- 1 **Title:** Benthic food resources and the condition of bentho-demersal fish: spatial trends and relationships
- 2 in the northern shelf of the Bay of Biscay.
- 3
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19 Abstract:

20 To understand the trophic functioning of continental shelves, it is necessary to evaluate the factors that 21 drive production at different levels. For bentho-demersal fish, one such factor is spatial variability in the 22 biomass of their macrobenthic prey. In this study, we investigated the distribution of the macrobenthos 23 in the northern part of the continental shelf of the Bay of Biscay in order to determine how variability in 24 these prey resources might affect the bentho-demersal fish that feed on them. Due to the scarcity of 25 available field data, an empirical graph model and a biochemical model were used to produce different 26 indices of macrobenthic biomass, which were then used to generate three maps illustrating the 27 distribution of this biomass in the Bay of Biscay. In general, values from these maps showed high 28 Spearman correlations with the limited data available, but also discrepancies at greater depths where 29 field data are quite poor. Globally, a pronounced decline in macrobenthic biomass was observed from 30 the coast to offshore, but with particularly high values in deeper muddy sediments. Among eight fish 31 species analyzed, the four most known for their strongly benthivorous diet demonstrated positive 32 relationships between body condition and the macrobenthic biomass indices, with the strongest 33 associations detected for smaller-sized fish. This suggests that biomass index maps may be i) indicative 34 of the availability of benthic prey at the level of the continental shelf along a major coastal-offshore 35 gradient, and ii) useful for studying spatial variability in the transfer of energy or matter along the food 36 chain within this ecosystem. However, there is a clear need for macrobenthic data from field surveys at 37 the shelf scale, notably in deep circalittoral areas, to confirm or improve the accuracy of the spatial 38 biomass trends estimated by models.

39 Highlights:

- 40 Estimates of macrobenthic biomass exhibited coastal-to-offshore trend on the shelf.
- 41 The sparse field data macrobenthic biomass correlated with data from models.
- 42 In deeper areas, muddy sediments contained the greatest macrobenthic biomass.
- Body condition of benthivorous fish improved with increased macrobenthic biomass.
- Diet literature suggests biomass maps reflect food resources beyond macrofauna.
- 45 **Key words:** Macrobenthic biomass, fish, body condition, continental shelf, food resource, Bay of
- 46 Biscay

48 Introduction

49 The northwest Atlantic shelf is a vast area whose location between the continent and the ocean creates 50 strong environmental gradients that influence ecological processes such as matter production and 51 transfer (Hall, 2002). Within this ecosystem, benthic invertebrates play a preeminent role as a link 52 between primary producers and top predators (Chen et al., 2021), particularly bentho-demersal fish. 53 From the coast to the shelf break, strong declining gradients are typically observed in levels of nutrients, 54 organic matter, and primary production, and it is generally assumed that the biomass and/or production 55 of benthic invertebrates follow the same trend (Hall, 2002; Wei et al., 2010). Given the role of these 56 invertebrates as prey for bentho-demersal fish (Amezcua et al., 2003; Armstrong, 1982; Day et al., 57 2019), it is likely that such spatial trends in food resources will have repercussions for fish populations 58 across the continental shelf.

59 The northern part of the continental shelf of the Bay of Biscay offers an ideal location to study spatial 60 trends in these prey resources--particularly in the macrobenthic component--and the consequences for 61 the bentho-demersal fish that depend on them for food (e.g., Lagardère, 1987; Le Loc'h and Hily, 2005). 62 In this area, there is a strong terrigenous influence from the numerous small coastal rivers that flow onto 63 the coastal fringe as well as two larger rivers, the Vilaine and the Loire (Borja et al., 2019). Given this, it 64 is logical to expect a coastal-to-offshore gradient in the production or biomass of benthic invertebrates. 65 However, to confirm this kind of trend, it would be necessary to have data covering the entire study area, as was done in the North Sea by Reiss et al. (2010). Unfortunately, the available data on macrobenthic 66 67 biomass in the Bay of Biscay are scattered in time and mainly restricted to coastal areas or to specific 68 shelf habitats (e.g., mid-shelf fishing grounds) (Kopp et al., 2015; Le Loc'h et al., 2008; Rigolet et al., 69 2014; Robert, 2017). In such a case, though, it may be possible to estimate biomass using established 70 models, such as the empirical model developed by Chassé and Glémarec (1976) based on depth and 71 sedimentary substrates. More recently, mechanistic models like ERSEM (Kay et al., 2018) have also 72 shown promise for predicting the spatial variability of macrobenthic biomass (e.g., Timmermann et al., 73 2012).

Higher-quality habitats, such as those offering more food resources, are generally associated with
improved body condition among predators (Engelhard et al., 2013; Liao et al., 1995). Therefore, it may
be that patterns of abundance or scarcity of macrofaunal food resources across the shelf could translate

77 into spatial variability in the body condition of bentho-demersal fish predators. Within the Bay of Biscay, 78 bentho-demersal fish communities have been well described (Souissi et al., 2001), and the diets of most 79 species are known (e.g., Rault et al., 2017; Day et al., 2020). With this information, it is possible to 80 characterize the extent to which each species might be dependent on macrobenthic invertebrates. 81 Moreover, thanks to the EVHOE sampling surveys (Laffargue et al., 2020), biometric data are available 82 for multiple bentho-demersal fish species across several years. By analyzing patterns in macrobenthic 83 biomass and the body condition of bentho-demersal fish-and taking into account the diet plasticity of 84 these fish with respect to benthic macrofauna-it might be possible to detect the influence of spatial 85 trends in benthic food resources on higher trophic levels. Such information can then be used to shed 86 light on the broader trophic functioning of continental shelves.

