors, and can therefore be viewed as transitional between the Grande Vasière and the deeper areas near the shelf edge. Glémarec (1969) described this area as a barrier with coarser substrate that limits the offshore extent of the Grande Vasière. As in the present study, he identified sandy communities with high densities of *Leptometra celtica* as observed also in our study, marking a transition between the "Grande Vasière" mudsand bathyal muds further offshore.

The northern outer shelf assemblages are located in the deep circalittoral area (around 552 553 145 m deep) and are characterized by a deep mixed layer. This layer is caused by the 554 presence of a frontal zone (K. Koutsikopoulos & Le Cann, 1996) where the interaction of 555 barotropic tide with the slope results in surface-water mixing (Simpson & Sharples, 2012). 556 This promotes the upwelling of cold, nutrient-rich waters from great depths and thus feeds 557 strong primary production (Sharples et al., 2009). The influence of this frontal structure 558 was previously observed in a study of zooplanktonic secondary production on the northern 559 shelf of the Bay of Biscay (Vandromme et al., 2014). In the context of our study, this 560 phenomenon would be more likely to affect the structural parameters (biomass, 561 abundance, and diversity) of invertebrate communities, and to a lesser extent those of the 562 bentho-demersal ichthyofauna. Indeed, here we found that, compared to the assemblages 563 further from edge, the outer shelf boasted significantly higher invertebrate biomass, 564 confirming previous observations by Le Loc'h (2004). This suggests that the unique 565 hydrodynamic processes in this area are a major driver structuring the shelf-edge 566 ecosystem from plankton to benthos.

567 Despite their proximity to the shelf break, the southern outer shelf assemblages differ from 568 the northern ones and do not benefit from the upwelling observed on the Armorican shelf 569 break (Koutsikopoulos and Le Cann, 1996). They are located in a deep circalittoral area 570 (around 135 m deep) and are less faunistically distinct than communities on the northern 571 outer shelf. Prior to our study, the only types of epibenthic megainvertebrates that had 572 been described in this area were the decapod crustaceans (Lagardère, 1973). Here, we 573 observed that the widespread brittle-star *Ophiura ophiura*, known to have no sediment 574 preference (Boos et al., 2010), was the most indicative species. For fish, the species with 575 the highest indicator value was the flatfish *Lepidorhombus boscii*, which was also 576 proposed to be indicative (as a 'secondary species') in this area 20 years earlier (Souissi 577 et al., 2001). No major determining environmental drivers were identified for either the fish 578 or benthos communities.

4.3 Spatial structuration of meso-scale benthic ecosystems across European

579

580 continental shelves

The main benthic features identified in our study can be found in other areas of the North Atlantic European continental shelf system, and reflect certain common characteristics of benthic ecosystem functioning. Although assemblages may differ in terms of faunal composition, similar key species are found in comparable habitats in all European areas. The convergence of spatial structures between the various ecosystem components mega- and macroinvertebrates and bentho-demersal fish—has also been widely reported, and seems to originate from the effects of common driving factors.

588 Shallow coastal assemblages similar to the north coast assemblages of our study are 589 observed in the Celtic Sea, the English Channel, and the North Sea, which all feature 590 indicator species similar to those found in the shallower area of our study site, such as the 591 crustaceans Pagurus bernhardus and Liocarcinus holsatus (Ellis et al., 2013; Hinz et al., 592 2004; Reiss et al., 2010) and the fish Solea solea, Trisopterus luscus, and Merlangius 593 merlangus (Vaz et al., 2007) or Buglossidium lutem (Reiss et al., 2010). The presence 594 and abundance of these species reflect terrigenous influences, specifically related to the 595 proximity of estuaries (Ruiz-Castillo et al., 2019; Vaz et al., 2007).

Along the Welsh coastline (Kaiser et al., 2004) or in the south of the North Sea (Ellis et al., 2011), surveys have identified fish species with affinity for sandy substrate, such as *Echiichthys vipera* (Reiss et al., 2010) which here was detected as an indicator species of the south coast assemblage in the Bay of Biscay. As in the present study, assemblages with this species on the Welsh coastline were also located in areas with high tidal amplitude (Horrillo-Caraballo et al., 2021) and therefore high bottom energy.

