

9 January 2020

Subject: Revision of MS No.: bg-2019-304

Dear Associate editor,

We would like to thank both reviewers for their thorough revision of the manuscript.

We addressed the main concerns regarding Figure 3 and Table A1 and elaborated on certain parts of the methodology. Figures were redone to make sure the same dataset was used and the text was corrected whenever necessary. We argument the use of the ind/100m metric due to the limitation of the data collection, which is explained in more detail below.

All our other detailed replies to the reviewers' and associate editor's queries, the changes made to the manuscript (with line references of the revised manuscript) and a marked-up manuscript version are enclosed,

Sincerely,

Daphne Cuvelier

Associate Editor Comments to the Author:

Dear authors,

2 reviewer comments have been received for your manuscript that you have both responded to. Both reviewers have made substantial comments that need to be addressed and incorporated during the revision of your manuscript. This should be possible considering your replies.

A few additional comments (in addition to the reviewer's comments) from my side are:
I agree with the reviewers that Figure S1 presents essential information on the size of seamounts, water depths and locations of the transects. Therefore it should be moved into the article. However, important information is not accessible from the current figure, because axes and color code labels are not readable (even if zooming into the PDF) due to too low resolution of the figure.

A: Figures are embedded in a pdf as requested when submitting a manuscript for Biogeosciences, which is why their resolution is not that high. We have increased the letter type so that the numbers are readable and increased the scales so that the size of the seamounts can be deduced. We will provide high resolution figures when the manuscript is accepted.

During the revision, please take care to significantly improve the quality of all figures (Fig.7 has a similar problem) that labels and annotations become readable.

A: See comment above. We reorganised figure7 to enhance its readability whilst embedded in the pdf.

It may be useful to add a table in the supplement containing specific information for each seamount raised by R#1 (i.e. size, summit and base depth, average slopes at sampling/video depths and average current direction/velocity (e.g. sampling site upstream or downstream), etc).

A: All this information can be found in the text and in the improved Fig 1. Slope can be calculated using the depth gradient and the length of the transect.

Sampling sites were all downstream – this is mentioned in L83-84 of the revised ms.

R#1, comment 3: The distance between the ROV laser spots on the videos (I believe it is 50 cm for the parallel beams, but please confirm with the ROV team of Kiel6000) should enable estimating the width of the visual field and hence allow providing an area for the transects, i.e. ind/m². This would increase comparability.

A: This was considered previously and extensively, but lasers were not always visible during the entirety of the seamount dives. Changing camera pan and tilt and forward-looking view also introduce bias in any possible surface calculations (e.g. perspective view). Additionally, the non-constant travel speed, and changing ROV altitude complicates the surface calculations. In order to counter this, we considered subsampling images at a predefined distance interval (e.g. every 10 m) and extrapolate the surface covered. Nevertheless, as stated before, lasers were not visible/operational during the entire dive and would thus only increase over-or underestimation of the surface covered and therefore of the faunal densities. Overall, the ind/100m appears a more

correct representation of the patterns observed and therefore the *in situ* reality and allowed for an in-detail comparison to the nodule field dives, which was one of the main objectives of this study.

R#1, comment L.288-295: This is an important point and I agree with the reviewer that more discussion on possible reasons would be helpful. For example, the seafloor POC flux in APEI3 seems to be reduced by ~30% compared to the investigated license areas (see biogeochemistry paper of Volz et al. (2018) Deep-Sea Research I 140).

A: We mitigated the overall statement that the nodule coverage would be the main driving factor and added more on possible other factors at play using the reference proposed. L329-331.

R#2, comment 3: Where is Fig. R1 you mention in your reply?

A: It was attached as a supplementary figure during the first review round and can be found in the "reply to reviewer 2" .pdf. We have inserted the figure in the appendix (Fig. A1) and referred to it in Section 3.2.

Reviewer 1:

R1: My main concern is that no environmental data are presented. In addition to missing hydrographical data, there is no detailed description of the sampling sites, particularly the seamounts, such as size, summit and base depth, inclination of slopes/general bathymetry, or current field. More important, no information is given on habitat types encountered along the transects. It is well known that substrate can vary considerably at short distances at seamounts, and of course megafaunal communities are strongly associated with substrate type. This is briefly mentioned in the discussion, but I do not understand why this information is not provided and analysed in the results. It should easily be available from the video footage.