The work presented here aimed to (1) describe spatial variability in macrobenthic biomass at the shelf scale in the Bay of Biscay based on data from models, (2) compare these data to the limited field data available, and (3) assess whether macrofaunal biomass data generated by such models exhibit a relationship with the body condition of bentho-demersal fish, and might thus be representative of the underlying availability of food resources.

92 **2. Materials and Methods**

93 2.1. Macrobenthic biomass indices



2.1.1. From an empirical biomass model (Chassé and Glémarec, 1976)



95

96 Figure 1: Graph model from Chassé and Glémarec (1976). Y-axis corresponds to logged bathymetric
 97 depth, and x-axis to granulometry divided into broad substrate categories.

98

99 The first model used to generate macrobenthic biomass indices was based on a graph of
100 biomass isopleths created from grab samples in the Bay of Biscay (Chassé and Glémarec, 1976) (Figure
101 1). Depth data came from depth maps (GEBCO Bathymetric Compilation Group, 2023). This model is
102 subsequently referred to as the "graph" model.

Biomass values were digitized from this graph (Figure 1) for each substrate category (silted heterogeneous sand (SHV), heterogeneous gravel (GH), clean/muddy gravel (GP/GV), coarse sand (SG), fine sand (SF), muddy fine sand (FV), sandy mud (VS), and pure mud (VV)), and for each bathymetric stratum (with a step of 10 m between 0 and 100 m depth, then from 100 to 200 m). Once rasterized, each cell was assigned a biomass value (in g.m⁻² dry weight (DW)) corresponding to the lower-bound isopleth. The maximal cell value was 32 g.m⁻² DW, associated with shallow muddy fine sand (FV). Finally, macrobenthic biomass values were assigned to each sub-section (substrate category
and bathymetric stratum), corresponding to the averaged biomass value over the proportion of cells.
This process provided an average value of macrobenthic biomass for each substrate category and
bathymetric stratum.

113 The first substrate map used was the 'Atlas du littoral français: Atlas des fonds meubles du 114 plateau continental du Golfe de Gascogne [Atlas of the French coastline: Atlas of the soft bottoms of the 115 continental shelf of the Bay of Biscayl' obtained from the same study as the graph model described 116 above (Chassé and Glémarec, 1976). The map of biomass indices resulting from this first substrate map 117 is hereafter referred to as the 'Biomass₁₉₇₆ map'. The second substrate map used is more recent than 118 the first (Vasquez et al., 2023) and is based on a more-extensive dataset describing the nature of the 119 seabed. We adapted the substrate categories in this map to correspond to those of the biomass graph 120 model based on a comparative analysis of the two substrate maps (results are in Supplementary 121 Materials 1). The second map of biomass indices is subsequently referred to as the 'Biomass₂₀₂₃ map'.

122 Some approximations were necessary for certain areas (5% of the shelf surface for Biomass₁₉₇₆ 123 and Biomass₂₀₂₃ maps) where the substrate x bathymetry intersection was not represented in the graph 124 model, and biomass data were not available. These areas were approximated by another substrate with 125 similar levels of macrobenthic biomass and similar trends along the depth gradient. For the Atlas 126 substrate maps, biomass values for muddy/clean gravel (GP/GV) were approximated by biomass from 127 coarse sand (SG) below 80 meters depth, and biomass values from muddy heterogeneous sand (SHV) 128 were approximated by biomass from muddy fine sand (FV) below 50 meters depth. For the more-recent 129 substrate maps, biomass values of gravel and coarse sand from the graph model were averaged 130 (GP/GV and SG) for the "coarse substrate" category. Descriptions of the substrate categories of the 131 graph model and the two substrate maps are available in Supplementary Materials 2.

132

2.1.2. From a biogeochemical model (POLCOMS-ERSEM)

The second set of modeled data used to generate a macrobenthic biomass index came from a mechanistic biogeochemical model. The data originated from the outputs of the POLCOMS-ERSEM model, a regional high-resolution coupled hydrodynamic-ecosystem model for low trophic levels, driven by a reanalysis of data from the period 2000–2015 (Kay et al., 2018). POLCOMS-ERSEM links a general ocean circulation model, POLCOMS (the Proudman Oceanographic Laboratory Coastal Ocean Modelling System) and a marine ecosystem model, ERSEM (European Regional Seas Ecosystem Model; Butenschön et al., 2016). It covers the European shelf seas, with a horizontal resolution of 0.1° (around 11 km). As we are interested in the biological part of the model, we will refer to it subsequently as the ERSEM model. In this model, three groups represent the bulk of macrobenthic biomass, and they were initially calibrated using data estimated by Bryant *et al.* (1995) based on macrobenthic biomass values in the North Sea. We selected deposit- and suspension-feeding groups as the macrobenthic components of interest (the only other group, 'meiobenthos', was not considered).

Biomass values were summed from all suspension and deposit feeders in ERSEM to create a macrobenthic biomass index. Macrobenthic biomasses were averaged over the last 10 years (2006– 2015) to minimize potential yearly effects. The macrobenthic biomass index from ERSEM was expressed in gC/m² and the map of these values is referred to here as the "Biomass_{ERSEM} map".

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2. 2. Biological field data

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2.2.1. Biomass of macrobenthic invertebrates

Macrobenthic biomass indices from the three biomass maps described above were compared to field data collected in the circalittoral area (*i.e.*, less than 15 m depth) of the northern part of the shelf of the Bay of Biscay. Table 1 provides a summary of the various surveys from which data on macrobenthic biomass were obtained; the spatial distribution of sampled stations (from 2 to 5 grabs per station) is presented in Figure 2. As seen in this figure, these data are very sparse over both time and space, and no single survey covers the full extent of the continental shelf.

Table 1: Summary of different macrobenthic surveys in which biomass data were collected, with the gear used, the time of the year, the year sampled, the average bathymetry (and its range), and the number of stations for each.