602 Further offshore, previous studies of other European shelves have identified assemblages 603 that appear to be similar to the transitional ones we identified between shallow and deeper 604 areas of the Bay of Biscay. For example, Ellis et al. (2013) described a wide mid-shelf 605 invertebrate assemblage in the Celtic Sea featuring the polychaete Hyalinoecia tubicola 606 and the crab *Liocarcinus holsatus*. In the same area, Mérillet et al. (2019) reported a broad 607 mid-shelf fish assemblage with widespread species like Trisopterus minutus. These widespread assemblages do not seem to be shaped by any clearly identified physical 608 609 processes and, given their faunistic composition, can be thought of instead as transitional 610 groups.

Assemblages typical of muddy areas corresponding to the Grande Vasière of our study site also appear to be common along European shelves, for example in the North Sea close to the English coast (Rees et al., 1999) and in the Celtic Sea offshore of the Bristol Channel (Ellis et al., 2013; Rees et al., 1999). These mud-associated fauna seem to be located in areas where strong primary production occurs in spring blooms (González-Gil et al., 2022; Seguro et al., 2019).

Likewise, the outer shelf assemblages found in our study also have their equivalents on other European shelves. Species that were found here to be representative of northern outer shelf assemblages in the Bay of Biscay (e.g., *Lepidorhombus whiffiagonis*, *Ophiotrix* *luetkeni*, or *Argentina sp.*) are commonly found along shelf edges in other European areas like the Celtic Sea (Ellis et al., 2013; Mérillet et al., 2019). These areas are all under the influence of shelf-break upwellings which are known to enhance local biological production (Kossack et al., 2023). This influence can also extend to the more internal assemblages of the continental shelf, for example through high densities of crinoids such as *Leptometra sp.*, as found in deep areas of the Mediterranean shelf (Colloca et al., 2004).

4.4 From spatial assemblages to "benthiscapes": towards ecological modeling and ecosystem-based management

628 The correspondence of spatial distributions among different components of benthic 629 communities is a common feature of continental shelves. For example, in the Celtic Sea similar spatial patterns were noted for assemblages of epifaunal invertebrates and fish 630 631 (Ellis et al., 2013; Mérillet et al., 2019). Farther to the north, the benthic communities of 632 the North Sea shelf were found to be structured into mesoscale assemblages comparable. 633 to some extent, to those in the Bay of Biscay, with a strong correlation between 634 communities of epifaunal invertebrates and demersal fish as well as infauna (Reiss et al., 635 2010). Beyond the European shelves of the North Atlantic, close similarities in mesoscale 636 community structure have also been reported for demersal fish and epibenthic 637 invertebrates in the Mediterranean Sea, in the Gulf of Lions (Gaertner et al., 1999) and 638 along the Italian shelf break (Colloca et al., 2004). Even in regions with stronger 639 environmental gradients, such as the Barents Sea, communities of benthic invertebrates 640 and fish were strongly associated with each other, and co-varied with depth and the 641 ice/temperature gradient (Johannesen et al., 2017). This high degree of spatial similarity 642 between epifaunal invertebrates and fish does not necessarily imply direct interactions 643 (e.g., trophic linkages) between these ecological components. Instead, it may be the result of the strong convergence of intense environmental drivers at the scale of continental
shelves that shape and structure different biological communities in similar ways.