A: Regarding the reviewer's comment on the description of the sampling sites: There was a supplementary figure (Fig. S1) from which the size of the seamount, the depth of the base and bathymetry can be deduced and which we now have incorporated into Figure 1 (as requested by the other reviewer and associate editor).

We have added information on the amount of hard substrata under the form of 3 categories: (1) Predominant soft substrata (<40% hard substrata), (2) mix or transition (between 40 and 60% hard substrata) and (3) predominant hard substrata (>60% hard substrata), annotated per 10m distance unit. This has been added to the methods section (L121-124). The amount of hard substrata was linked with faunal observations in higher taxonomic groups. No significant correlations between substratum type and faunal abundances were found for ROV02, ROV04 and ROV09. This is most likely due to the amount of hard/mix/soft substrata, which were almost equally distributed over the transects (varying between 16-39%, 30-51% and 15-38%, respectively). This highlights the need for a more in-detail assessment of substratum type, which is currently underway as the geomorphology of the seamount transects is being modelled (based on backscatter data) and investigated in detail, but this falls outside the scope of the current manuscript.

Based on the preliminary substratum results, a paragraph was added in section 3.1 L206-218

Concerning the hydrographical properties, a link with the POC-flux as known for the CCZ area was elaborated, also following the associate reviewer's suggestion (L328-332). Following the comments of the other reviewer, we have added the predominant current direction at the CCZ in the Methods section.

R1: There are also some methodological issues. The basic problem, as also admitted by the authors in the discussion, is that only a very limited number of rather short transects without replications are available and that transects at the seamounts and at the nodule fields were taken at different depths; in the case of Mann Borgese Smt the depth sampled was nearly 3000 m less than on the corresponding nodule field, and hence the data are hardly comparable. Although the depth difference at the other sites was much smaller, it may also limit the comparability of the data. This is mentioned in the discussion, but the consequences should be elaborated in more detail, and it makes the conclusion that "seamounts appear inadequate as refuge areas to help maintain nodule biodiversity" disputable.

A: In name of all the co-authors, I think we were very cautious in our conclusions, recognising the sampling shortcomings as well as the limited amount of data. We purposefully stated "*Based on our current knowledge; seamounts appear inadequate as refuge areas to help maintain nodule biodiversity.*" And then urged on for more proper sampling to adequately corroborate or refute observations done here. We recognised the shortcomings throughout the manuscript. We think it is important to take a look at the entirety of the sentences written and to not take parts of them out of context.

Moreover, because of the unknown impacts and extent of mining, we could speculate that only the communities living outside a certain range in distance across the seafloor and upwards in the water column, will be protected from the mining plumes (and other impacts). Hence, it is relevant to also investigate "shallower" areas of the seamounts, as there is a possibility that recolonisation will start from these somewhat shallower areas.

R1: I am also not convinced that the quantification of the samples is correct. In section 2.1, the authors state that the altitude of the ROV was "kept constant whenever possible". Apart from not providing the information at which target altitude the ROV was kept, and whether this was the same at all transects, the authors inform in section 2.2 that, due to varying altitude as well as pan and tilt of the camera, "surface coverage" could not be used for standardisation and instead just transit length was used. However, since the field of view and thus the number of visible objects per unit transit section depend on the altitude and angle of the camera, the standardisation to 100 m transit sections, without taking into account the varying field of view, could strongly bias the results. Generally, the methods section has to be improved with much more detail.

A: Target altitude was 2m above seafloor and travel speed ~0.2m/s, though interrupted by sampling actions, instrument check-ups, exploration, object avoidance (in the case of the uphill seamount transects) etc. L100-102

Regarding the comment that ind/100m might not be a good metric, we would like to clarify that the length of the transects was calculated only taking into account the parts of the dive when the ROV was visualising the seafloor. The parts of the dives where the ROV was higher up in the water column (i.e. >10m altitude) and/or not visualising the seafloor (e.g. Transiting or checking ROV parts

or instruments) whilst travelling forward, were omitted out of these calculations, thus granting a best estimative possible and allowing for comparison.