Survey	Sampling gear	Time	Bathymetry	Ν	Reference
RetroB	Van Veen grab	March (2008)	20.1 (15-25.6)	17	(Le Bris, 2008)
Nurse2008	Van Veen grab	August - September (2008)	21.8 (16.3-35.3)	20	(Brind'Amour, 2008)
PLOOPS 1	Van Veen grab	July (2009)	27.6 (23-35)	18	(Dubois, 2010a)
PLOOPS 2	Van Veen grab	February - November (2010)	25.5 (21.3-32.8)	24	(Dubois, 2010b)
Isobaie 2	Van Veen grab	June (2010)	77.3 (18.9-154)	11	(Le Bris, 2010)
FEBBE 1	Day grab	May - August (2013)	101.3 (78-113)	18	(Laffargue, 2013a)
FEBBE 2	Day grab	August (2013)	101.9 (91-113)	23	(Laffargue, 2013b)
APPEAL1	Hamon grab	May (2018)	60.7 (53-74)	15	(Le Loc'h and Grall, 2018a)
APPEAL 2	Hamon grab	August (2018)	59.2 (52-73)	15	(Le Loc'h and Grall, 2018b)
APPEAL 4	Hamon grab	September (2019)	109.5 (98.4-125.1)	12	(Le Loc'h and Grall, 2019)

161 RetroB and Nurse2008 were two surveys performed in a coastal bay (Bay of Vilaine) 162 (Brind'Amour, 2008; Le Bris, 2008), in a shallow circalittoral area under the terrigenous influence. 163 PLOOPS 1 and 2 surveyed a coastal area in the Bay of Concarneau (Dubois, 2010a; Dubois, 2010b), 164 which, although it also features a shallow circalittoral fringe, has less terrigenous influence compared to 165 the Bay of Vilaine. APPEAL 1 and 2 focused on the deeper limit of the shallow circalittoral fringe, to the 166 south of the island of Groix (Le Loc'h and Grall, 2018a; Le Loc'h and Grall, 2018b). APPEAL 4 and 167 FEBBE 1 and 2 surveyed a deep circalittoral muddy bank, the so-called "Grande Vasière" (Laffargue, 168 2013a; Laffargue, 2013b; Le Loc'h and Grall, 2019). Finally, Isobaie 2 covered an coastal-to-offshore 169 transect from the Loire estuary and Bay of Vilaine towards the shelf break (Le Bris, 2010).



- 170
- Figure 2: Locations of macrobenthic field observation stations on the northern shelf of the Bay of Biscay.
 Different surveys are represented with different colors. The light gray shaded area is the mid-shelf mud
 belt known as the "Grande Vasière".

Each survey counted taxa down to the lowest taxonomic level, and biomass was measured for most of the taxa, sampled stations, and studies. When individual mass was not available, it was approximated using values from the closest phylogenetic relative from the same station or from a neighboring station of the same survey. Biomass values in wet weight were converted into ash-free dry 178 weight using Brey's conversion factors (www.thomas-brey.de/science/virtualhandbook). For each 179 station, observed macrobenthic biomass was the average of the values obtained from two to five 180 replicates.

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2.2.2. Data on individual bentho-demersal fish

182 Individual biometric data (total length and mass) from autumnal EVHOE surveys (between 2008 183 and 2022, onboard Thalassa R/V) were used to assess the body condition of a limited selection of 184 bentho-demersal fish species. Body condition is ultimately a reflection of a fish's habitat quality, and 185 consequently also of food resource availability (Cavraro et al., 2019; Karlson et al., 2018; Liao et al., 186 1995; Schloesser and Fabrizio, 2019). We examined most of the benthivorous species for which 187 sufficient data were available across the continental shelf (Supplementary Materials Figure SM4 and 188 SM5). Four demersal species were considered: whiting (Merlangius merlangus), striped red mullet 189 (Mullus surmuletus), poor cod (Trisopterus minutus), and pout (Trisopterus luscus). Four benthic taxa 190 were also considered: scaldfish (Arnoglossus spp.), small-spotted catshark (Scyliorhinus canicula), 191 common sole (Solea solea), and thickback sole (Microchirus variegatus). Arnoglossus spp. includes 192 both A. laterna and A. imperialis, which are assumed to have a similar diet (Castro et al., 2013), and the 193 combination of the two made it possible to cover a wide coastal-to-offshore gradient.

- 194 **2.3. Data analysis**
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2.3.1. Comparison of macrobenthic biomass indices from models

196 Once we had generated the maps of the three macrobenthic biomass indices, we compared the 197 spatial patterns presented. Each map was standardized and gridded with the 0.1° x 0.1° ERSEM grid, 198 and each grid cell was assigned an average biomass index value (weighted by the surface area of each 199 biomass value within the grid cell). Biomass_{ERSEM}, Biomass₁₉₇₆, and Biomass₂₀₂₃ were then compared 200 using Spearman correlations. Specifically, for a given cell, a buffer area of 5 cells was considered 201 (resulting in a 25-cell window) and Spearman correlations were computed for the values of each 202 biomass map in this window, allowing us to highlight potential differences in spatial trends between the 203 different maps. Negative values meant that the two maps in question depicted opposite trends in 204 macrobenthic biomass. All analyses were conducted using R 4.3.1 (R core team, 2023); maps were 205 processed using the raster and sf packages (Hijmans, 2023; Pebesma, 2018).

206 2.3.2. Comparison between field data on macrobenthic biomass and biomass indices207 from models

Macrobenthic biomass indices from the three maps generated here were compared to values of macrobenthic biomass from field datasets. For this, the nearest macrobenthic biomass values from each of the three maps were assigned to each field station sampled, and these values were averaged within a 500-m buffer surrounding the station. Values from biomass index maps and field values were log-transformed and normalized so that the final data ranged between 0 and 1. Then, Spearman correlations were calculated between macrobenthic biomass values from field data and indices from models.

