

Benthic habitat mapping of Langness Marine Nature Reserve in the Isle of Man

A dissertation submitted in partial fulfilment of the requirements
for the degree of Master of Science (MSc) in Marine Environmental Protection

Bangor University



PRIFYSGOL
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Submitted in October 2024.

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1 For submission to *Estuarine, Coastal and Shelf Science*

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3 **Benthic habitat mapping of Langness Marine Nature Reserve in the Isle of Man**

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11

12 **Abstract**

13 A survey on the benthic habitats and communities of the Langness Marine Nature Reserve in the Isle
14 of Man was carried out in 2024 Summer. The area was first designated as an inshore conservation
15 zone in 2016, as maerl beds and an eelgrass meadow were found, which are habitats of conservation
16 interest, followed by re-designation as Marine Nature Reserve (MNR) in 2018. As the MNR
17 is comparatively new to the others, baseline habitat mapping of it has yet to be done. This study
18 therefore aimed to map the benthic habitats and communities within the MNR, to provide baseline
19 information for future management and monitoring efforts. A benthic imagery survey was conducted
20 using a towed sledge, in which still images and footage of benthos were taken. The encountered
21 habitats were classified into biotopes using both a statistical approach and a classification system.
22 The survey results have identified 6 distinct benthic biotopes within the area of the MNR, with depth,
23 the type of substratum and hydrodynamic regime as primary abiotic factors shaping the communities.
24 No habitats of conservation interest have been identified. Although patches of maerl colonies have
25 been found in the northern regions of the MNR, they are mostly composed of dead nodules. No large
26 colonies of species of conservation interest or commercial species have been found.

27 **Keywords:**

28 Benthic Habitat; Mapping; Habitat classification; Marine Protected Area; Isle of Man

29 **1. Introduction**

30 Coastal benthic habitats are a key component of the marine ecosystem and are crucial to human
31 survival. Apart from being a hub that nurtures a diversity of species, the ecosystem services offered,
32 such as provision of food and abiotic resources, recycling of nutrients, and carbon sequestration, have
33 justified their importance (Barbier et al., 2011; Hall, 2002; Snelgrove, 1999). However, benthic
34 habitats are frequently experiencing disturbance, either by natural events, such as hydrodynamic
35 activities and biological interactions (e.g. predation), or by human activities (Gray and Elliott, 2009;
36 Hall and Harding, 1997; Jennings and Kaiser, 1998; Reise, 1978).

37 Regarding human disturbance to benthic habitats, fisheries activities are one the major players in
38 bringing about adverse impacts to the benthic ecosystems. Unregulated fisheries activities could lead
39 to drawbacks such as homogenisation of habitats, a decline in species biomass, abundance and
40 diversity, as well as a shrink in the size structure and production of a community (Beukers-Stewart
41 and Beukers-Stewart, 2009; Dayton et al., 1995; Jennings and Kaiser, 1998; Kaiser et al., 2006, 2002).
42 Fortunately, since the late 20th century, the environmental management bodies around the globe have

43 started to bring in an ecosystem-based approach (EBA) for fisheries management (FAO, 2003). Apart
44 from the focus on protecting the targeted species, the EBA also considered the entire ecosystem
45 in which the species live, aiming to preserve its structure, function, and diversity (Davies et al., 2021).
46 Marine protected areas (MPAs) are established as part of the implementation of the EBA management
47 (Gell and Roberts, 2003; Halpern and Warner, 2002; Roberts et al., 2001). With MPAs, the marine
48 habitats in the protected sea areas are avoided from detrimental activities (Renn et al., 2024; Sala and
49 Giakoumi, 2018), protecting the targeted species as well as its associated habitats and species
50 (Mesnildrey et al., 2013).

51 Situated in the middle of the Irish Sea, the Isle of Man (IoM) is a self-governing UK Crown
52 Dependency with a territorial sea covering a total area of approximately 4000 km² (Fig. 1) (DEFA,
53 2023). With such geographical privilege, it is not surprising that the fisheries sector has been a key
54 player in the Manx economy for centuries (DEFA, 2023; Duncan and Emmerson, 2018). The king
55 scallop, *Pecten maximus*, is the main fishery in Manx water, with the queen scallop, *Aequipecten*
56 *opercularis*, coming second (DEFA, 2015; Duncan and Emmerson, 2018). The Department of
57 Environment, Food and Agriculture (DEFA) Fisheries Directorate is a governmental body in the IoM
58 responsible for the management and protection of its territorial sea, fisheries and their supporting
59 ecosystems (Duncan and Emmerson, 2018). To date, 10 Marine Nature Reserves (MNRs) (a kind of
60 MPA) have been established within the Manx inshore territorial waters (0-3 nm), protecting a total
61 area of 430.75 km², which occupy 10.8% of the entire Manx territorial sea (Fig. 1). The MNRs are
62 established for different reasons, including conservation purposes, fisheries management and
63 experimental research, but it varies among MNRs (DEFA, 2024a, 2017).

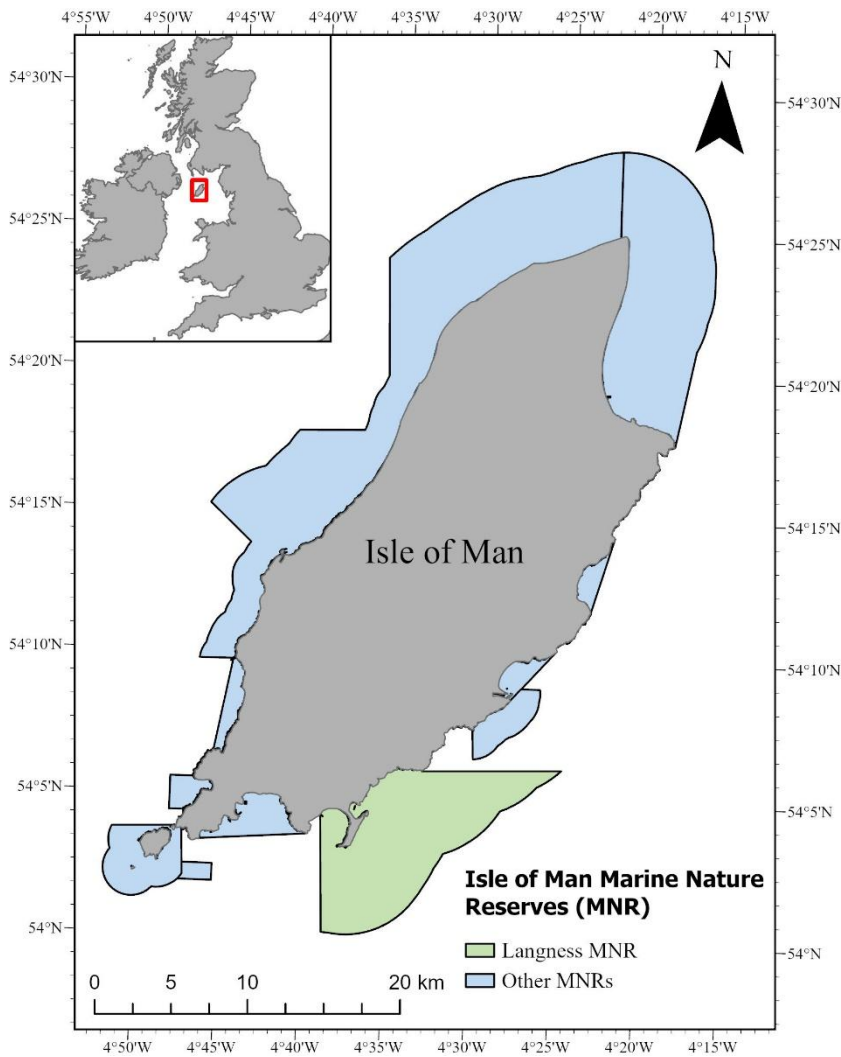


Fig. 1. A map of the Isle of Man, showing the area and location of the 10 Marine Nature Reserves (MNRs) designed in 2018. The location/area of the Langness MNR is shaded in green. The situation of the Isle of Man within the British Isles is shown on the smaller map at the top-left corner, indicated with a red box.

The coastal area of the Langness Peninsula was first in the spotlight in 2008. Thanks to its diverse rocky reef habitat, eelgrass meadow, and diverse sand/mud habitats, it has been identified as one of the candidate Marine Nature Reserve (MNR) sites in the ‘*Manx Marine Nature Reserve Project*’ (DEFA, 2010; Howe, 2018; Thomas et al., 2018). Followed by the implementation of the ‘*Inshore Marine Zoning Plan for the 0-3 Nautical Mile Area of the Isle of Man Territorial Sea*’ in 2016, the area has been designated as a ‘conservation zone’, with mobile fishing gear prohibited and habitats of conservation importance (for Langness, it is the maerl beds and an eelgrass meadow) protected (DEFA, 2017, 2016). On 1st September 2018, the area was re-designated as Langness Marine Nature Reserve (MNR) (Fig. 2) under the *Wildlife Act 1990*, becoming part of the 10 inshore MNRs network. General restrictions in MNR, such as bottom-towed fishing gear, are implemented in Langness MNR, but a specific management plan for the MNR is still under preparation as the MNR is relatively new (DEFA, 2018a, 2018b, 2017).

80 To protect the sensitive/vulnerable habitats and their associated species, it is necessary to know their
81 distribution in the first place. Benthic habitat mapping with the inclusion of biotope classification is
82 therefore being regarded as a useful tool for the management of marine-based resources (Fraschetti
83 et al., 2024; Harris and Baker, 2012). As mentioned, the Langness MNR is relatively new, the baseline
84 habitat mapping has therefore yet to be done. Though the general distribution of the benthic habitats
85 within the Manx territorial sea was known, thanks to the previous coarse-scale survey (Hinz et al.,
86 2008), habitats and species with restricted distribution are yet to be identified. Thus, surveys on a
87 finer scale are required to be done in areas of conservation interest, such as the Langness MNR.

88 Given the aforementioned need, this study therefore aims to map the benthic habitats and
89 communities within the MNR on a finer scale, determining the type, distribution and extent. The
90 completion of the mapping work would provide baseline information for future management and
91 monitoring efforts of the MNR, such as the assignment of conservation or fisheries management zone,
92 just as in other MNRs.

93 **2. Material and methods**

94 ***2.1 Study Site***

95 The study area, the Langness MNR, comprised a sector of the southeastern inshore waters of the IoM.
96 It extends eastward from Santon Head (54° 06.0000' N, 04° 33.0000' W) and southward from
97 Castletown Harbour (54° 04.3998' N, 04° 39.0000' W), out to 3 nautical miles from shore at an
98 astronomical high tide, which covered a total area of 88.67 km² (DEFA, 2024a, 2024b) (Fig. 2) .

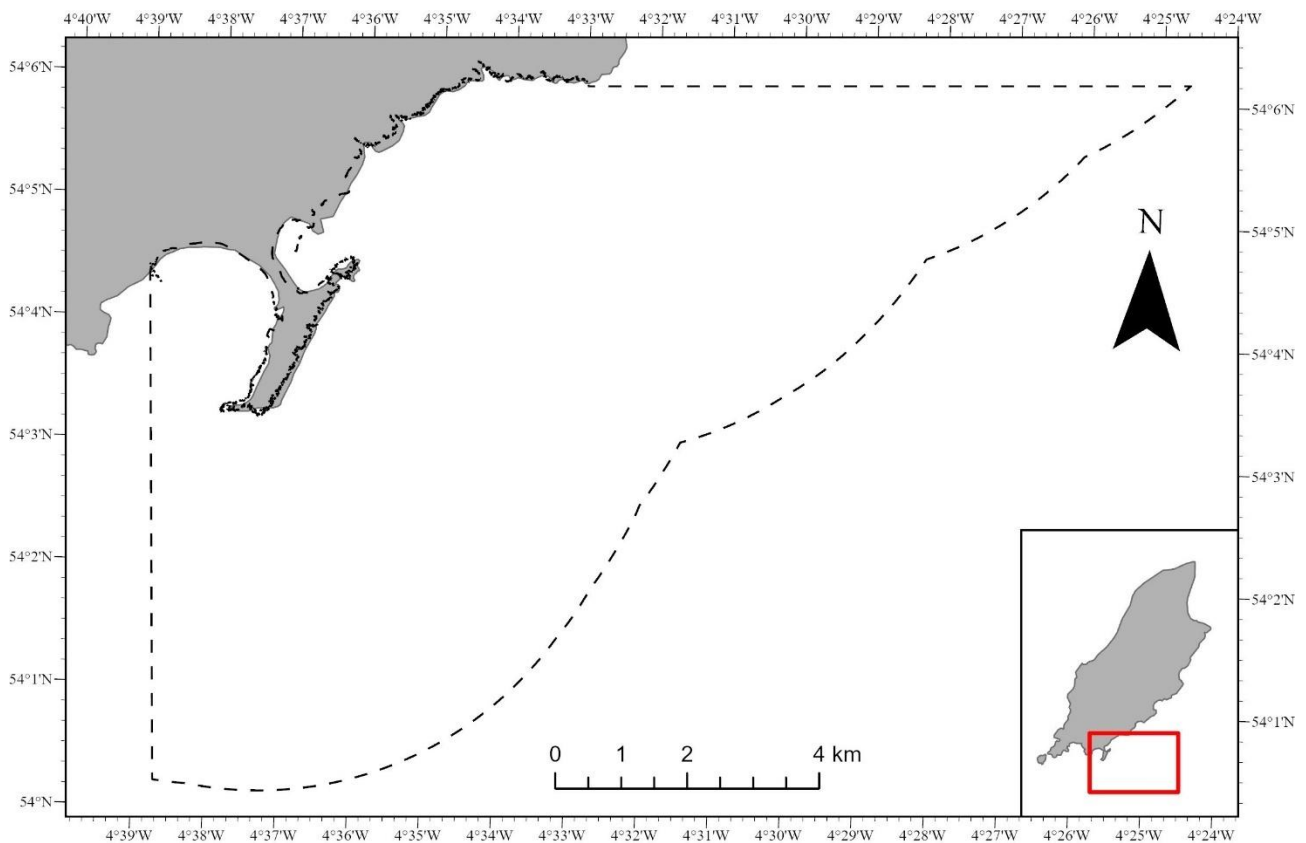


Fig. 2. A map showing the area of the Langness Marine Nature Reserve (MNR). Its situation within the Isle of Man's inshore waters is indicated on the small map at the bottom-right corner with a red box.