The strong convergence of the spatial distributions of megainvertebrates and bentho-646 647 demersal fish in the Bay of Biscay provides evidence for the existence of different benthic 648 habitats characterized by different substrates and hydrological processes, as well as 649 shelf-scale environmental gradients (coast-to-offshore and south-to-north). We propose 650 defining these benthic habitats as "benthiscapes". In our study, these habitats appear to 651 be consistent with the "hydrological landscapes" described by Planque et al. (2004), which 652 appear to structure both the benthic and the pelagic systems. In the pelagic domain, the 653 main factors shaping the spatial structure of biological communities are the influences of 654 freshwater, water column stratification, and bottom temperature (Petitgas et al., 2018), 655 similar to many of the processes highlighted in the present work. These commonalities 656 strengthen the "meta-ecosystem" vision of the continental shelf proposed by Petitgas et 657 al. (2018). The benthic system is also known to be strongly connected to water column 658 processes (Piepenburg et al., 1997), and the biological components of benthic 659 ecosystems are fundamental to the transfer of matter from the water column to sediments 660 (Zhang et al. 2021). These components are in turn fundamentally shaped by the physical 661 structure of continental shelves, particularly in terms of nutrient sources. Previous work 662 by Ruzicka et al. (2018) defined a few of the major physical processes structuring various shelf ecosystems from the inner to the mid- and outer shelf. Similarly, the benthiscapes 663 664 identified in our study reflect spatially determined alterations in ecosystem functioning that 665 go beyond simple taxonomy-based structures. Here, these structural patterns have been 666 described very briefly in terms of differences in ecological parameters such as biomass 667 gradients. The spatial variability we observed raises the question of differences in trophic functioning between the different benthiscapes, which is all the more important given the correlations we observed (mostly with respect to abundance and biomass) between communities of invertebrates and fish. Such patterns suggest that ecological functioning varies spatially in the same way for both benthos and fish; further investigation of this issue will improve our understanding of organic matter fluxes at the scale of the continental shelf.

674 One application of the concept of benthiscapes—benthic habitats characterized by unique 675 ecological assemblages, distinct functions, and potentially differing responses to natural 676 and anthropogenic pressures—could be in the development and improvement of 677 strategies for ecosystem-based management. The definition of spatially coherent benthic 678 entities, based on this study and others, could assist in efforts to monitor seafloor 679 biodiversity and understand and manage anthropogenic perturbation at the scale of 680 continental shelves (e.g., Lauria et al., 2020). In this context, the benthiscapes described 681 in our study could therefore be considered as one of the fundamental structures to develop 682 tools for spatial planning. However, our knowledge of seafloor ecosystem functioning in 683 the Bay of Biscay remains incomplete (Galparsoro et al., 2014), and further studies are 684 needed to fill the gaps on the different biological communities of this area. The need is 685 especially pressing given the significance of the natural resources of the Bay of Biscay 686 (e.g., sand extraction, offshore wind farms, fishing for benthic and demersal species), 687 which are already heavily utilized and likely to be increasingly important in the future 688 (Boussarie et al., 2023; Lavialle et al., 2023).

689 **4 CONCLUSION**

690 This study investigated the overall spatial structure of ecological communities on the 691 continental shelf of the Bay of Biscay through the identification and characterization of 692 assemblages of epibenthic megainvertebrates and bentho-demersal fish. These 693 assemblages were faunistically similar to communities described on other European 694 continental shelves. The spatial delineation of biological communities into distinct 695 assemblages was highly similar between fish and invertebrates, suggesting that the 696 continental shelf ecosystem is made up of different spatial habitats that are predominantly 697 structured by environmental drivers such as bottom temperature, sediment type, bottom 698 energy, and primary production. Here, we propose the term "benthiscapes" for such 699 habitats.

The benthiscapes characterized in this study differed from each other with respect to multiple structural parameters, particularly biomass, which was highest at the coast and close to the continental slope. This suggests that the trophic functioning of this ecosystem is dependent not only on terrestrial inputs from major rivers but also on the influence of offshore upwelling.

The concept of benthiscapes could be particularly useful for applications in the areas of ecosystem-based management and marine spatial planning. Further characterization of these spatially distinct meso-scale entities could lead to an improved understanding of continental shelf ecosystems and the essential ecosystem functions they provide.