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2.3.3. Relationship between body condition and macrobenthic biomass indices

Fish body condition was assessed using the body condition factor developed by Le Cren (1951), which represents 'the deviation of an individual from the average weight for length' in a given population. Le Cren's condition factor, K_n, compensates for allometric changes, enabling comparisons of body condition between small and large individuals. Indeed, the response of body condition to macrobenthic resource availability may vary in a size-dependent manner due to trophic shifts and diet plasticity.

For each species, log length-mass relationships (LMR) were fitted using individual biometric data. Observations with a Cook's distance greater than 4/n (n = number of observations) were considered outliers and removed (Altman and Krzywinski, 2016). For each species, homogeneity of variance and the normal distribution of residuals were checked visually.

K_n is the observed individual mass divided by the predicted mass according to each species'
 LMR. To take into account ontogenic variations, individuals were split into two size groups, to reflect the
 trophic size thresholds reported in the literature for six of the fish species (Table 2).

Finally, we created mixed-effect linear models to assess the relationship between the body condition factor and the macrobenthic biomass indices. One model was computed for each of the eight fish taxa and the three macrobenthic biomass index maps (Biomass₁₉₇₆, Biomass₂₀₂₃, and Biomass_{ERSEM}), and fitted using the Ime4 R package (Bates et al., 2015). Models were computed with station as a random effect to account for intra-station correlations (Zuur et al., 2009) and also included a size category effect (*i.e.*, Small or Large), with the exception of the models for *Arnoglossus* spp. and 234 Microchirus variegatus. Each station was assigned the average value of macrobenthic biomass within 235 a 5000-m buffer around its location. The sensitivity of results to this buffer was explored, and no major 236 differences were observed when comparing results from 1-km to 50-km buffers. Macrobenthic biomass 237 indices were scaled to allow the comparison of their respective effects on K_n. Chi-squared tests were 238 used to evaluate these results against the null hypothesis (*i.e.*, a model without a biomass effect). During model evaluation, mixed-effect linear models were refitted using maximum likelihood, and only 239 240 significant models (p-value < 0.05) were kept. Additionally, the variance explained by selected mixed-241 effect linear models was computed using conditional R-squared and marginal R-squared values; 242 conditional R-squared included random and fixed effects of the model, while marginal R-squared 243 corresponded to the variance explained only by the fixed effect (Zuur et al., 2009).

Since the spatial extent of the biomass maps varied, the amount of data per species and size class available for the mixed-effect linear models also differed. The quantity of data used for each mixedeffect linear model and each macrobenthic biomass model is provided in Table SM6.

247 Table 2: Size thresholds used to account for ontogenic shifts in different fish species.

Species	Size shift	Trophic shift	Reference	
Arnoglossus spp.	-		(Schückel et al., 2011)	
Merlangius merlangus	20	Shift from benthic invertebrates to fish.	(Hislop et al., 1991)	
Microchirus variegatus	-		(Amezcua et al., 2003)	
Mullus surmuletus	18	Shift from benthic invertebrates to shrimp,	(Bautista-Vega et al.,	
		ophiurids, and polychaetes.	2008)	
Seuliorhinus conicula	22	Shift from small invertebrates to larger	(Šantić at al. 2012)	
Scynonninus camcula		decapods, cephalopods, and fish.	(Santic et al., 2012)	
	30	Shift toward epibenthic decapods and higher	(Fanelli et al., 2022)	
Solea Solea		trophic level.		
	25	Shift from benthic invertebrates towards	(Armstrong, 1982)	
msoplerus iuscus		shrimp and fish.		
	17	Shift from small crustaceans to larger	(Armstrong 1092)	
msopierus minutus	17	decapods and fish.	(Annstrong, 1962)	

249 3. Results

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3.1. Maps of macrobenthic biomass indices



- 253 254 Figure 3: Maps of macrobenthic biomass indices obtained from the graph and ERSEM models. White areas in the Biomass₁₉₇₆ and Biomass₂₀₂₃ maps correspond to hard bottom substrate, where soft bottom 255 macrobenthic biomass could not be assessed. Gray lines represent the 100- and 200-meter isobaths.
- 256
- 257 The Biomass₁₉₇₆ and Biomass₂₀₂₃ maps highlighted high values of macrobenthic biomass along 258 the coast, in concentrated areas characterized by shallow patches of mud, sandy mud, and muddy sand

substrate. Instead, index values from the Biomass_{ERSEM} map were higher along a coastal fringe ranging
from the Pertuis-Charentais to the peninsula of Quiberon (Figure 3).

261 The Biomass_{ERSEM} map depicts a decreasing gradient in biomass towards the shelf break, inversely 262 related to water depth, while the Biomass₁₉₇₆ and Biomass₂₀₂₃ maps highlight patches of substrate at 263 moderate depths that support higher macrobenthic biomass. Overall, the Biomass₁₉₇₆ and Biomass₂₀₂₃ 264 maps were highly correlated (Spearman correlation = 83.09%), while the Biomass_{ERSEM} map exhibited 265 a weaker correlation with the two others (49.82% and 62.84%, respectively). The most substantial 266 differences were noted between the Biomass₁₉₇₆ and Biomass_{ERSEM} maps; patches corresponding to 267 certain habitats (e.g., the large mud belt in the central part of the shelf, or at the edge of the continental 268 shelf) exhibited opposing trends in macrobenthic biomass between these two maps (Figure 4).