2.2 Habitat mapping

A grid system with a grid square size of 1km^2 each was used in this project to allocate sampling stations across the area of the MNR. The sampling stations were roughly 1km apart. At each station, a sledge with still image and video capability was deployed and towed along the seabed to record the benthos present. The sledge is composed of a metal frame on skids, with cameras and light fixed at the centre and oriented to face the seabed (Fig. 3). Two cameras in waterproof housing were used in this survey, namely a Canon EOS 400D, set to take a flashed photo every 10 seconds [Field of View (FOV) = $44 \times 29 \text{ cm}$], and a GoPro HERO3 for recording continuous video footage (FOV = $\sim 62 \times 35 \text{ cm}$).

The sampling work took place over 4 days in June 2024 by onboarding the IoM Government's Fisheries Patrol Vessel, the Barrule, and completed 49 transects within the MNR. At each sampling station, the sledge was towed along the seabed at approximately 1 knot for around 10 minutes, to obtain a 10-minute video clip and 60 still photos for each transect. Position data, including GPS coordinates, time and vessel speed, were automatically logged every 30 seconds throughout the survey journey and manually recorded at the beginning and end of each tow to allow geo-referencing

117 of the captured photographs. Water depth was also noted down at the start, middle and end of each
118 tow.



119
120 **Fig. 3.** Photograph of the equipment, the sledge, used for collection of the benthic imagery data. Both the cameras and
121 light were fixed on the raised unit at the centre.

122 **2.3 BRUVS sampling**

123 Baited remote underwater video systems (BRUVS) were used for the identification of mobile species
124 present within the MNR. Each of the BRUVS is composed of a rectangular metal frame with three
125 GoPro cameras (either HERO3 or HERO3+, settings: 1080p, 60/50/30 fps, Wide FOV) attached on
126 top, viewing at three different angles (top-down, top-45°left, and top-45°right) (Fig. 4). Bait (herring,
127 *Clupea harengus*) packed in a mesh bag was fixed at the bottom of the frame (in the centre), to
128 produce a scent for attracting mobile species. A total of 20 BRUVS were deployed, and each was left
129 on the seabed for at least an hour. At each deployment site, GPS coordinates, water depth, as well as
130 start and end time of deployment were recorded.

131



Fig. 4. Photograph of the BRUVS, equipment used for collecting mobile species data. Three GoPro cameras at different viewing angles were fixed on top of the rectangular metal frame.

2.4 Analysis of images and videos

2.4.1 Visibility and Quality Assessment

To ensure data reliability, the still images and video footage captured were assessed for visibility and quality before any further analysis. The assessment was done by applying a standardised scoring system adapted from Hannah and Blume (2012) (Table 1). Based on the defined criteria, the images and videos were scored from 0 to 3; those that scored 0 on either visibility or quality were omitted from further analysis.

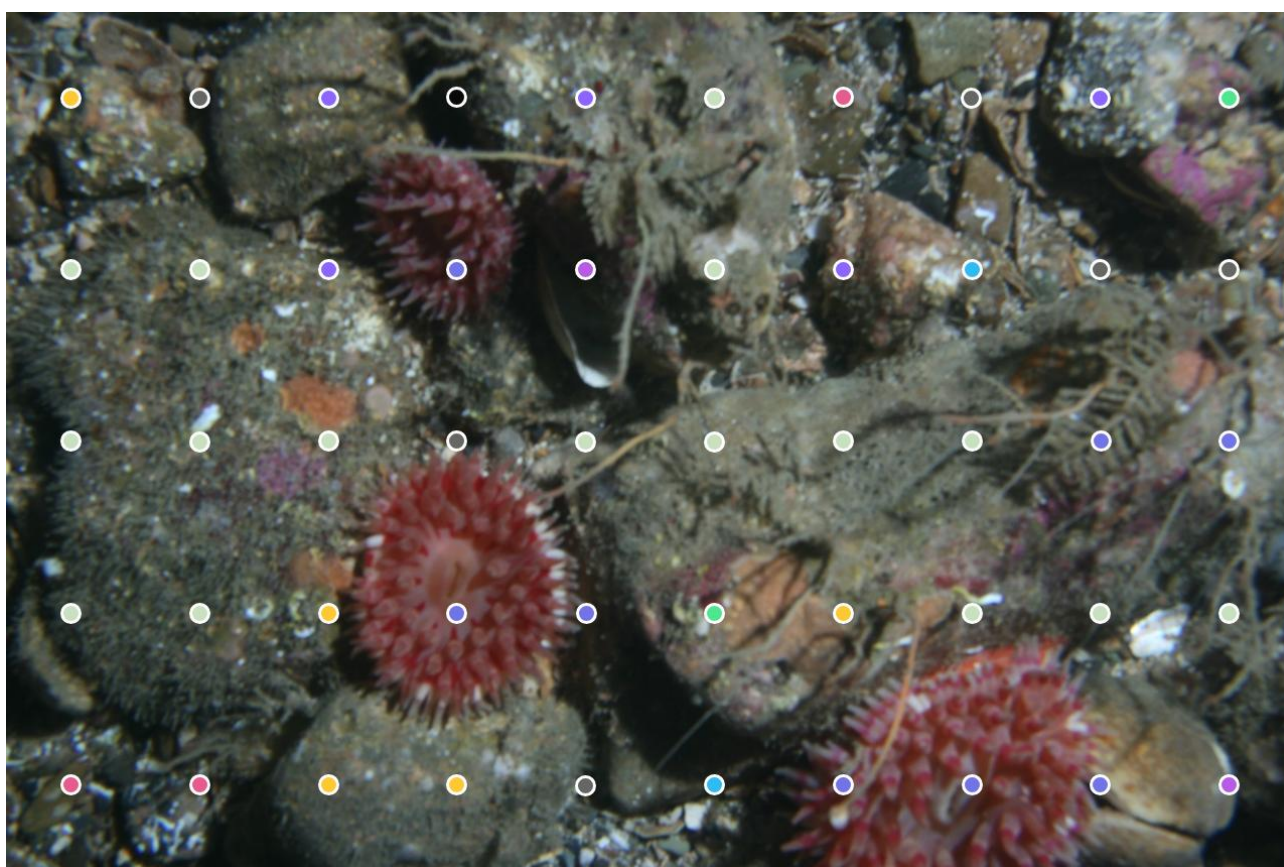
Table 1

The scoring system applied for the visibility and quality assessment of the images and videos captured. (Hannah and Blume, 2012).

Score	Visibility	Quality
0	The field of view is completely obscured by close-up species or suspended sediment.	The image/video is completely blurred or has major problems with lightning/viewing angle.
1	The field of view is greatly obscured (>50%) by close-up species or suspended sediment.	The image/video is greatly blurred (>50%) or has some problems with lightning/viewing angle.
2	The field of view is partially obscured (<50%) by close-up species or suspended sediment.	The image/video is greatly blurred (<50%) or has minor problems with lightning/viewing angle.
3	The field of view is clear/of insignificant obstruction.	The image/video is clear/of insignificant quality issues.

146 2.4.2 Analysis of images from the towed sledge

147 The still images were analysed using the image annotation function of a web-based application,
148 BIIGLE. (Langenkämper et al., 2017). Three types of data were extracted from the analysed images,
149 including 1) the percentage cover of the fauna, flora and physical benthic substrate by applying the
150 point sampling technique (Fig. 5) (Ninio et al., 2003; Ryan, 2004; Wakeford et al., 2008); 2) the
151 presence/absence of faunal and floral taxa; and 3) the abundance of countable epifauna. 5 images
152 were analysed per towed transect, and each was captured at an interval of 120 seconds apart, achieving
153 an even distribution of analysed images throughout the 10-minute tow. In case of poor visibility or
154 quality (scored 0), the image captured 10 seconds before or after was used as a replacement for the
155 image analysis.



156
157 **Fig. 5.** An example image showing the (regular) point sampling technique applied for obtaining the relative percentage
158 cover data. The fauna, flora, or physical benthic substrate found at each of the 50 points were identified, with each point
159 taken as a 2% cover. For this example, the Dahlia anemone, *Urticina felina*, has been identified at 6 points, thus covering
160 12% of this image.

161 All taxa encountered were identified to the lowest possible taxonomical level, using several photo
162 identification guides, namely Moen and Svensen (2004), Kay and Dipper (2009), Bunker et al. (2012),
163 Porter (2012), Sterry and Cleave (2012), Wood (2013), and Wood (2018). Broad descriptive
164 categories were used in the case of problematic taxa identification. Physical benthic substrates were

determined visually and described into board categories based on a simplified version of the Wentworth scale (Wentworth, 1922) (Table 2). The described substrates include boulders, cobbles, dead maerl, dead shells, shell fragments, pebbles, and granules.

Table 2

A simplified version of the Wentworth scale, used for physical benthic substrate determination (Wentworth, 1922).

Classification	Particle size (diameter)
Boulders	>256 mm
Cobbles	64 – 256 mm
Pebbles	4 – 64 mm
Granules	2 – 4 mm

2.4.3 Analysis of videos from the BRUVS

The videos were also analysed with BIIGLE (Langenkämper et al., 2017), using its video annotation function. The duration of the analysis for each video was standardised as 60 minutes. MaxN, a conservative, commonly used approach for the estimation of the total number of individuals from a species (Cappo, 2010; Whitmarsh et al., 2017), was applied to estimate the abundance of mobile species at each BRUVS station. For any species identified in a video, its MaxN is defined as the maximum number of individuals observed in a single frame (Ebner et al., 2009; Loiseau et al., 2016). Since each BRUVS unit has three analysed videos, for an identified mobile species, its abundance estimate at a BRUVS station was determined by referring to its maximum MaxN value obtained from the video analyses.

2.5 Data Analysis

All multivariate analyses were performed using functions in PRIMER 7 (details in the following sections), whereas the univariate analyses were done using the packages installed in RStudio. All presented maps were prepared in ArcGIS Pro.

2.5.1 Biotope classification

A biotope was assigned to each of the benthic habitats/communities identified from the previous analyses of the geo-referenced still images. A biotope was assigned through two methods: 1) using statistical method (multivariate) in PRIMER 7; and 2) application of the habitat classification system, ‘The Marine Habitat Classification for Britain and Ireland’ (MHCBI) (Version 22.04) (JNCC, 2022).

190 2.5.1.1 Biotope classification in PRIMER 7

191 A CLUSTER analysis (hierarchical agglomerative method) with the inclusion of similarity profile
 192 (SIMPROF) tests (for defining significant clusters, $p < 0.05$) was performed on the similarity matrix
 193 of the square-root transformed percentage cover data (Clarke et al., 2014, 2008; Gordon, 1987). The
 194 sampling stations were grouped according to their similarity in benthic community assemblages,
 195 forming individual ‘clusters’, and each ‘cluster’ was defined as a distinctive biotope.

196 With the defined biotopes as a factor, ANOSIMs (Analysis of similarity) were run on the similarity
 197 matrixes of the square-root transformed percentage cover data, taxa presence/absence data, and
 198 countable epifauna abundance data to determine if the community assemblages of the defined
 199 biotopes are significantly different from each other (Clarke and Green, 1988). As a follow-up analysis,
 200 SIMPERs (Similarity percentage) were carried out on the aforementioned square-root transformed
 201 data, to identify the discriminating taxon that contributed to the distinctiveness of the biotopes
 202 (Clarke, 1993). A non-metric multidimensional scaling (nMDS) plot was eventually used to visualise
 203 the differences between biotopes (Kruskal and Wish, 1978).

204 2.5.1.2 MHCBI biotope classification

205 The defined biotopes in the MHCBI system are categorized in a 6-level hierarchical structure (Table
 206 3) (JNCC, 2022). Biotopes in the first 3 levels are defined based on physical parameters, including
 207 water depth, substrate type, wave energy and current energy. Further down the hierarchy,
 208 the biological community information is also included in defining the biotopes.

209 **Table 3**

210 An example of the application MHCBI system for biotope assignment.

Level	Category	Example	Code
Level 1	Environment	Marine	-
Level 2	Broad habitat type	Sublittoral sediment	SS
Level 3	Habitat complex	Sublittoral macrophyte-dominated communities on sediments	SS.SMp
Level 4	Biotope complex	Kelp and seaweed communities on sublittoral sediment	SS.SMp.KSwSS
Level 5	Biotope	Saccharina latissima and red seaweeds on infralittoral sediments	SS.SMp.KSwSS.SlatR
Level 6	Sub-biotope	Saccharina latissima and robust red algae on infralittoral gravel and pebbles	SS.SMp.KSwSS.SlatR.Gv

212 A biotope in the MHCBI system was assigned to each sampling station (analysed still image), based
 213 on both the measured or obtained abiotic and biotic data. For abiotic parameters, water depth was

214 measured during sampling, substrate type was determined during the analysis of still images, and
215 wave and current energy were obtained from the EMODnet Seabed Habitats portal (EMODnet, 2022).
216 As for the biotic component, both the still images and footage captured by the towed sledge within
217 a similar time frame (± 30 seconds) were used for determining the benthic taxon present. All biotopes
218 were assigned to the lowest possible level.

219 ANOSIMs followed by SIMPERs were done in a similar approach as mentioned in the previous
220 section but the factor was changed to the identified MHCBI biotopes. A nMDS plot was also used to
221 visualise the differences. In addition, the species richness (number of species, S), summed abundance
222 of countable epifauna and summed algal percentage cover of each sampling station were calculated.
223 The means were then tested for differences among biotopes using ANOVAs (Analysis of variance),
224 or Kruskal–Wallis tests, in case of failure to fulfil parametric assumptions of ANOVA (Rohlf, 2011).

225 *2.5.2 Creation of habitat maps*

226 Maps of the estimated extent of the identified biotopes (both classification methods) within the
227 Langness MNR boundary were created through Euclidean allocation analysis of point samples in
228 ArcGIS Pro. The biotope assigned to each sampling station was extrapolated to the cells nearby,
229 eventually forming a map of biotopes covering the area of the MNR.