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1028 **5. APPENDIX**

1029

1030 Table A.1: Summary of environmental variables analyzed

	Method of Variable name calculation		Time period averaged	Position in the water	the References		
Bottom energy		90th percentile	2010-2015	1 m from the seabed	(Accensi & Maisondieu, 2015; Caillaud et al., 2016)		
	Mixed layer depth	Mean	year-2; year- 1; year of sample	Surface of the water column	(EU-Copernicus, 2020b)		
	Dissolved oxygen	Minimum	year-2; year- 1; year of sample	Near-seafloor value	(EU-Copernicus, 2020a)		
	Net primary production	Mean	year-2; year- 1; year of sample	Depth where monthly maximum value was reached	(EU-Copernicus, 2020a)		
	Salinity	Standard deviation	year-2; year- 1; year of sample	Surface of water column	(EU-Copernicus, 2020b)		
	Seafloor temperature	Standard deviation	year-2; year- 1; year of sample	Seafloor	(EU-Copernicus, 2020b)		
103	1						
103	2						
103	3						
103	4						
103	5						
103	6						
103	7						
103	8						
103	9						
104	·0						

1041 Table A.2: Summary of structural indices for epibenthic megainvertebrate communities (average (standard-deviation))

Index	South coast	North coast	Inner shelf	Grande Vasière	Northern middle shelf	Northern outer shelf	Southern outer shelf
Shannon's diversity	1.86 (0.8)	2.69 (0.6)	2.66 (0.65)	2.64 (0.54)	2.8 (0.91)	3.08 (0.67)	1.76 (0.85)
Pielou's evenness	0.72 (0.18)	0.64 (0.14)	0.7 (0.17)	0.62 (0.13)	0.61 (0.2)	0.65 (0.14)	0.46 (0.23)
Taxonomic richness	7.5 (3.71)	20.26 (8.21)	15.64 (6.47)	20.09 (4.86)	24.83 (6.5)	27.8 (5.73)	16.71 (7.11)
Abundance (ind.10 ³ .km ⁻²)	1.87 (4.45)	20.29 (52.5)	4.23 (5.9)	8.3 (7.11)	19.18 (41.64)	20.28 (34.49)	18.31 (34.04)
Biomass (kg.km ⁻²)	54.36 (173.55)	249.75 (555.39)	43.81 (74.95)	78.15 (51.36)	91.75 (119.64)	132.14 (132.46)	96.82 (157.1)

1043

1044 Table A.3: Summary of structural indices for epi-bentho-demersal fish communities

1045 (average (standard-deviation))

Index	South coast	North coast	inner shelf	Grande Vasière	Northern middle shelf	Northern outer shelf	Southern outer shelf
Shannon's diversity	2.12 (0.67)	2.09 (0.61)	1.95 (0.84)	1.79 (0.69)	2.13 (0.6)	2.02 (0.54)	2.2 (0.66)
Pielou's evenness	0.6 (0.17)	0.5 (0.14)	0.52 (0.21)	0.43 (0.15)	0.51 (0.14)	0.52 (0.13)	0.56 (0.16)
Taxonomic richness	12.42 (4.41)	19.07 (4.6)	14.44 (4.8)	18.24 (3.67)	18.15 (2.99)	15.16 (3.14)	15.53 (3.53)
Abundance (ind.10 ³ .km ⁻²)	4.91 (5.88)	20.27 (23.28)	11.17 (16.91)	22.86 (29.76)	13.1 (9.68)	22.6 (136.18)	7.84 (6.58)
Biomass (kg.km ⁻²)	480.1 (728.79)	1302.27 (1391.75)	693.75 (914.57)	962.49 (554.5)	945.35 (564.44)	1039.93 (832.23)	760.45 (555.91)
8							





Figure A.1: Dendrogram inertia for epibenthic megainvertebrates





Figure A.2: Dendrogram inertia for bentho-demersal fish





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1056

Figure A.5: Variance partitioning of the effect of each force in shaping benthos
 communities. Physical forces are temperature, current, and mixed layer depth; chemical
 forces are oxygen, salinity, and net primary production.



1060

1061Figure A.6: Variance partitioning of the effect of each force in shaping fish community1062structure. Physical forces are temperature, current, and mixed layer depth; chemical1063forces are oxygen, salinity, and net primary production.