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- 273

3.2. Biomass indices from models versus field biomass data

274 Despite differences in absolute trends, positive Spearman correlations were detected between 275 the three indices of macrobenthic biomass and field data on macrobenthic biomass (60.4%, 66.9%, and 276 77.5% for Biomass₁₉₇₆, Biomass₂₀₂₃, and Biomass_{ERSEM}, respectively; Figure 5). Based on the ERSEM 277 model, the highest level of biomass was predicted for stations located at 15–20 m depth. Instead, the 278 graph model (Biomass₁₉₇₆ and Biomass₂₀₂₃) estimated the highest levels of biomass at a depth of 20-50

Figure 4: Maps of local Spearman correlations, with blue representing negative correlations and red positive correlations. Gray lines represent the 100- and 200-meter isobaths.

m, and similar, lower, levels between stations at 15–20 m and 50–75 m. Most of the stations that were
estimated to have high biomass levels (from Biomass₁₉₇₆ and Biomass₂₀₂₃) were sampled as part of the
PLOOPS surveys (Figure SM7). The bathymetric gradient previously noted in the Biomass_{ERSEM} index
is also evident in Figure 5, as stations tended to aggregate along the y-axis according to their
bathymetry. This pattern was less marked for the Biomass₁₉₇₆ and Biomass₂₀₂₃ indices, which generated
similar estimates of biomass at depths of 50–75 m and 75–100 m.



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Figure 5: Modeled macrobenthic biomass indices versus field data on macrobenthic biomass. Stations
 are colored according to their bathymetric range, with red shades for shallower depths and blue for
 deeper ones. The red dashed line corresponds to the isoline.

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291

3.3. Relationship between body condition and macrobenthic biomass indices

292 Of the numerous mixed-effect linear models we constructed to test the effect of the 293 macrobenthic biomass indices as predictors of body condition in various fish species, 15 detected 294 significant effects, *i.e.*, a macrobenthic biomass index explained significantly more variance than the null 295 model (Table 3). No such models were found for Trisopterus minutus and Scyliorhinus canicula, and 296 only one significant model was obtained for Microchirus variegatus and for Merlangius merlangus. 297 Instead, significant positive relationships were found between the three macrobenthic biomass indices 298 and body condition factors for the species Arnoglossus spp., Mullus surmuletus, Solea solea, and 299 Trisopterus luscus. The level of variance explained remained globally low, with the maximum observed 300 for the relationship between body condition in *Mullus surmuletus* and the Biomass₁₉₇₆ index (14.7%) 301 (Table 3). On average, Mullus surmuletus and Arnoglossus spp. were the two taxa with the highest 302 amount of variance explained by macrobenthic biomass indices.

Table 3: Summary of mixed-effect linear models that identified a significant relationship between macrobenthic biomass indices and body condition factors of fish. Chi2 and p-value correspond to the test of the model against a null model including only an intercept and the random effect of station. R2m is the marginal R-squared value and R2c is the conditional R-squared value.

Macrobenthic index	Species	R2m	R2c	Chi2	p-value
	Arnoglossus spp.	0.143	0.419	5.4	2.01e-02
Biomass	Mullus surmuletus	0.147	0.327	14.94	1.87e-03
Biointa33 ₁₉₇₆	Solea solea	0.038	0.197	18.46	3.53e-04
	Trisopterus luscus	0.066	0.183	8.11	4.38e-02
	Arnoglossus spp.	0.073	0.383	4.01	4.52e-02
Biomassa	Merlangius merlangus	0.014	0.183	9.22	2.65e-02
Biointa332023	Mullus surmuletus	0.085	0.313	15.54	1.41e-03
	Solea solea	0.034	0.18	19.84	1.83e-04
	Trisopterus luscus	0.072	0.229	20.18	1.56e-04
Biomassenser	Arnoglossus spp.	0.126	0.43	5.87	1.54e-02
DIOILIGSEKSEM	Microchirus variegatus	0.131	0.464	5.33	2.10e-02
	Mullus surmuletus	0.1	0.301	23.67	2.93e-05

	Solea solea	0.038	0.184	20.22	1.53e-04
-	Trisopterus luscus	0.085	0.238	22.64	4.80e-05

308 In all taxa in which a significant relationship was detected between body condition and macrobenthic 309 biomass, the relationship was positive; the only exception to this was found in *M. merlangus*. 310 Additionally, these relationships were stronger (steeper slope) for smaller-sized individuals, although 311 this pattern was identified for large individuals of *Mullus surmuletus* as well.



313 Figure 6: Significant linear mixed-effect models selected. Data used to fit models are shown here; 314 orange dots correspond to small individuals and blue triangles to larger ones.

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317 The only species for which K_n exhibited a steep negative relationship with macrobenthic biomass was 318 *Merlangius merlangus*. For *Solea solea*, instead, larger individuals seemed to show higher values of K_n, 319 with only a very weak (positive or negative depending on the biomass index) relationship with 320 macrobenthic biomass indices.

321 **4.** Discussion

322 The northern part of the shelf of the Bay of Biscay is a particularly productive ecosystem, 323 supporting both pelagic and benthic trophic pathways (Cresson et al., 2020) that provide essential 324 resources for fish species. One such resource is the macrobenthos; these organisms are a key food 325 source for bentho-demersal fish, which tend to be opportunists that will eat whatever benthic prey is 326 available (Day et al., 2019; Rault et al., 2017). However, spatial variability in macrobenthic biomass on 327 this shelf remains largely unexplored. To date, quantitative data on macrobenthic biomass in the Bay of 328 Biscay are limited to very coastal areas (e.g., Dutertre et al., 2013) or specific habitats (e.g., muddy 329 areas like the Grande Vasière; (Le Loc'h et al., 2008). The benthic characteristics of these areas have 330 been extrapolated to the entire shelf through ecosystem modeling (Lassalle et al., 2011). The present 331 study aims to fill this gap by analyzing the spatial distribution of macrobenthic biomass over the whole 332 northern shelf of the Bay of Biscay and evaluating the relationship between variations in this biomass 333 and the fish that feed (to a greater or lesser degree) on the macrobenthos. The lack of field data on 334 macrofaunal biomass in much of the study area prompted us to explore and use existing models, of 335 which two were particularly relevant to our objective and the marine region we were studying.