230 *2.5.3 Environmental variables*

231 The relationship between environmental variables and benthic community assemblage was assessed
232 using the BEST function (BIO-ENV routine) in PRIMER 7 (Clarke and Gorley, 2015). The
233 considered variables include water depth, the coverage of different substrate types, algal coverage,
234 wave energy, and current energy. Through the ‘BEST’ analysis, the variable(s) that ‘best’ correlate
235 with the community assemblage were identified.

236 *2.5.4 Data from the BRUVS*

237 An ANOSIM with MHCBI biotopes as a factor was run on the similarity matrix of the square-root
238 transformed mobile species abundance data to test for significant differences in community
239 assemblage between biotopes. The MHCBI biotope associated with each BRUVS station was
240 determined based on the created biotope map. A SIMPER was also carried out on the aforementioned
241 square-root transformed data as a follow-up analysis to identify the discriminating mobile species
242 that contributed to the distinctiveness of the biotopes. An nMDS plot was eventually applied for
243 visualization of the differences in community assemblage between biotopes. In addition, the species

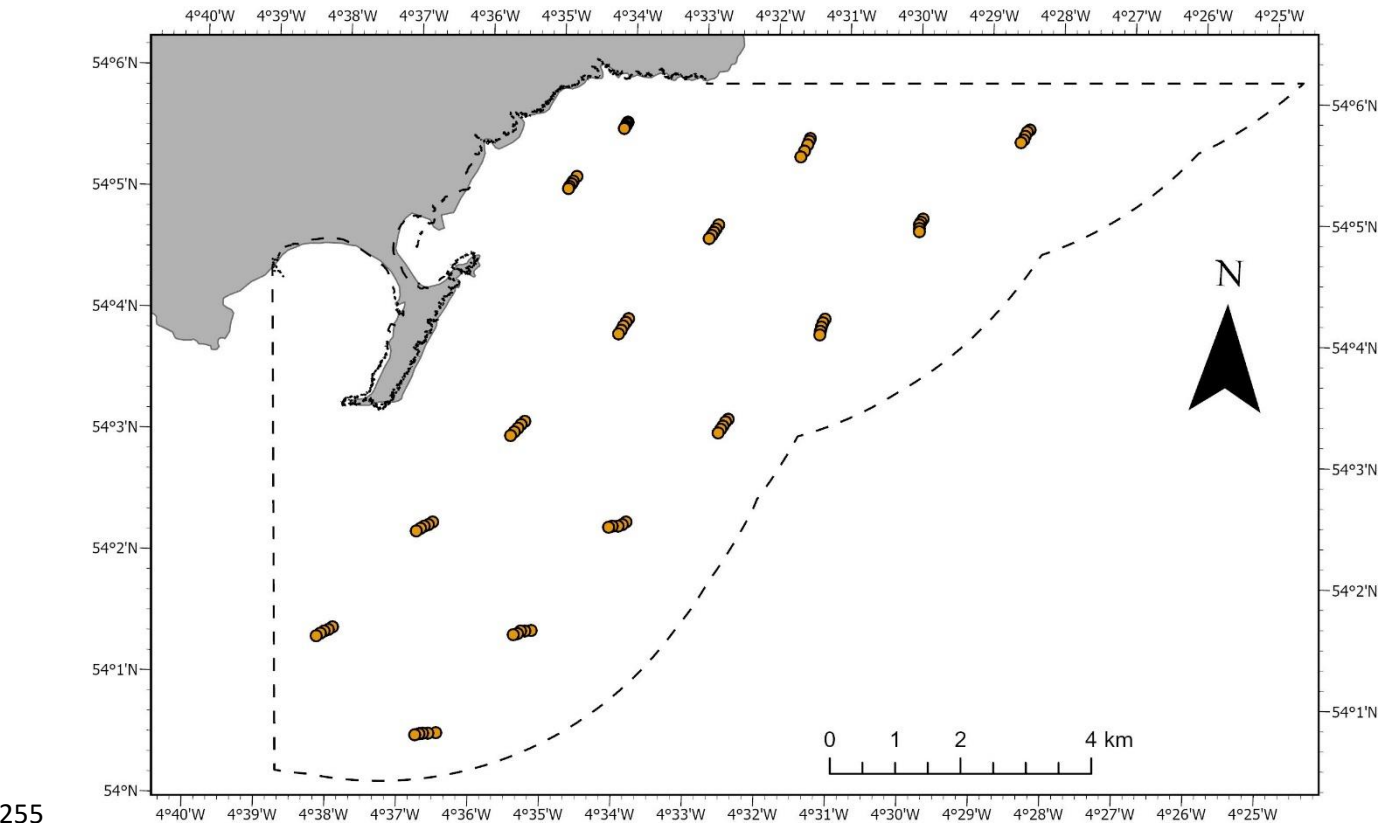
244 richness (number of species, S) of each sampling station was calculated and the mean was then tested
245 for differences among biotopes using the Kruskal–Wallis test.

246 A ‘BEST’ analysis was also done to identify which of the three environmental variable(s), namely
247 water depth, wave energy, and current energy, ‘best’ correlate with the structure of community
248 composition.

249 **3. Results**

250 **3.1 Towed sledge sampling**

251 Due to time constrain, 75 images captured from 15 towed transects (Fig. 6) were selected for analysis
252 to obtain percentage cover and species data. The images were of acceptable quality only, most had
253 minor focus issues due to gear malfunctioning. Only 4% of the images scored 3 in both visibility and
254 quality while the majority (~90%) scored 2 in quality and 3 in visibility.



255 **Fig. 6.** A map showing the location of the 75 analysed still images taken along the 15 towed transects in Langness MNR.

257 **3.1.1 Identified taxa**

258 A total of 82 taxa were identified from the analysed still images (supplementary data, Table A1),
259 including 18 algae (22%), 16 cnidarians (20%), 15 molluscs (18%), 8 arthropods (10%), 6 bryozoans
260 (7%), 6 sponges (7%), 4 echinoderms (5%), 3 algae/hydroids/bryozoans mix (4%), 2 annelids (2%),

261 2 ascidians (2%), and 2 teleosts (2%). More than 80% of the taxa were identified to at least family
262 level, 73% were identified to species or genus level, and the remaining taxa were categorised into
263 broad descriptive categories.

264 Species richness in individual images ranged from 3 to 18 taxa, with an average of 10 taxa per image.
265 7 taxa were commonly encountered, each found in more than 40% of the analysed still images. These
266 taxa include brownish-green encrusting bryozoan, Serpulidae spp., Corallinaceae spp., muddy surface
267 turf, *Parasmittina trispinosa*, *Glycymeris glycymeris*, and *Steromphala cineraria*. If only the
268 countable epifauna is considered, the top five most common taxa include *Glycymeris glycymeris*
269 (65%), *Steromphala cineraria* (48%), *Galathea intermedia* (32%), *Gibbula magus* (29%), and
270 *Calliostoma zizyphinum* (19%).

271 Taxa with conservation and commercial importance were identified in Langness MNR, namely maerl
272 and queen scallop, *Aequipecten opercularis*. The maerl has a restricted distribution, it was only
273 present in the northern region of the MNR, both inshore and offshore. Live maerl was present in 9%
274 of the analysed still images, with an average cover of $0.56 \pm 0.18\%$, while dead maerl was sighted in
275 40% of the images and averaged at $4.70 \pm 0.54\%$ cover. As for the queen scallop, only a single
276 individual was recorded from the analysed still images, in which the sampling station was situated in
277 the centre of the MNR.

278 3.1.2 Environmental variables

279 The water depth (below CD) of the sampling stations ranged from 19.1m to 38.7m, with an average
280 of $32.0 \pm 0.57\text{m}$ (Fig. 7a). The substratum across the sampling area was heterogenous, with granules,
281 pebbles, cobbles, boulders, shells, and coarse sediments. Most of the sampling areas have substrates
282 mainly composed of pebbles and cobbles, with some degree of shells and coarse sediments. The
283 strength of the current and wave in these areas was generally weak to moderate. In areas with
284 substratum consisting of boulders, e.g. the coastal waters in front of Langness Peninsula and Port
285 Grenaugh, the current or wave strength was comparatively stronger (Fig. 7b).

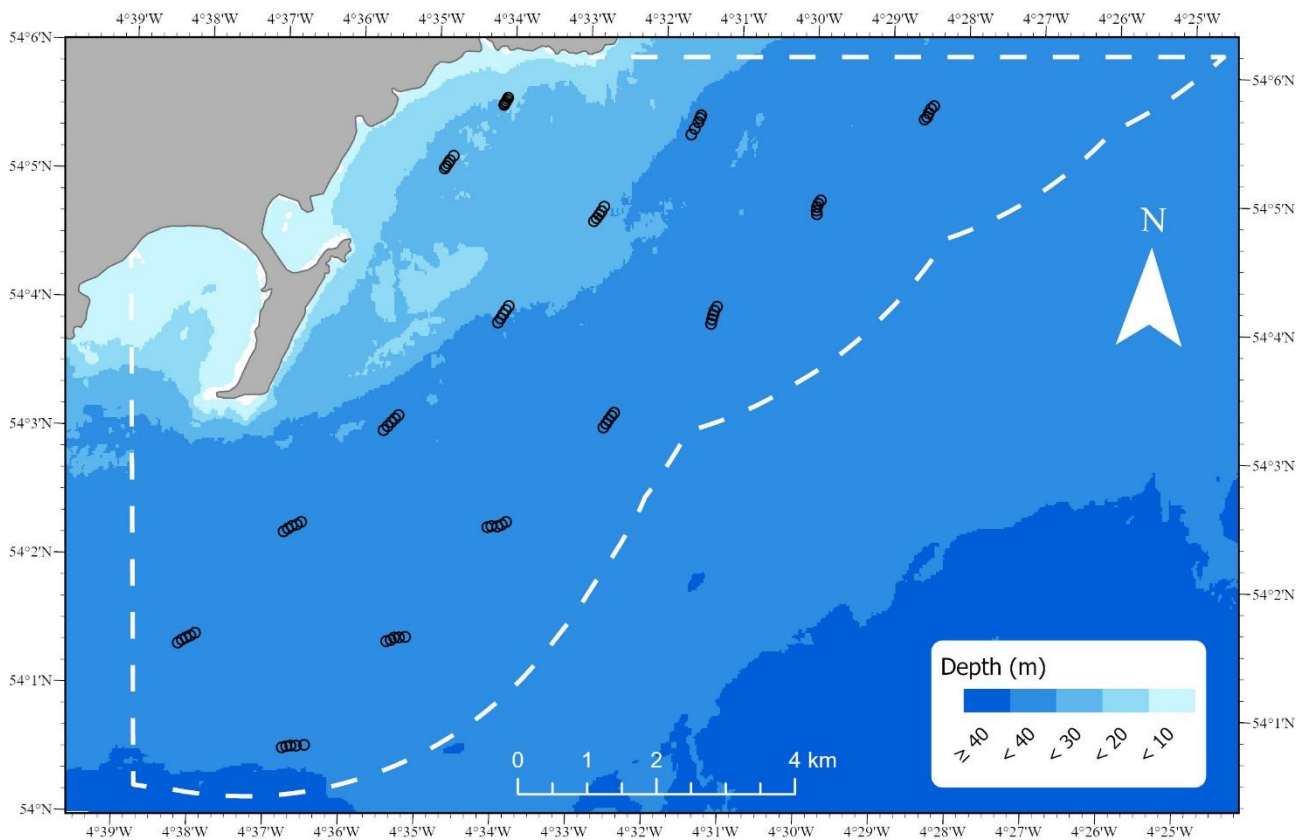


Fig. 7a. A map showing the bathymetry (below CD) across the areas of the Langness MNR (EDINA, 2020), with locations of towed sledge sampling stations (only those with image analysis done) overlaid on top.