336 The empirical "graph model" from Chassé and Glémarec (1976) is based on depth and sediment 337 typology, with the assumption that hydro-sedimentary characteristics drive ecological processes and 338 features of macrobenthic communities (Glémarec, 1969; McArthur et al., 2010; Reiss et al., 2015). The 339 graph model was defined on the basis of a limited number of sediment classes, reflecting the substrate 340 distribution presented in the "Atlas des fonds meubles du plateau continental Nord Gascogne" [Atlas of 341 the French coastline: Atlas of the soft bottoms of the continental shelf of the Bay of Biscay] (Chassé and 342 Glémarec, 1976). This modeling strategy is comparable to the habitat suitability modeling framework 343 adopted by Degraer et al. (2008), since the map involved an initial biophysical dataset and spatial 344 interpolation from this dataset with physical data maps.

In the ERSEM model, which is part of the POLCOMS-ERSEM biophysical model, macrobenthic biomass of suspension and deposit feeders is the direct result of flows from ecological processes (primary production, organic matter sedimentation) that supply the seafloor. Bentho-pelagic coupling models like the ERSEM model have already demonstrated their ability to accurately represent macrobenthic biomass at a small scale (Ehrnsten et al., 2019). This model also incorporates other

350 processes, such as organic matter ingestion or bioturbation (Butenschön et al., 2016)-but not the 351 remobilization of buried organic matter (Butenschön et al., 2016)-that could boost macrobenthic 352 biomass (Qin et al., 2010) and benefit the benthos during periods of low input (Zhang et al., 2019). 353 However, the ERSEM model is only based on muddy sediments and, unlike the graph model, does not 354 account for spatial variability in substrate types, despite the critical role these characteristics play in the 355 modeled processes and in determining benthic invertebrate biomass. Indeed, the nature of a substrate 356 reflects numerous features and processes (e.g., permeability, deposition, resuspension, organic 357 content, among others) (Chen et al., 2018; Idier et al., 2010; Pace et al., 2021) that can influence 358 macrobenthic characteristics and biomass (Ehrnsten et al., 2019; Pearson and Rosenberg, 1978).

359 The graph and ERSEM models reproduced biomass gradients that were largely similar to each 360 other, particularly in the overall decrease from the coast to the shelf break and the presence of 361 heterogeneous coastal zones with high biomass. This gradient is indicative of a significant terrigenous 362 influence, one whose intensity increases with proximity to river outlets (Loire and Vilaine estuaries), 363 particularly in the southeastern region of the study area. However, there were substantial differences 364 between the two types of models. The two maps derived from the graph model showed discontinuities 365 in this biomass gradient, with distinct patches of high biomass values, particularly in the muddlest zone 366 (the Grande Vasière). With the graph model, the use of the most recent substrate map only marginally 367 changed the distribution of biomass; this was unsurprising given that these maps largely reproduce 368 substrate data from previous versions and retain high local uncertainties due to their extremely 369 heterogeneous spatial sampling coverage and limited sedimentary field data (Kaskela et al., 2019; 370 Pierrejean et al., 2022). Instead, the biomass distribution provided by the ERSEM model was much less 371 patchy, illustrating the more continuous nature of the spatial distribution of the variables used to estimate 372 biomass, especially because this model does not take into account sedimentary substrate features.

Based on an exhaustive census of surveys and research projects carried out over the last decade, we gathered all available field data on spatial variability in macrobenthic biomass in the study area. A preliminary analysis of the data highlighted not only spatial gaps (especially between 120 and 200 m) but also heterogeneity in the sampling gear used (3 types) and in the years (10 years from 2008 to 2019) and months/seasons (from March to November) in which samples were collected. Keeping in mind these limitations, when these data were standardized and examined at a broad scale, they seemed to describe a depth gradient roughly comparable to those we estimated from models (Spearman

380 correlations ranging from 60% to 80%). Even the limited amount of field data we were able to find is 381 useful in understanding the strengths or deficiencies of the models. For example, when comparing 382 coastal areas, the ERSEM model seemed to better reflect the reality of the observed biomass difference, 383 which was lower in the Bay of Concarneau (PLOOPS surveys) than in the Bay of Vilaine (RetroB and 384 Nurse2008 surveys). This difference was reversed in the graph model, which seemed to underestimate 385 macrobenthic biomass in pure muddy substrates such as those found in the Bay of Vilaine (Le Bris and 386 Glémarec, 1995). This may reflect the fact that terrigenous influences, which are known to increase 387 benthic biomass (Deubel et al., 2003; Escobar-Briones and Soto, 1997; Zuschin and Stachowitsch, 388 2009), were not taken into account in the graph model, which only considered depth and sediment type. 389 This point was discussed by Dutertre et al. (2013), who explained that in the shallow part of the Bay of 390 Biscay, although 26% of macrobenthic diversity patterns could be explained by substrate, 391 hydrodynamics, and depth, the inclusion of parameters quantifying riverine influence (Vilaine and Loire) 392 made it possible to explain an additional 16%.

For both types of models, there is a notable deficiency at the continental shelf edge. The absence of a peak in macrobenthic biomass at the shelf edge seems to indicate that the two models ignore the hydrological context of the shelf break, where biological production is stimulated by upwelling (Simpson and Sharples, 2012; Vandromme et al., 2014). Although limited field data were available in this area, some of the deepest field observations could indicate the presence of a peak; such a pattern would be consistent with estimates made by Le Loc'h *et al.* (2008) at three sampling stations in the slope area, which they compared to the highly productive mud zone in the middle of the continental shelf.