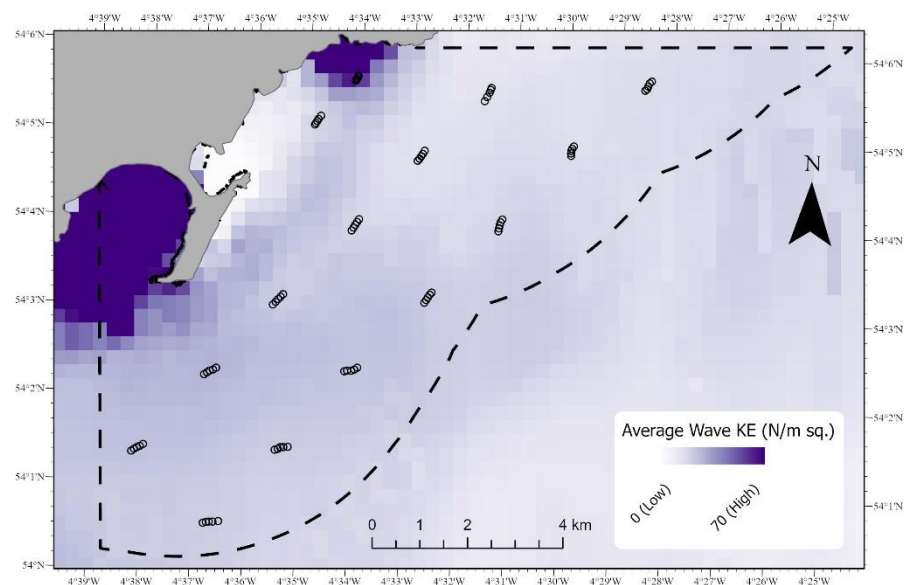
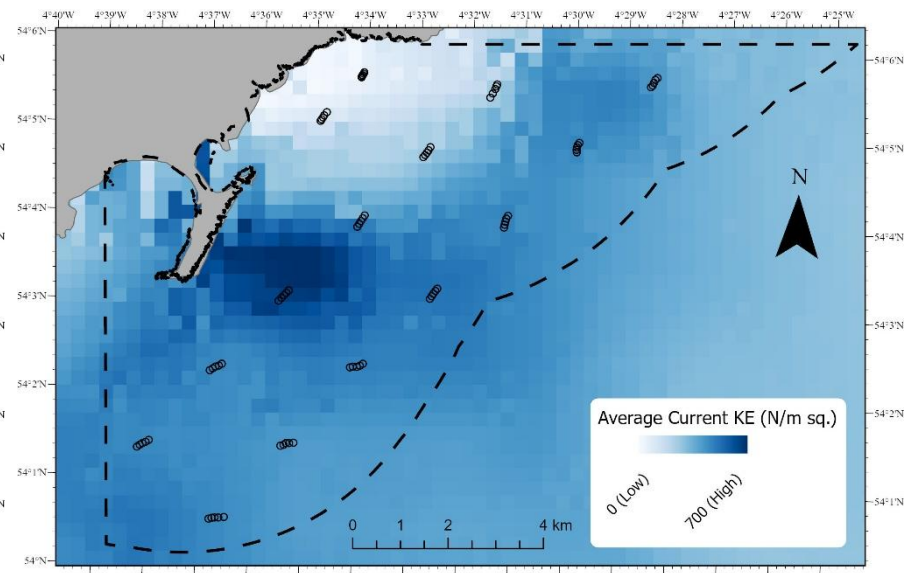
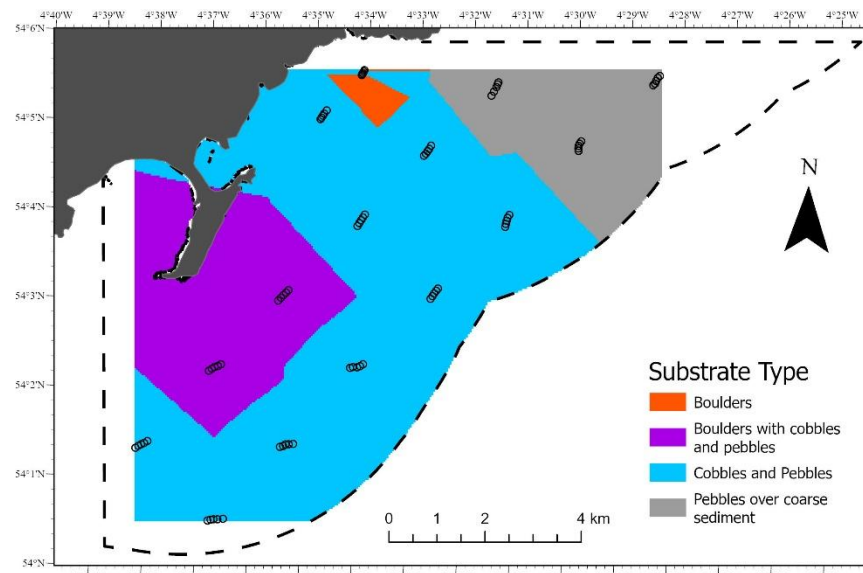
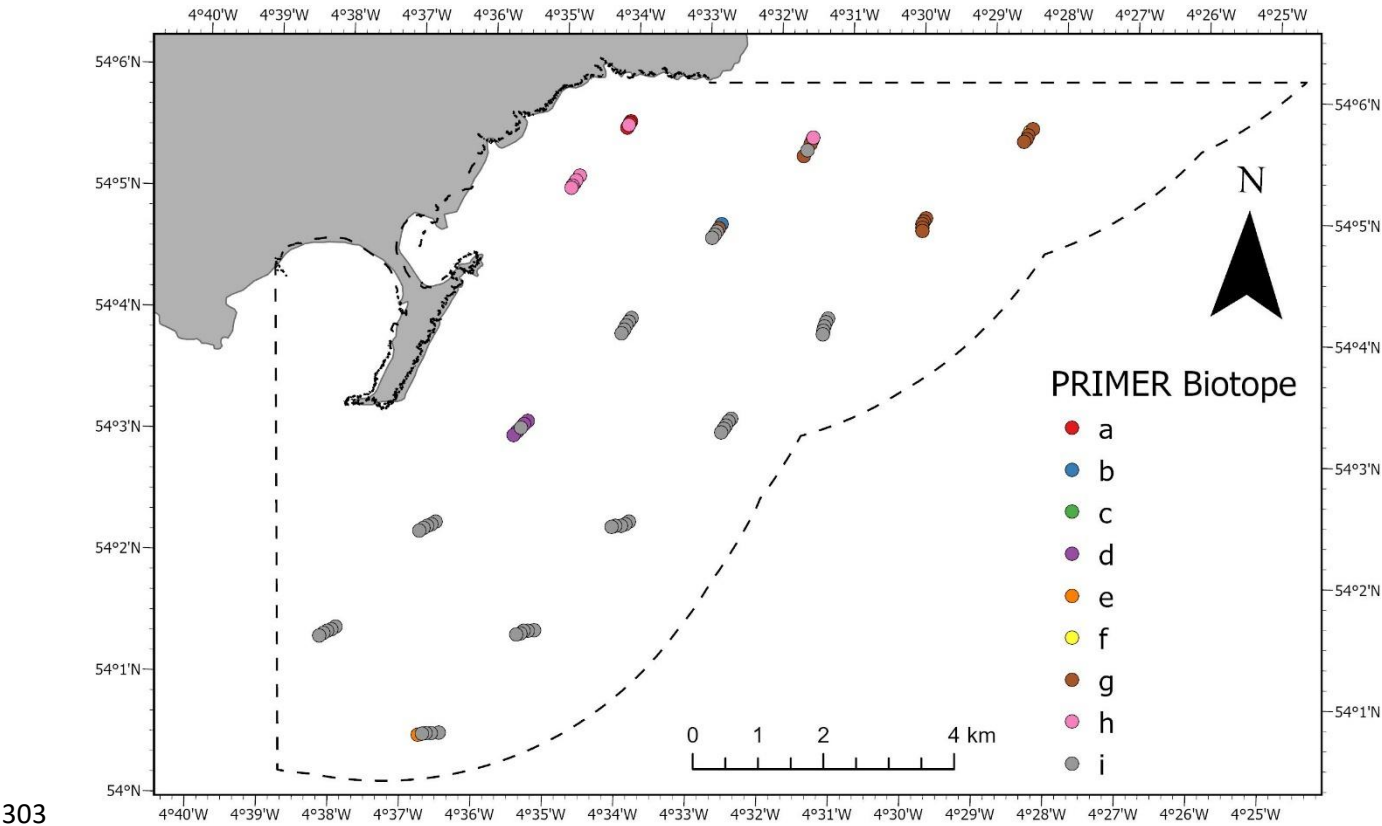


Fig. 7b. A map showing the substrate type, current energy (N/m^2) (EMODnet, 2022), and wave energy (N/m^2) (EMODnet, 2022) across the areas of the Langness MNR, with locations of towed sledge sampling stations (only those with image analysis done) overlaid on top.

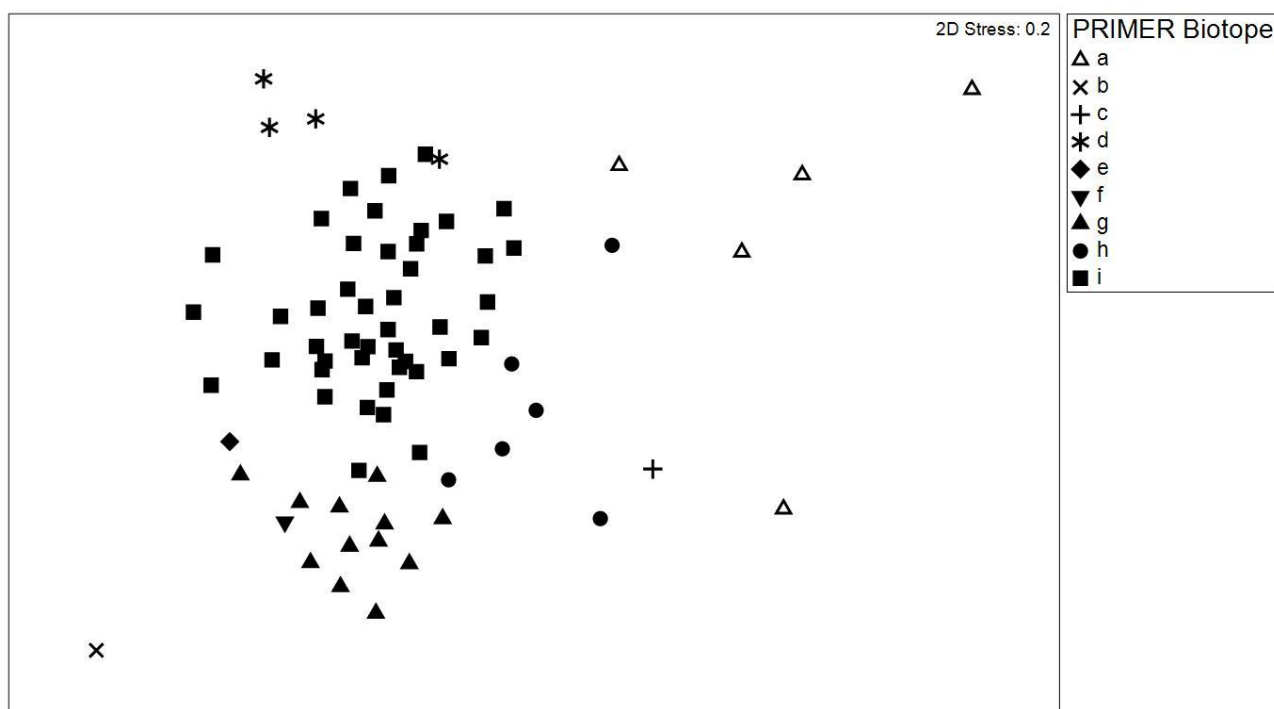
293 3.1.3 PRIMER biotopes

294 Based on the percentage cover data, the PRIMER statistical approach has identified 9 clusters of
295 benthic communities (supplementary data, Fig. A1) in Langness MNR. Each ‘cluster’ is defined as a
296 distinct biotope, named with an alphabet (a-i). The associated PRIMER biotope of each sampling
297 point (analysed still image) and their distribution within the Langness MNR boundary is shown on
298 the map in Figure 8.

299 ANOSIMs have confirmed the significant difference in community structure among biotopes, in
300 terms of both the percentage cover data ($R = 0.724$, $p = 0.001$) (Fig. 9) and taxa presence/absence
301 data ($R = 0.431$, $p = 0.001$). The summary of each of the identified biotopes is provided in Table 4
302 with the inclusion of the SIMPER analysis results.



303
304 **Fig. 8.** A map showing the associated PRIMER biotope of each sampling point (analysed still image) and their distribution
305 within the Langness MNR boundary.



306

307

308

309

Fig. 9. An nMDS plot showing the relationship between communities of each sample. The plot was created using percentage cover data, with each sample assigned a symbol according to their associated biotope. Points in close proximity indicate high similarity with each other while points further apart indicate low similarity.

310 **Table 4**

311 A summary of the benthic biotopes identified in Langness MNR using the PRIMER statistical approach. The table below
 312 includes the total number of sampling sites (still images used) for each biotope, the average similarity in percentage cover
 313 data among sampling sites within each biotope (full SIMPER analysis result of the percentage cover data is available in
 314 the supplementary data, Table A2), and the discriminating taxa for each biotope (the taxa that contributed to both within
 315 biotope similarity and between biotopes dissimilarity in terms of community composition), based on the taxa
 316 presence/absence data.

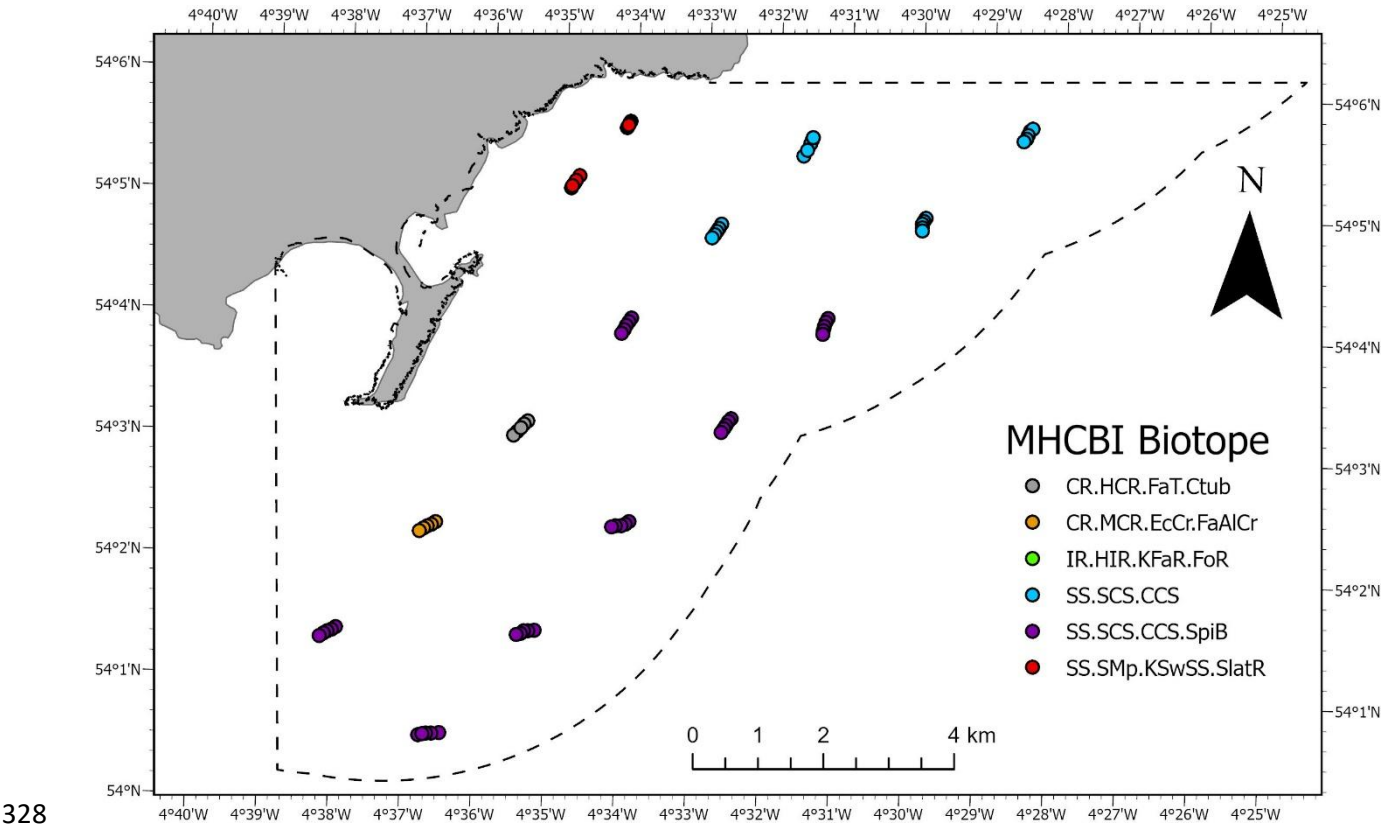
PRIMER Biotope	No. of sampling sites (still images used)	Average similarity	Discriminating taxa
a	5	43%	Branching red seaweed, <i>Clavelina lepadiformis</i> , <i>Delesseria sanguinea</i> , Flat brown seaweed, Mixed turf of algae with bryozoan and/or hydroid, <i>Parasmittina trispinosa</i> , <i>Phyllophora</i> spp., <i>Plocamium</i> spp., <i>Schizomavella</i> spp., <i>Steromphala cineraria</i> , Thin flat red seaweed
b	1	100%	<i>Calliostoma zizyphinum</i> , <i>Galathea intermedia</i> , <i>Halecium</i> spp., <i>Hydrallmania falcata</i> , <i>Nemertesia antennina</i> , <i>Pagurus bernhardus</i> , <i>Pomatoschistus</i> spp.
c	1	100%	Serpulidae spp.
d	4	61%	Anomiidae spp., <i>Corynactis viridis</i> , <i>Crisia</i> spp., <i>Schizomavella</i> spp., <i>Steromphala cineraria</i> , <i>Tubularia indivisa</i>
e	1	100%	Small pinkish crab, <i>Synarachnactis lloydii</i>
f	1	100%	<i>Steromphala cineraria</i>
g	12	69%	<i>Gibbula magus</i> , <i>Glycymeris glycymeris</i> , <i>Steromphala cineraria</i>
h	6	61%	Brownish green encrusting bryozoan, <i>Glycymeris glycymeris</i> , <i>Hapalidiaceae</i> spp., <i>Nemertesia antennina</i> , Serpulidae spp.
i	44	60%	<i>Galathea intermedia</i> , <i>Glycymeris glycymeris</i> , Mixed turf of bryozoan and hydroid, Muddy surface turf, <i>Steromphala cineraria</i>

317

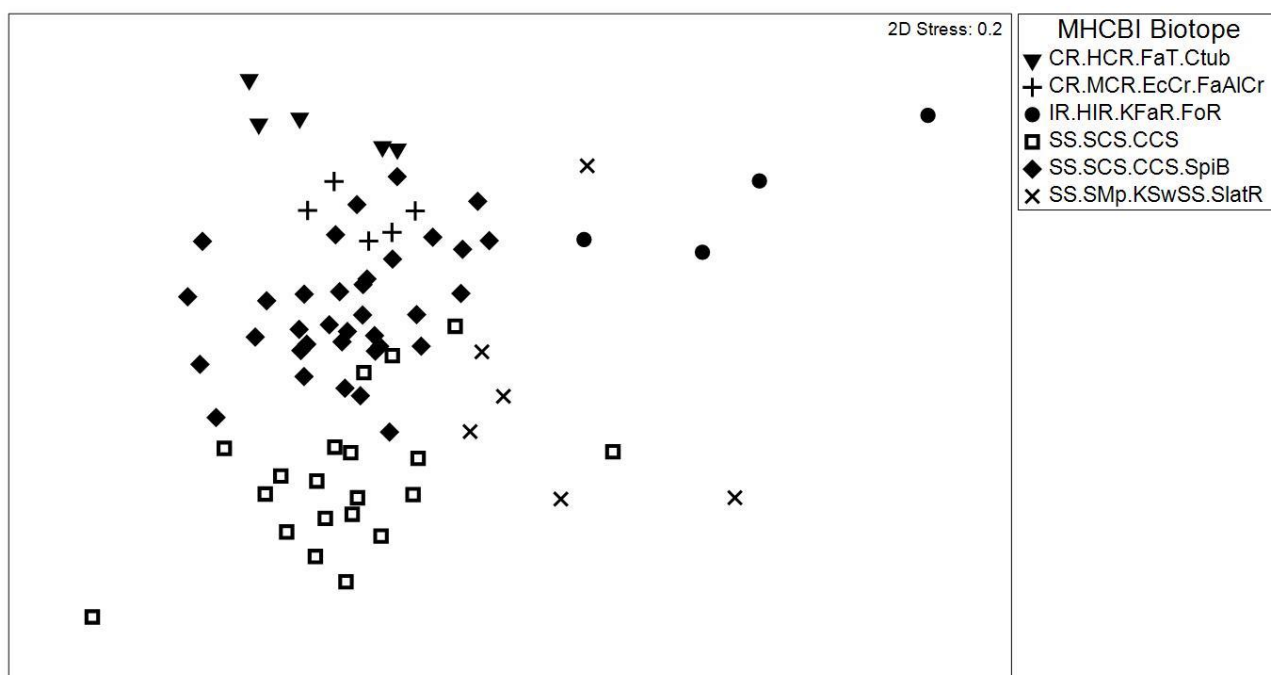
318 3.1.4 MHCBI biotopes

319 Based on the percentage cover data, the MHCBI classification system approach has identified 6
 320 distinct biotopes in Langness MNR. The associated MHCBI biotope of each sampling point (analysed
 321 still image) and its distribution within the Langness MNR boundary are shown on the map in Figure
 322 10.

323 ANOSIMs have confirmed the significant difference in community structure among biotopes, in
 324 terms of the percentage cover data ($R = 0.585$, $p = 0.001$) (Fig. 11), taxa presence/absence data ($R =$
 325 0.447 , $p = 0.001$), and countable epifauna abundance data ($R = 0.505$, $p = 0.001$). The summary of
 326 each of the identified biotopes is provided in Table 5 with the inclusion of the SIMPER analysis
 327 results.



328 **Fig. 10.** A map showing the associated MHCBI biotope of each sampling point (analysed still image) and their distribution
 329 within the Langness MNR boundary.
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Fig. 11. An nMDS plot showing the relationship between communities of each sample. The plot was created using percentage cover data, with each sample assigned a symbol according to their associated biotope. Points in close proximity indicate high similarity with each other while points further apart indicate low similarity.

335 **Table 4**

336 A summary of the benthic biotopes identified in Langness MNR using the MHCBI classification system approach. The
 337 table below includes the total number of sampling sites (still images used) for each biotope, the average similarity in
 338 percentage cover data among sampling sites within each biotope (full SIMPER analysis result of the percentage cover
 339 data is available in the supplementary data, Table A3), and the discriminating taxa for each biotope (the taxa that
 340 contributed to both within biotope similarity and between biotopes dissimilarity in terms of community composition),
 341 based on the taxa presence/absence data.

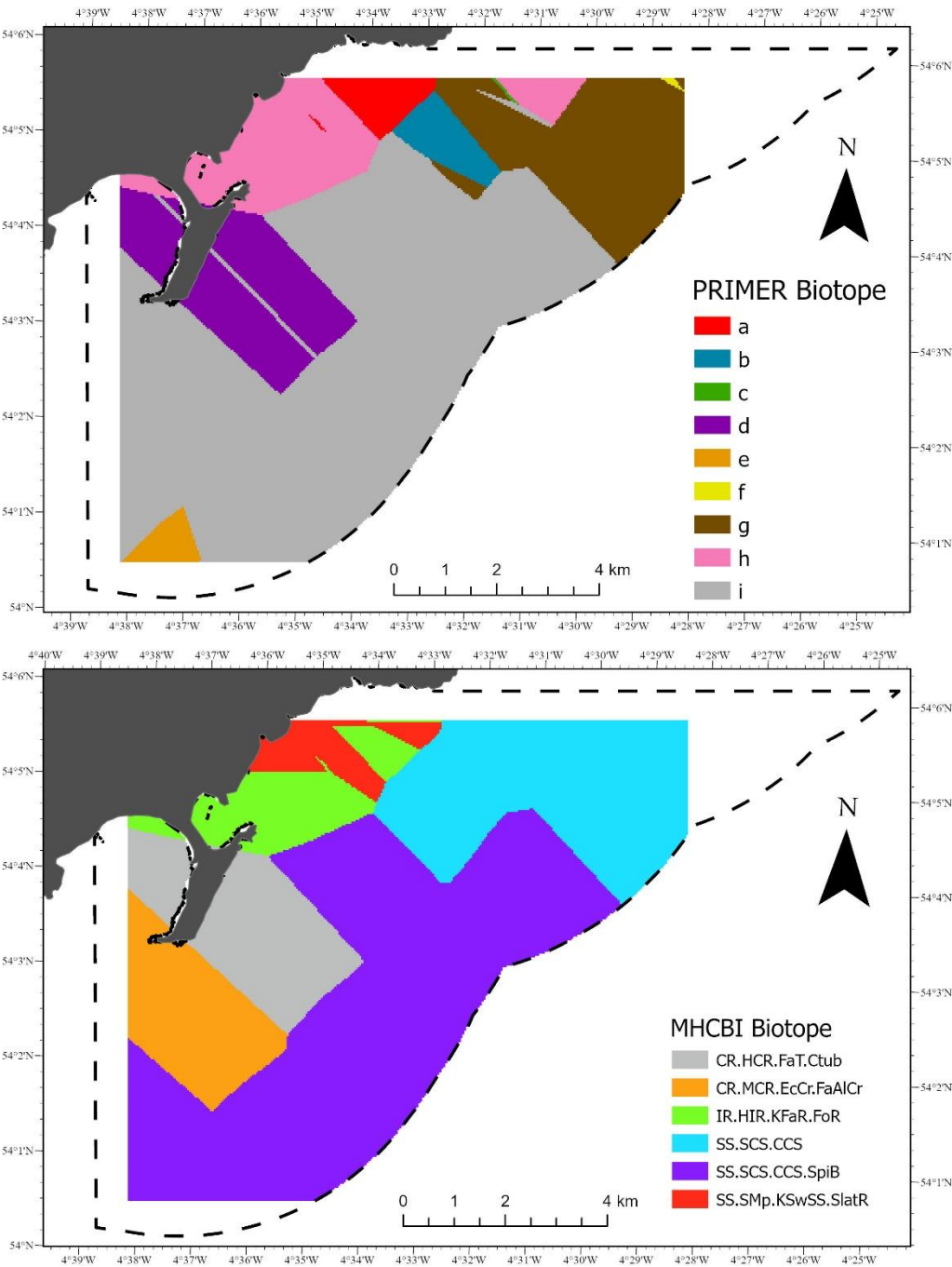
MHCBI Biotope	Biotope no.	No. of sampling sites (still images used)	Average similarity	Discriminating taxa
CR.HCR.FaT.Ctub Tubularia indivisa on tide-swept circalittoral rock	1	5	59%	Anomiidae spp., <i>Balanus</i> spp., <i>Corynactis viridis</i> , <i>Crisia</i> spp., Mixed turf of bryozoan and hydroid, <i>Schizomavella</i> spp., <i>Steromphala cineraria</i> , <i>Tubularia indivisa</i>
CR.MCR.EcCr.FaAlCr Faunal and algal crusts on exposed to moderately wave-exposed circalittoral rock	2	5	69%	<i>Balanus</i> spp., <i>Calliostoma zizyphinum</i> , <i>Corynactis viridis</i> , <i>Tubularia indivisa</i>
IR.HIR.KFaR.FoR Foliose red seaweeds on exposed lower infralittoral rock	3	4	48%	<i>Delesseria sanguinea</i> , <i>Dictyota dichotoma</i> , Mixed turf of algae with bryozoan and/or hydroid, <i>Nemertesia antennina</i> , <i>Parasmittina trispinosa</i> , <i>Phyllophora</i> spp., <i>Schizomavella</i> spp.
SS.SCS.CCS Circalittoral coarse sediment	4	20	59%	<i>Glycymeris glycymeris</i> , <i>Parasmittina trispinosa</i> , <i>Steromphala cineraria</i>
SS.SCS.CCS.SpiB Spirobranchus triqueter with barnacles and bryozoan crusts on unstable circalittoral cobbles and pebbles	5	35	60%	Corallinaceae spp., <i>Galathea intermedia</i> , <i>Glycymeris glycymeris</i> , Mixed turf of bryozoan and hydroid, Muddy surface turf, <i>Parasmittina trispinosa</i> , <i>Steromphala cineraria</i>
SS.SMp.KSwSS.SlatR Saccharina latissima and red seaweeds on infralittoral sediments	6	6	52%	Brownish green encrusting bryozoan, <i>Calliostoma zizyphinum</i> , <i>Gibbula magus</i> , Hapalidiaceae spp., Mixed turf of algae with bryozoan and/or hydroid, Muddy surface turf, <i>Nemertesia antennina</i> , <i>Plocamium</i> spp., Serpulidae spp., <i>Steromphala cineraria</i>

342

343 3.1.5 Summary of biotope classifications

344 In summary, the two biotope classification approaches have identified 9 (PRIMER) and 6 (MHCBI)
345 biotopes respectively. The associated biotope of each sampling point was extrapolated to its nearby
346 areas, creating maps showing the estimated distribution of biotopes within the Langness MNR
347 boundary (Fig. 12). Considering the suitability of the approach to the data collected, the MHCBI
348 classification system has been adopted for the remaining analysis of the data (to be further explained
349 in discussion).