400 By analyzing the relationship between fish body condition and macrobenthic biomass indices, 401 our aim was to explore the impact of the distribution of food resources on the shelf as indicated indirectly 402 by the macrobenthic component. In our study, the relationship between macrobenthic biomass indices 403 and body condition factor (K_n) was significant and positive for four of the eight bentho-demersal fish 404 species tested. These relationships were similar for all models of macrobenthic biomass and for most 405 of the fish species examined, which emphasizes the importance of the coastal-to-offshore gradient in 406 the spatial distribution of macrobenthic food resources. It also suggests bottom-up processes in which 407 food availability may be a determining factor, as seen in other contexts (Brosset et al., 2015; Laurel et 408 al., 2001; Porath and Peters, 1997).

409 In many cases, the presence or absence of significant relationships can be explained by the 410 trophic or life-history characteristics of the species analyzed. Among the different species studied, 411 whiting (Merlangius merlangus), poor cod (Trisopterus minutus), and small-spotted catshark 412 (Scyliorhinus canicula) exhibited a negative or non-existent relationship between body condition and 413 biomass indices. For whiting, this could be explained by the fact that the diet of small individuals (less 414 than 22 cm) is both pelagic and benthic, while that of larger individuals is also based on pelagic and 415 demersal fishes (Day et al., 2019). This feeding characteristic is shared by the poor cod, which has a 416 very generalist diet featuring macrobenthic, suprabenthic, and pelagic prey, including fish at larger sizes 417 (Armstrong, 1982). The small-spotted catshark is also thought to feed on pelagic resources such as 418 euphausiids (Rodríguez-Cabello et al., 2007), which are often found near the continental slope of the 419 Bay of Biscay (Peña et al., 2019), where the fish in our study were mainly found. For these species, the 420 absence of significant relationships with macrobenthic biomass may reflect dietary compensation with 421 pelagic prey when benthic food resources are less available.

422 Compared to the other flatfish studied, the common sole (Solea solea) did not exhibit a strong 423 positive relationship between body condition and macrobenthic biomass indices, even though it is mostly 424 benthivorous (Lagardère, 1987; Rault et al., 2017). For small sole individuals, whose distribution is more 425 coastal than that of large individuals, it is difficult to find an explanation for this weak relationship. For 426 large individuals, which are mainly found further offshore (Figure SM5), the relatively high body condition 427 scores in an area of poorer macrobenthic biomass may be related to the fact that these individuals have 428 accumulated reserves in more coastal areas during the summer period (Rijnsdorp, 1990), before 429 migrating offshore in the autumn (where they were sampled) to reproduce a few months later (Le Bec, 430 1983). This may not be the case for the other fish that did show a relationship between their body 431 condition and macrobenthic biomass, such as Arnoglossus spp. and M. surmuletus, which are known 432 to reproduce later than the common sole (Gibson and Ezzi, 1980; N'Da and Déniel, 1993).

In the case of pout (*Trisopterus luscus*), only the smaller individuals demonstrated a strong positive relationship between body condition and macrobenthic biomass. The small individuals of this species are known to feed mainly on macrobenthic and suprabenthic invertebrates (Castro et al., 2013), while the diet of larger individuals includes fish (Armstrong, 1982). Scaldfishes (*Arnoglossus* spp.) and the two age classes of striped red mullet (*M. surmuletus*) also exhibited strong relationships between body condition and macrobenthic biomass indices. Scaldfishes feed mainly on suprabenthic

invertebrates (Castro et al., 2013), but also on macrofauna and even benthic fish (Déniel, 1981). Striped red mullet feed on epibenthic crustaceans (*e.g.*, Galatheidae and Portunidae) but mainly rely on more infaunal prey such as bivalves or polychaetes (Pavičić et al., 2018; Serrano et al., 2003). Unlike these other species, thickback sole (*M. variegatus*) demonstrated a relationship only between its body condition and the macrobenthic biomass index from the ERSEM map. Its diet relies heavily on benthic macroinvertebrates (Amezcua et al., 2003; King and Fives, 1990) and it is difficult to explain why the relationship was not established using the data from the other two maps.

446 With the diets of these species in mind, it seems justifiable to assume that the macrobenthic 447 biomass maps also represent a wider range of benthic prey, such as suprabenthic prey which are less 448 effectively sampled by grabs (Sorbe, 1981). The various relationships observed between fish body 449 condition and biomass maps illustrate both differences among fish species in their dependence on 450 benthic resources as well as variability in the availability of such resources. Certain areas (e.g., coastal 451 habitats or mudflats like the Grande Vasière) offer better feeding conditions for benthic-feeding fishes. 452 In such areas, competition between benthic feeders is reduced through processes such as individual 453 specialization (Sánchez-Hernández et al., 2021). Areas with scarcer benthic resources, such as deeper 454 zones (e.g., in the vicinity of the shelf edge), may instead experience heightened intra- and interspecies 455 competition (Holbrook and Schmitt, 1989). This could drive adaptive strategies such as shifts to (i) prey 456 with lower energy content, with consequences for fish growth and body condition (Cormon et al., 2016; 457 Hiddink et al., 2016; Kamimura et al., 2021) and/or (ii) prey at higher trophic levels as suggested by the 458 isotope study of Schaal et al. (2016). This highlights the trophic plasticity of fish with respect to conditions 459 of resource availability or competition (Pelage et al., 2022). Our study supports the idea that demersal 460 fishes generally depend on trophic resources at spatial scales that are smaller than that of the 461 continental shelf, a hypothesis that has already been corroborated by different approaches (e.g., 462 Chouvelon et al., 2012). Our body condition results are also consistent with the trends reported by 463 Schaal et al. (2016) regarding the trophic levels of demersal fishes along the coastal-to-offshore isotopic 464 gradient, which were characterized by a trophic shift that those authors hypothesized to be due to lower 465 prey abundance offshore.