350

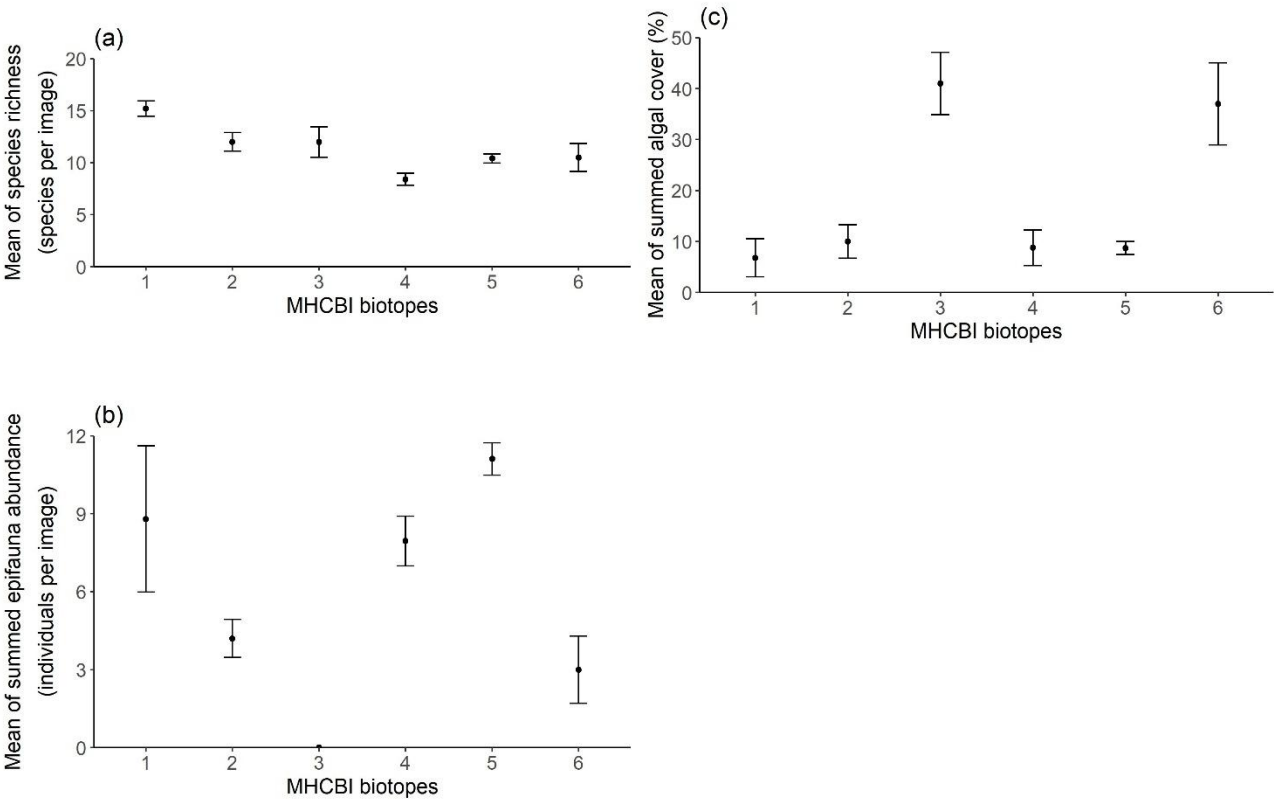


351

352 **Fig. 12.** Maps showing the estimated distribution of PRIMER biotopes (above) and MHCBI biotopes (below) within the
353 Langness MNR boundary.

354 3.1.6 Patterns among the MHCBI biotopes identified

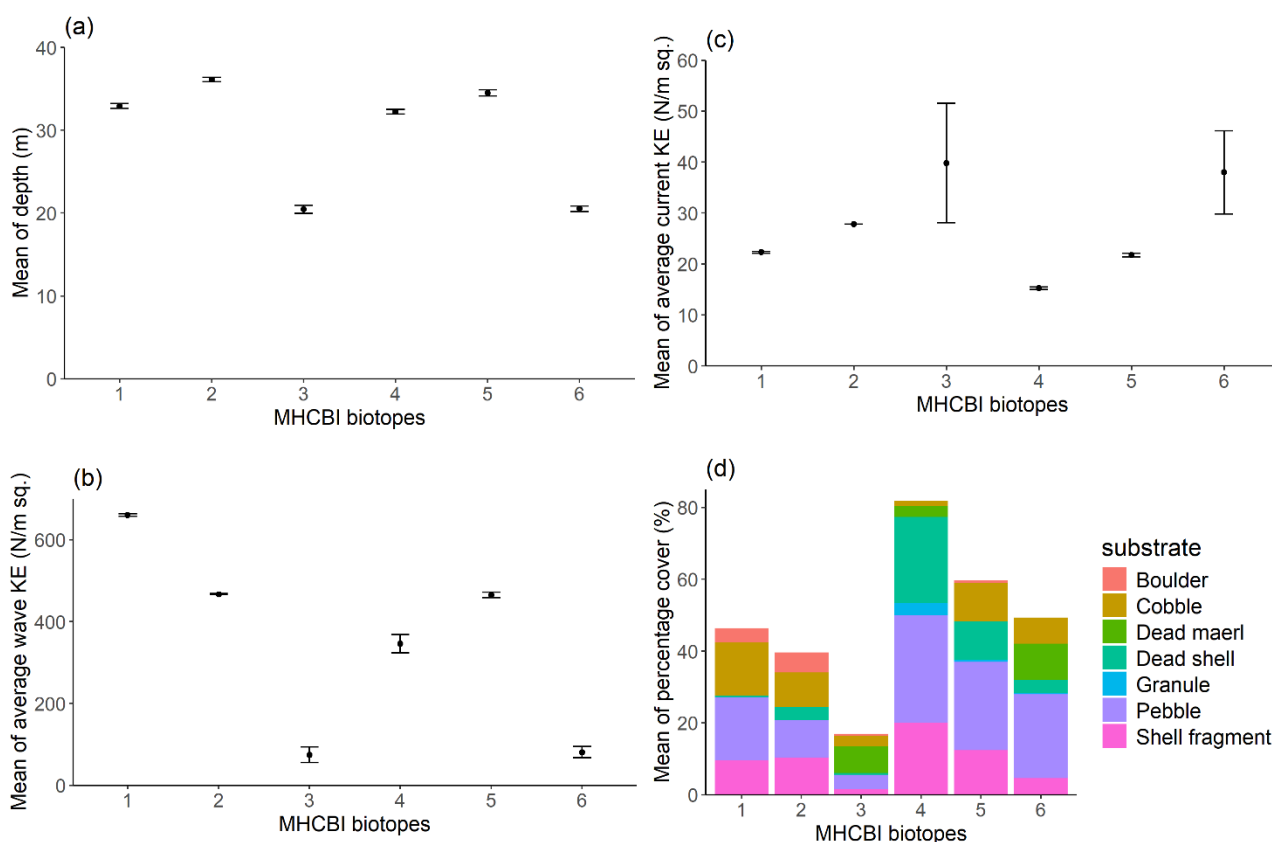
355 There was a significant variation in mean species richness among the MHCBI biotopes identified
356 ($F_{5,70} = 6.56, p < 0.001$), ranging from 8 taxa per image in biotope 4 to 15 taxa per image in biotope
357 1 (Fig. 13). In addition, a significant difference in countable epifauna abundance ($\chi^2 = 32.8, df = 5,$
358 $p < 0.001$) and algal percentage cover ($\chi^2 = 24.4, df = 5, p < 0.001$) was also found across the MHCBI
359 biotopes. Biotope 5 has the highest mean in the summed abundance of the countable epifauna (11
360 individuals per image), while the mean of the summed algal percentage cover peaked in biotope 3
361 (41%) (Fig. 13).



362 **Fig. 13.** Graphs showing the means ($\pm$ S.E.) of (a) species richness, (b) summed countable epifauna abundance, and (c)
363 summed algal percentage cover (%) across the 6 MHCBI biotopes identified. Biotope numbers align with those used in
364 Table 4.
365

366 Environmental variables also varied across the MHCBI biotopes identified, including water depths
367 (below CD), coverage of different substrate types, average current energy, and average wave energy
368 (Fig. 14). Species richness appeared to be relatively more correlated to the percentage cover of the
369 substrate types. Biotopes with a substratum of higher portions of cobbles and boulders, e.g. biotopes
370 1 and 2, have displayed higher species richness (Fig. 13 and 14). Taking all 11 environmental
371 parameters into consideration, the ‘BEST’ analysis using the taxa presence/absence data has found

372 that the community assemblage of the biotopes was best described by 5 factors, including water depth,
 373 percentage cover of boulders and pebbles, and average energy of current and wave (correlation of
 374 0.623).



375
 376 **Fig. 14.** Graphs showing the means ($\pm$ S.E.) of (a) water depth (m), (b) average wave energy ((N/m²), (c) average current
 377 energy ((N/m²), and (d) percentage cover of substrates (%) across the 6 MHCBI biotopes identified. Biotope numbers
 378 align with those used in Table 4.

379 3.2 BRUVS Sampling

380 Due to time constraints and gear malfunctioning, of the 20 BRUVS deployed, only data from 10
 381 BRUVS were analysed. The deployment sites of these BRUVS have a water depth (below CD)
 382 ranging from 21.8m to 37.3m, consist of 4 types of substrata, and were spread across 5 types of
 383 MHCBI biotopes identified previously (Fig. 15).

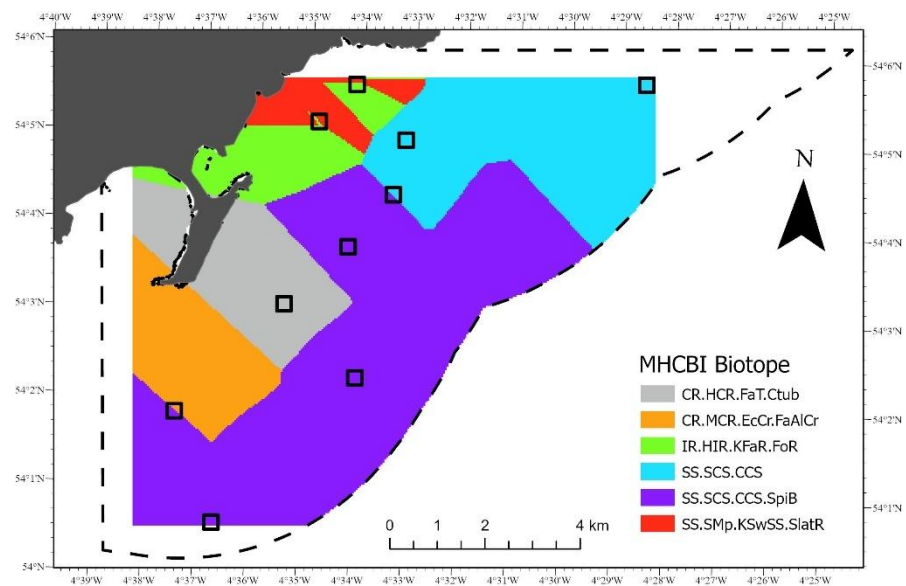
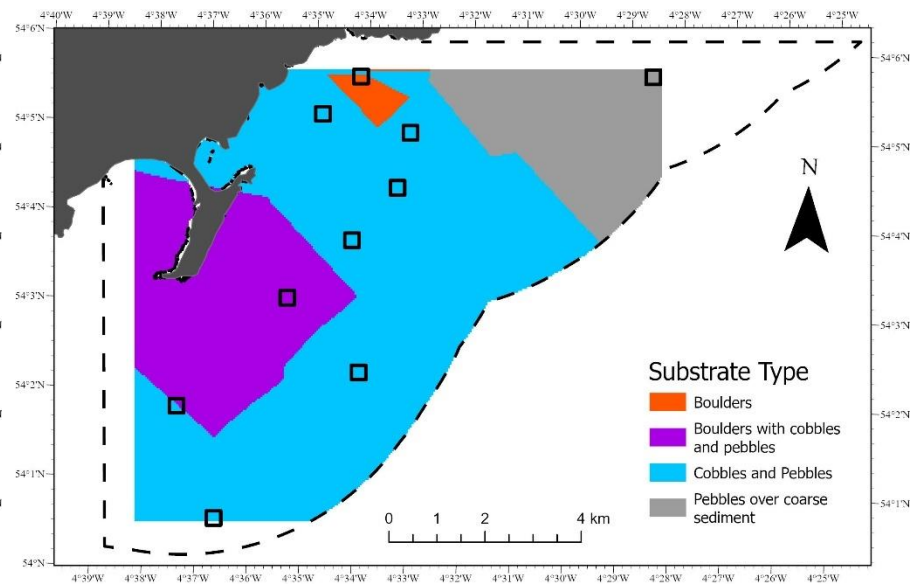
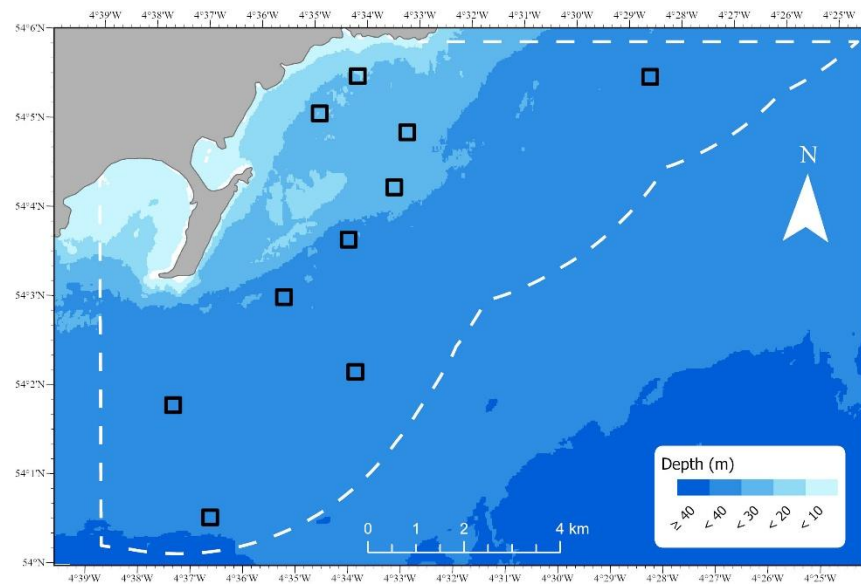


Fig. 15. Maps showing the bathymetry (below CD) (EDINA, 2020), substrate type, and identified MHCBI biotopes across the areas of the Langness MNR, with locations of BRUVS deployment (only those with video analysis done) overlaid on top.

Mobile species communities associated with each MHCBI biotope identified were significantly different ($R = 0.846$, $p = 0.001$) (Fig. 16), but the communities did not differ significantly when substrate types were used for the categorization of habitats ($R = 0.262$, $p = 0.167$). The species richness did not differ significantly across the MHCBI biotopes ($\chi^2 = 4.25$, $df = 4$, $p = 0.374$), but a relatively higher means was found in biotope 4 (12 taxa per footage) and biotope 6 (10 taxa per footage), in which their substratum was mainly composed of pebbles and cobbles, with some degree of coarse sediment (Fig. 15).

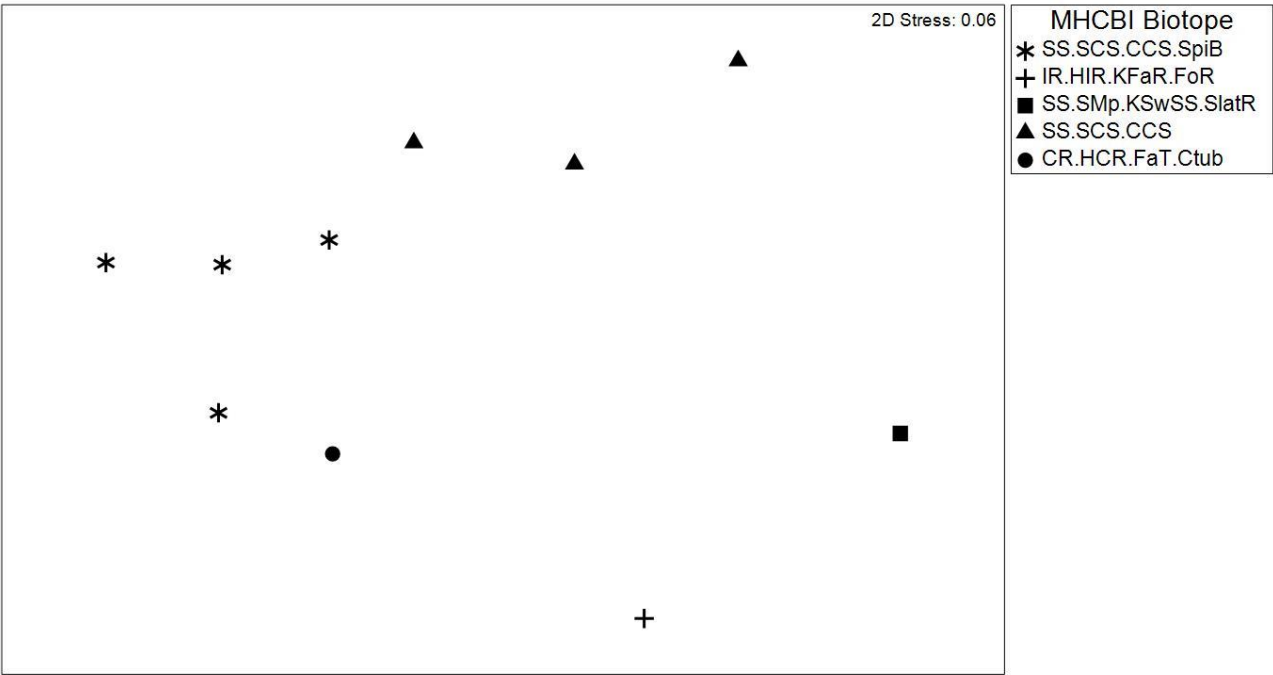


Fig. 16. An nMDS plot showing the relationship between mobile species communities of each BRUVS sample. The plot was created using the abundance data (MaxN), with each sample assigned a symbol according to its associated biotope. Points in close proximity indicate high similarity with each other while points further apart indicate low similarity.

A total of 41 taxa were identified from the analysed BRUVS footage (appendix, Table A5), including 15 teleosts (37%), 13 arthropods (32%), 6 molluscs (15%), 4 echinoderms (10%), 2 elasmobranchs (5%), and 1 ctenophore (3%). The most frequently encountered taxon was found to be the small-spotted catshark, *Scyliorhinus canicular*, which was present in every analysed BRUVS footage, with an average of 1.70 ± 0.40 individuals per footage and a maximum abundance of 4 individuals recorded in two BRUVS (in biotopes 4 and 6). The other elasmobranch recorded, the nursehound, *Scyliorhinus stellaris*, was only sighted in two BRUVS footages (in biotopes 1 and 6), with an abundance of 2 and 3 individuals respectively. The brown shrimp, *Crangon crangon*, was the second-most frequently encountered taxon, recorded in 8 of the BRUVS footage (in biotopes 1, 3, 4 and 5). It averaged at 9.75 ± 2.91 individuals per footage and a maximum abundance of 25 individuals was sighted in one BRUVS (in biotope 5). Other commonly sighted taxa, each recorded in 4 or more BRUVS footage,

include the edible sea urchin, *Echinus esculentus*, common whelk, *Buccinum undatum*, butterfish, *Pholis gunnellus*, Sea gooseberry, *Pleurobrachia pileus*, and Bernhard's hermit crab, *Pagurus bernhardus*. In addition, though only recorded in 2 (in biotopes 3 and 6) and 3 (in biotopes 4 and 5) BRUVS respectively, the poor cod, *Trisopterus minutus*, and netted dog whelk, *Tritia reticulata*, both have a relatively high mean abundance, averaging at 64.0 ± 57.0 and 20.7 ± 14.5 individuals per footage respectively. The poor cod has a maximum abundance record of 121 individuals in one BRUVS (in biotope 6) while that of the netted dog whelk was 49 individuals (in biotope 5).

Taking the environmental variables into account, in terms of a single factor, it was found that the mobile species communities were best described by the average energy of current in the area (correlation of 0.452). With an additional factor of average wave energy included, the correlation increased further (correlation of 0.495).

4. Discussion

Langness MNR contains a range of benthic habitats, from pebbly sand to algal-dominated stony rocky substrate. Most of the MNR areas are characterised by coarse sediments and the commonest taxa were encrusting species, such as Serpulidae spp. and *Parasmittina trispinosa*. Dead maerl and encrusting coralline algae were frequently encountered, both appeared in more than 40% of the analysed images; conversely, live maerl were rarely sighted and only recorded in 7 images. In comparison to other MNRs, the epifauna species richness in Langness was higher than Laxey and Niarbyl Bay and comparable to Port Erin Bay, Ramsey Bay, Douglas Bay (Garratt et al., 2022a, 2022b, 2022c, 2022d, 2022e).

Environmental processes are major determinants of marine communities and habitats. The hydrodynamic regime affects the distribution of different types of sediment, forming different substratum within an area. The varied substratum created different habitats and thus shaped different communities of species (Connor et al., 2004). This study has identified the substrate composition, water depth, and strength of wave and current energy as the most important factors in shaping the composition of benthic communities. It is believed that a further increase in variation of these factors would result in more significant differences among the communities identified. Meanwhile, it is important to note that environmental factors are interactive, as mentioned, and they do not act independently. The scope of this study is only to identify the factors that best correlate with the community data, not to study the effect of individual factors.

Environmental drivers of BRUVS communities were similar and also had a combination of water depth, strength of wave and current energy as the best descriptor of community composition. Though

442 the community composition was significantly different across the MHCBI biotopes, they did not
443 differ significantly across different types of substrata, and no association between species richness
444 and algal cover has been recognised. The condition might be explained by the mobile nature of the
445 species. As mentioned in Kaiser et al. (2020), mobile taxa are less associated with environmental
446 conditions when compared with the sessile component of a community. The mobile taxa identified in
447 the BRUVS communities, such as fishes and crustaceans, might not have specific habitat
448 requirements, but they rely on hydrodynamic activities to assist their movements, migration, and
449 foraging activities.

450 This study aimed to create a map of the distribution of biotopes, in which a biotope is assigned based
451 on biotic and abiotic factors (Connor et al., 2004). The PRIMER approach for biotope classification
452 only takes account of the percentage cover data of the analysed still images. Though the substrate
453 information is included, other environmental variables were excluded, as well as other taxa that were
454 absent in the percentage cover data. Conversely, the MHCBI classification system approach has
455 considered several environmental data, including depth and energy of wave and current, and has
456 included both the percentage cover and tax presence/absence data from the analysed images, for the
457 assignment of biotopes. In addition, with such an approach, the information obtained from the video
458 footage can also be considered for a more accurate assignment of the biotope, since still images only
459 provide sections of the seabed, whereas the continuity of video footage can offer a broader view of
460 the seabed. Given the MHCBI classification system approach can cope with variations in habitats and
461 communities, it is regarded as a more suitable approach for biotope assignment in this study.

462 Due to gear malfunctioning, the majority of still images (~90%) captured had minor focus issues and
463 thus, were not of the best quality for analysis. This issue had an impact on species identification, as
464 some taxa encountered could only be identified to family level or even assigned to broad descriptive
465 categories due to the blurriness of the image. It is believed that this might have affected the number
466 of taxa identified, the recognition of the overall community, and potentially the accuracy of biotope
467 assignments. However, since the issue was caused by equipment failure, not much could be done to
468 mitigate the impact for the current study. In future studies, it is suggested to carry out several in-water
469 trials for the survey equipment before the actual survey, to ensure that the settings are correct and the
470 gear itself is functioning properly. As for this study, if possible, it would be ideal to redo some of the
471 tows, to get a more accurate picture of the habitats and communities present in the area.

472 Though the identification level for the same taxa was consistent throughout the data analysis process,
473 there was a variation in the taxonomical levels for the identification of different taxa. Some were
474 identified to species level, while to the other extreme, some were only able to be grouped into broad

475 descriptive categories, especially for bryozoans, hydroids, and algae. Several species are likely to be
476 present in the same group for these categories. Thus, if all encountered taxa have been identified to
477 species level, the assignments of biotopes might differ.

478 One of the drawbacks of benthic imagery surveys is that they failed to account for the presence of
479 small epifauna as well as infaunal community (Beisiegel et al., 2017). Since infauna are a key
480 component of benthic communities (Elliott, 1994), and a large proportion of the sampling stations in
481 this study have sediment-based substratum, the inclusion of benthic grab sampling would help
482 complete the identification of all taxa present in a particular benthic community, improving the
483 accuracy of the eventual biotope assignments.

484 Constrained by time, it was only possible to conduct 1 tow in each sampling grid, and though still
485 images of benthic habitats were taken at frequent intervals of 10 seconds in each tow, only 1 image
486 was selected for analysis every 120 seconds. Many previous studies elsewhere on coastal benthic
487 communities (e.g. Hernández-Fernández et al., 2019; Wahl, 2001) have found that the assemblage,
488 coverage, and diversity of benthic fauna could vary on a small spatial scale. As such, in this study,
489 the single tow done in each sampling grid and the limited amount of analysed still images might not
490 have captured all habitats and communities present within the area of the grid. It is therefore suggested
491 that future studies could focus on areas of particular interest, complete multiple tows in a single grid,
492 and analyse more still images per tow, aiming to obtain a more representative assessment of the
493 habitats and communities present.

494 **Declaration of competing interest**

495 The authors declare that they have no known competing financial interests or personal relationships
496 that could have appeared to influence the work reported in this paper.

497 **Acknowledgements**

498 I would like to thank Professor Stuart Jenkins, Matthew Garratt, Dr Isobel Bloor, Dr Matthew
499 Coleman and Dr Peter Duncan for their knowledge, guidance, and support provided throughout my
500 MSc project. Many thanks to all the boat crew on Barrule, particularly Rob Annett, for their assistance
501 and guidance during the data collection process. Thanks to the staff at DEFA for their help and
502 hospitality during my stay. Many thanks to Maya Harries for helping with part of my data extraction
503 work. I would also like to all the staff of the School of Ocean Sciences, who have provided me loads
504 of support throughout my MSc course, particularly Professor Stuart Jenkins, Dr Craig Robinson, and

505 Dr Martin Skov. Finally, I want to thank my coursemates and parents, who have been incredibly
506 supportive over the year.

507 **Appendix A. Supplementary data**

508 Supplementary data to this article is attached at the end.

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Supplementary data

Table A1

List of taxa identified from the benthic still images taken in the Langness MNR, Isle of Man.

Phylum	Taxon	Phylum	Taxon
Annelids	<i>Lanice conchilega</i>	Molluscs	<i>Aequipecten opercularis</i>
	Serpulidae spp.		<i>Anomiidae</i> spp.
Arthropods	<i>Galathea intermedia</i>		<i>Glycymeris glycymeris</i>
	<i>Pagurus bernhardus</i>		Buccinidae spp.
	<i>Pagurus prideaux</i>		<i>Calliostoma zizyphinum</i>
	<i>Pandalus</i> spp.		<i>Colus</i> spp.
	<i>Pisidia longicornis</i>		<i>Coryphella</i> spp.
	<i>Polybius depurator</i>		<i>Gibbula magus</i>
	Small pinkish crab		<i>Nucella lapillus</i>
	<i>Balanus</i> spp.		<i>Ocenebra erinaceus</i>
Bryozoans	<i>Crisularia plumosa</i>		Rissoiidae spp.
	<i>Parasmittina trispinosa</i>		<i>Steromphala cineraria</i>
	<i>Reptadeonella violacea</i>		<i>Trivia monacha</i>
	<i>Schizomavella</i> spp.		<i>Acanthochitona</i> spp.
	<i>Crisia</i> spp.		Greyish shell gastropod
	Brownish green encrusting bryozoan	Sponges	<i>Dysidea fragilis</i>
Ascidians	<i>Ascidia</i> spp.		<i>Hemimyscale columella</i>
	<i>Clavelina lepadiformis</i>		<i>Tethya aurantium</i>
Teleosts	<i>Pomatoschistus</i> spp.		Dark brownish encrusting sponge
	<i>Taurulus bubalis</i>		Orange colonial sponge
Cnidarians	<i>Corynactis viridis</i>	Turf complex	Yellow colonial sponge
	<i>Cylista elegans</i>		Mixed turf of algae with bryozoan and/or hydroid
	<i>Epizoanthus</i> spp.		Mixed turf of bryozoan and hydroid
	<i>Synarachnactis lloydii</i>	Algae	Muddy surface turf
	<i>Urticina felina</i>		<i>Bonnemaisonia asparagoides</i>
	<i>Abietinaria abietina</i>		Corallinaceae spp.
	<i>Halecium</i> spp.		<i>Cryptopleura ramosa</i>
	<i>Halopteris</i> spp.		<i>Delesseria sanguinea</i>
	<i>Hydrallmania falcata</i>		<i>Dictyota dichotoma</i>
	<i>Kirchenpaueria pinnata</i>		Hapalidiaceae spp.
	<i>Nemertesia antennina</i>		<i>Heterosiphonia plumosa</i>
	<i>Nemertesia ramosa</i>		<i>Laminaria hyperborea</i>
	<i>Tubularia indivisa</i>		<i>Nitophyllum punctatum</i>
	<i>Alcyonium digitatum</i>		<i>Phycodrys rubens</i>
	Muddy branching hydroid		<i>Phyllophora</i> spp.
	Short stalky hydroid		<i>Plocamium</i> spp.
Echinoderms	<i>Crossaster papposus</i>		<i>Rhodophyllis</i> spp.
	<i>Henricia</i> spp.		Branching red seaweed
	<i>Echinus esculentus</i>		Flat brown seaweed
	<i>Psammechinus miliaris</i>		Fluffy colonial red seaweed
			Fluffy green seaweed
			Thin flat red seaweed