467 **5.** Conclusion

468 By providing visual representations of the spatial distribution of benthic resource availability, the 469 macrobenthic biomass maps analyzed here could be valuable tools in the study of spatial variability in 470 trophic networks, bentho-pelagic coupling, and matter fluxes on the continental shelf. The two models 471 we used to estimate macrobenthic biomass had different strengths and limitations based on the different 472 modeling assumptions employed. Nevertheless, both models, which are based on different ecological 473 hypotheses, revealed similar coastal-to-offshore distribution trends, with notable differences in the mid-474 shelf and near-coastal areas. Unfortunately, few of Europe's continental shelves (e.g., in the North Sea; 475 Reiss et al., 2010) have been sampled thoroughly enough to effectively reveal benthic biomass 476 gradients on a large spatial scale. These data are crucial for understanding certain processes affecting 477 marine populations or, more recently, for the development of ecosystem models (e.g., Lassalle et al., 478 2011; Corrales et al., 2022; Lopez De Gamiz-Zearra et al., 2023), especially in deeper areas where 479 macrobenthic samples are limited (e.g., beyond the Grande Vasière in the Bay of Biscay).

480 To better understand the functioning and status of fish populations, our study highlighted the 481 importance of recognizing spatial variability in the macrobenthic resources on which these fish depend. 482 The coastal-to-offshore gradient in macrobenthic biomass, and its relationship with body condition in 483 certain bentho-demersal fish, raises an important question about the viability of deeper populations. This 484 question is all the more crucial in the context of climate change: for example, deepening mechanisms 485 have been clearly demonstrated in bentho-demersal fish on the continental shelves of the northeast 486 Atlantic (e.g., Dulvy et al., 2008) and their impact on the populations of the species concerned has yet 487 to be fully assessed.

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Figure SM1: Relative proportion of the area of each Atlas map substrate polygon contained in each Euseamap substrate polygon.

Table SM2: Different substrate categories of different substrate maps and the correspondence between
 them and the graph model.

Graph mod	el Atlas biosedimentary substrate categories	Euseamap substrate
substrate categorie	s	categories
	Roches Paléozoïques ; Roches Tertiaires	Rock or other hard substrata;
		Seabed; Worm reefs
GP-GV	Graviers Envasés; Graviers Envasés Calcaires;	Coarse substrate
	Graviers Propres ; Graviers Propres Calcaires	
GH	Graviers Sableux Hétérogènes; Graviers Sableux	
	Hétérogènes Calcaires	
SG	Sables Grossiers ; Sables Grossiers Calcaires	Coarse substrate
SF	Sables Dunaires ; Sables Dunaires Calcaires ; Sables	Sand
	Fins ; Sables Fins Calcaires	
SHV	Sables Envasés Hétérogènes; Sables Grossiers	
	Envasés	
FV	Sables Fins Envasés ; Sables Fins Envasés Calcaires	Muddy sand
VS	Vases Sableuses ; Vases Sableuses Calcaires	Mixed sediment
		Sandy mud
VV	Vases ; Vases Calcaires	Fine mud

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Figure SM3: The location of sampling stations from various surveys along the continental shelf of the
Bay of Biscay. Panel A depicts values of the macrobenthic biomass index obtained from the graph model
based on 'Atlas' substrate maps, B shows values from the graph model based on a recent substrate
map, and C is the average biomass of suspension and deposit feeders estimated by the POLCOMSERSEM model between 2006–2015. Gray lines represent the 100- and 200-meter isobaths.



Figure SM4: Spatial distribution of available data on bentho-demersal fish condition. Gray lines represent the 100- and 200-meter isobaths.



870 Figure SM5: Relative proportion of 'Small' and 'Large' size-class categories in the available data on 871 bentho-demersal fish condition. Gray lines represent the 100- and 200-meter isobaths.

- 873 874 875 Table SM6: Number of samples included in the analysis, based on the biomass model used---"graph
 - model" (Biomass₁₉₇₆, Biomass₂₀₂₃) or ERSEM model (Biomass_{ERSEM})—and the fish size categories, small (S) or large (L).

Species	Number of individuals
	Biomass ₁₉₇₆ : 45
Arnoglossus spp.	Biomass ₂₀₂₃ : 67
	Biomass _{ERSEM} : 67
	Biomass ₁₉₇₆ : 253 (S) & 975 (L)
Merlangius merlangus	Biomass ₂₀₂₃ : 268 (S) & 1192 (L)
	Biomass _{ERSEM} : 268 (S) & 1192 (L)
	Biomass ₁₉₇₆ : 32
Microchirus variegatus	Biomass ₂₀₂₃ : 48
	Biomass _{ERSEM} : 48
	Biomass ₁₉₇₆ : 95 (S) & 91 (L)
Mullus surmuletus	Biomass ₂₀₂₃ : 211 (S) & 123 (L)
	Biomass _{ERSEM} : 211 (S) & 123 (L)
	Biomass _{1976:} 12 (S) & 683 (L)
Scyliorhinus canicula	Biomass ₂₀₂₃ : 35 (S) & 1094 (L)
	Biomass _{ERSEM} : 35 (S) & 1098 (L)
	Biomass ₁₉₇₆ : 418 (S) & 121 (L)
Solea solea	Biomass _{2023:} 490 (S) & 140 (L)
	Biomass _{ERSEM} : 490 (S) & 140 (L)
X	Biomass ₁₉₇₆ : 217 (S) & 204 (L)
Trisopterus luscus	Biomass ₂₀₂₃ : 274 (S) & 309 (L)
	Biomass _{ERSEM} : 274 (S) & 310 (L)



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878 Figure SM7 Biplot of modeled macrobenthic biomass compared to observed macrobenthic biomass.

879 Stations are colored according to their respective survey.