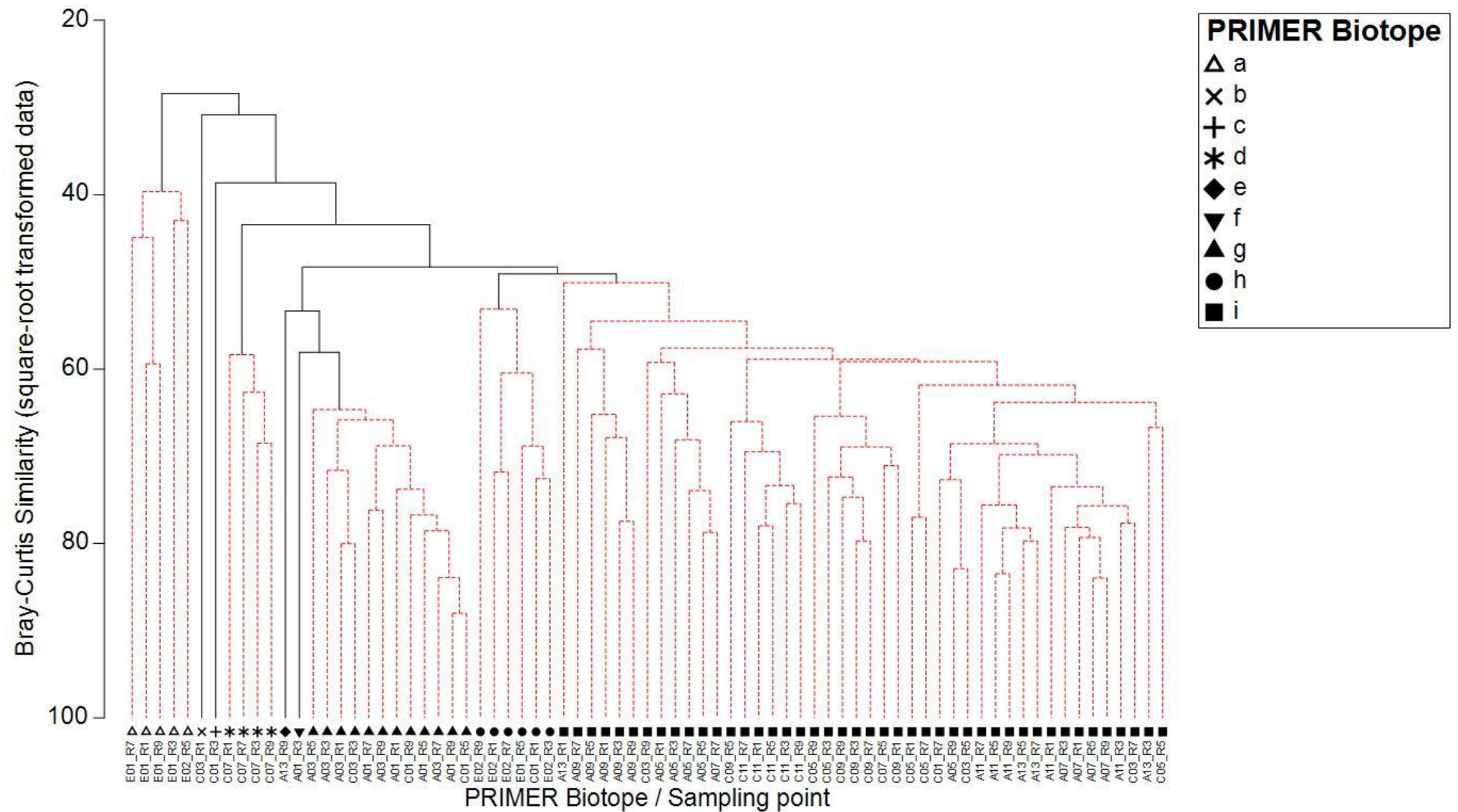


Fig. A1. A dendrogram showing the result of the CLUSTER analysis with the inclusion of similarity profile (SIMPROF) tests ($p < 0.05$) on the similarity matrix of the square-root transformed percentage cover data. Each resultant ‘cluster’ is defined as a distinctive biotope.

Table A2

Results of the SIMPER analysis on the community composition (percentage cover data) of the biotopes identified through the PRIMER statistical approach. The taxa that contributed the most to the similarities within the biotope are listed, with the percentage cut-off set at 90%.

Taxa / Substratum	Av.%cover	Av.Sim	Sim/SD	Contrib%	Cum.%
Biotope a - average similarity: 42.99%					
Muddy surface turf	2.99	8.59	3.46	19.99	19.99
Mixed turf of algae with bryozoan and/or hydroid	3.51	7.69	1.71	17.88	37.86
Corallinaceae spp.	3.16	6.86	2.51	15.95	53.81
Pebble	2.36	3.79	1.02	8.82	62.64
Branching red seaweed	1.78	3.79	1.14	8.81	71.45
<i>Phyllophora</i> spp.	1.95	3.24	0.94	7.53	78.98
<i>Plocamium</i> spp.	1.88	2.61	0.62	6.07	85.05
Thin flat red seaweed	1.41	1.7	0.58	3.96	89.01
<i>Heterosiphonia plumosa</i>	1.59	1.03	0.32	2.39	91.4
Biotope d - average similarity: 61.48%					
Pebble	4.39	11.48	4.87	18.67	18.67
Cobble	3.95	9.35	2.68	15.21	33.88
<i>Corynactis viridis</i>	3.47	6.7	2.05	10.89	44.78
Mixed turf of bryozoan and hydroid	2.51	5.75	2.81	9.35	54.12
Shell fragment	2.51	5.71	3.09	9.3	63.42
Muddy surface turf	1.71	4.58	6	7.46	70.88
Anomiidae spp.	1.56	4.29	25.73	6.97	77.85
<i>Tubularia indivisa</i>	1.79	3.12	0.91	5.07	82.92
Boulder	1.47	2.4	0.89	3.91	86.83
Serpulidae spp.	1.32	2.2	0.91	3.59	90.41
Biotope g - average similarity: 69.46%					
Pebble	5.71	21.35	5.05	30.73	30.73
Shell fragment	5.05	19.17	6.97	27.59	58.33
Dead shell	4.6	15.26	2.45	21.98	80.3
Granule	1.35	3.31	1.06	4.77	85.07
Corallinaceae spp.	1.26	2.96	0.82	4.27	89.33
Dead maerl	1.44	2.88	0.67	4.14	93.48
Biotope h - average similarity: 60.69%					
Pebble	4.97	14.62	3.28	24.09	24.09
Dead maerl	3.8	11.33	4.81	18.68	42.77
Corallinaceae spp.	3.11	8.9	3.66	14.67	57.44
Shell fragment	2.78	6.73	2.59	11.1	68.54
Cobble	2.56	6.24	1.35	10.28	78.82
Dead shell	1.62	3.45	1.35	5.69	84.51
Muddy surface turf	1.62	3.39	1.35	5.58	90.09
Biotope i - average similarity: 59.66%					
Pebble	4.61	12.75	2.37	21.38	21.38
Shell fragment	3.16	8	2.02	13.41	34.79
Cobble	2.88	7.24	1.96	12.14	46.93
Muddy surface turf	2.69	6.95	1.99	11.64	58.57
Serpulidae spp.	2.3	6.08	2.43	10.18	68.76
Corallinaceae spp.	2.6	6.04	1.4	10.13	78.89
Dead shell	2.49	4.74	0.95	7.94	86.83
Brownish green encrusting bryozoan	1.81	3.83	1.06	6.42	93.25

Table A3

Results of the SIMPER analysis on the community composition (percentage cover data) of the biotopes identified through the application of the MHCBI classification system. The taxa that contributed the most to the similarities within the biotope are listed, with the percentage cut-off set at 90%.

Taxa / Substratum	Av.%cover	Av.Sim	Sim/SD	Contrib%	Cum.%
CR.HCR.FaT.Ctub - average similarity: 59.30%					
Pebble	4.08	10.33	4.48	17.42	17.42
Cobble	3.65	8.45	2.94	14.26	31.68
<i>Corynactis viridis</i>	3.34	6.93	2.52	11.69	43.37
Shell fragment	2.9	6.46	2.81	10.89	54.27
Muddy surface turf	2	4.83	5.38	8.15	62.41
Anomiidae spp.	1.53	4.29	28.17	7.24	69.66
Mixed turf of bryozoan and hydroid	2.01	3.45	1.03	5.81	75.47
Boulder	1.74	3.22	1.08	5.43	80.9
Corallinaceae spp.	2.09	3.1	0.97	5.23	86.13
Serpulidae spp.	1.34	2.63	1.16	4.44	90.57
CR.MCR.EcCr.FaAlCr - average similarity: 69.23%					
Serpulidae spp.	3.44	10.15	8.24	14.66	14.66
Muddy surface turf	3.2	9.54	6.78	13.79	28.45
Pebble	3.2	9.38	6.59	13.54	41.99
Shell fragment	3.15	8.68	3.96	12.53	54.52
Cobble	3.02	8.31	3.78	12.01	66.53
Brownish green encrusting bryozoan	2.81	7.56	2.97	10.92	77.46
Corallinaceae spp.	2.99	7.48	3.09	10.8	88.25
Boulder	2.04	4	1.03	5.78	94.04
IR.HIR.KFaR.FoR - average similarity: 47.81%					
Mixed turf of algae with bryozoan and/or hydroid	4.32	11.89	4.13	24.87	24.87
Muddy surface turf	3.31	10.33	19.98	21.61	46.48
Corallinaceae spp.	3.03	8.44	5.6	17.65	64.13
<i>Phyllophora</i> spp.	2.7	6.69	2.7	13.98	78.12
Pebble	1.67	2.72	0.86	5.69	83.81
<i>Plocamium</i> spp.	1.41	1.61	0.41	3.36	87.17
Dead maerl	1.88	1.43	0.41	2.99	90.16
SS.SCS.CCS - average similarity: 58.75%					
Pebble	5.13	16.56	2	28.19	28.19
Shell fragment	4.25	14.02	2.54	23.86	52.06
Dead shell	4.45	13.54	2.01	23.05	75.11
Corallinaceae spp.	1.91	3.63	0.9	6.18	81.28
<i>Glycymeris glycymeris</i>	1.1	2.45	0.81	4.16	85.45
Granule	1.3	2.17	0.69	3.7	89.14
Dead maerl	1.21	1.97	0.54	3.36	92.5
SS.SCS.CCS.SpiB - average similarity: 59.65%					
Pebble	4.68	13.17	2.35	22.08	22.08
Shell fragment	3.13	7.87	1.8	13.2	35.28
Cobble	2.97	7.64	2.01	12.81	48.1
Muddy surface turf	2.57	6.39	1.74	10.72	58.82
Serpulidae spp.	2.27	6.31	3.01	10.58	69.4
Dead shell	2.62	5.16	0.97	8.65	78.05
Corallinaceae spp.	2.33	5.01	1.14	8.4	86.45
Brownish green encrusting bryozoan	1.72	3.65	1.03	6.11	92.56

Taxa / Substratum	Av.%cover	Av.Sim	Sim/SD	Contrib%	Cum.%
SS.SMp.KSwSS.SlatR - average similarity: 51.84%					
Pebble	4.76	13.93	4.62	26.87	26.87
Corallinaceae spp.	3.39	8.09	2.18	15.61	42.48
Cobble	2.46	6.19	1.36	11.94	54.42
Dead maerl	2.77	5.78	1.17	11.15	65.57
Shell fragment	1.8	3.44	1.27	6.63	72.2
Muddy surface turf	1.67	3.41	1.32	6.58	78.79
Dead shell	1.62	3.16	1.32	6.1	84.89
Hapalidiaceae spp.	1.45	2.24	0.74	4.33	89.21
Mixed turf of algae with bryozoan and/or hydroid	1.14	1.88	0.78	3.63	92.84

Table A4

List of taxa identified from the BRUVS footages taken in the Langness MNR, Isle of Man.

Phylum	Taxon	Phylum	Taxon
Arthropods	<i>Atelecyclus rotundatus</i>	Teleosts	<i>Gobius paganellus</i>
	<i>Cancer pagurus</i>		<i>Labrus bergylta</i>
	<i>Cragon cragon</i>		<i>Labrus mixtus</i>
	<i>Crangon allmanni</i>		<i>Pholis gunnellus</i>
	<i>Inachus spp.</i>		<i>Pollachius pollachius</i>
	<i>Liocarcinus spp.</i>		<i>Pomatoschistus flavescens</i>
	<i>Necora puber</i>		<i>Pomatoschistus spp.</i>
	<i>Pagurus bernhardus</i>		<i>Syngnathus schlegeli</i>
	<i>Pagurus prideaux</i>		<i>Trisopterus minutus</i>
	<i>Eurynome aspera</i>	Ctenophore	<i>Pleurobrachia pileus</i>
	<i>Munida rugosa</i>	Echinoderms	<i>Henricia spp.</i>
	<i>Pisidia longicornis</i>		<i>Marthasterias glacialis</i>
	<i>Xantho spp.</i>		<i>Echinus esculentus</i>
Elasmobranchs	<i>Scyliorhinus canicula</i>		<i>Ophiothrix fragilis</i>
	<i>Scyliorhinus stellaris</i>	Molluscs	<i>Bittium reticulatum</i>
Teleosts	<i>Blenniiformes spp.</i>		<i>Buccinum undatum</i>
	<i>Blennius ocellaris</i>		<i>Calliostoma zizyphinum</i>
	<i>Chelidonichthys cuculus</i>		<i>Neptunea antiqua</i>
	<i>Ctenolabrus rupestris</i>		<i>Nucella lapillus</i>
	<i>Diplecogaster bimaculata</i>		<i>Tritia reticulata</i>
	<i>Gobiidae spp.</i>		