We agree with the reviewer's observation that changing altitude and angle of the camera bias the observations, but by excluding the parts of the transect that were too high up, it is the best approximation possible with the data at hand.

Throughout the paper, we acknowledge the importance of performing standardising video transects and recognise the shortcomings of our study to this respect. However, on seamounts we cannot fly straight all the time, it does not necessarily prevent comparisons between transects, because the bias is systematic across transects. Moreover, while this shortcoming may pose limitations to quantitative comparisons, it does not preclude qualitative comparison between habitats, which are the main objective of this study.

Information on ROV altitude and transects length calculations were added in the Methods section with the following paragraph:

"For the transect length calculation for each dive, we omitted all parts of the video footage in which the ROV was at an altitude of >10m, or sections where the ROV was not visualising the seafloor (e.g. during transiting or inspecting ROV parts or instruments). Visualisation of ancient disturbance tracks were omitted as well, as these fell out of the scope of the article." L128-132

R1: Further, it is not clear how the investigations made in this study relate to those by Vanreusel et al (2016) who also presented results for epifaunal communities in the CCZ, comparing APEI, BGR, GSR and others. Obviously the sampling was done on the same cruise using the same gear. Were the same nodule field transects analysed? If yes, this has to be justified, the additional value of this study as compared to Vanreusel et al (2016) has to be demonstrated (apart from the additional seamount transects) and any overlap and differences in the analysis indicated. If not, a thorough comparison between the results of both studies is necessary.

A: Vanreusel et al. was based on a subset of the BGR, GSR and APEI3 nodule field videotransects/dives analysed here (they investigated 2740 m of the 6100m nodule fields transects as presented here or 44% of our study), and, as stated by the reviewer, they did not study the seamounts. Moreover, and most importantly, Vanreusel et al. did not identify individuals to morphospecies level but stayed at a higher taxonomical level such as Actiniaria, Echinoidea etc.

The following paragraph has been added in the ms for clarification purposes, in methods 2.2. L115-120

"A subset of the nodule field transects from BGR, GSR and APEI3 was presented by Vanreusel et al. (2016), and correspond to 44% of what we studied here and limited organism identification to a higher taxonomic level (Order (e.g. Alcyonacea) or Class (e.g. Ophiuroidea)). In our study, the entire transects (100%) were annotated to morphospecies level, allowing a more detailed comparison between seamounts and nodule fields."

Specific comments

R1: Use consequently "Ophiuroidea, Asteroidea" etc. instead of "ophiuroids, asteroids" etc.

A: Ok

R1: Abundances given in the text are not always consistent with those presented in Tab. A1. I did not check all entries, but two examples caught my eye: A total abundance of 89.2 ind/100 m is given for ROV10 in Tab. 1 and in the text, but summing up all observations in Tab. A1 results in ca. 67 ind/100 m. Another example: For Porifera, numbers given in the text match those in Tab. A1 for Rüppel and Senckenberg, but those presented for Heip and Mann Borgese are much lower than in the table (3 vs. 7.5 and 0.68 vs. 1.9, respectively). This has to be checked and resolved.

A: We carefully and meticulously checked all the densities, abundances and number of observations, to make sure they were correct and verified that the correct dataset was used for all figures and graphs.

R1: Line 42: Insert common definition for "seamount" and citation

A: We added the definition from the glossary of the International Seabed Authority: *"Seamounts are defined as isolated sub-surface topographic feature, usually of volcanic origin, of significant height above the seafloor (International Seabed Authority (ISA), 2019)"* on L42-43

<https://www.isa.org.jm/scientific-glossary/>

R1: 73: When did the sampling take place?

A: In 2015, this was added on L74

R1: 96: What is the difference between "exploration and opportunistic sampling"? More detail is needed.

A: The words chosen are rather self-explanatory. Explorative dives are dives when a site is visited for the first time and observations made during the dive are key to decide what happens during its course, e.g. sampling when the occasion presents itself or just imagery sampling. It was also mentioned in the text of the first submitted ms (L113-114) and now in L125-126.

R1: 98: What does "whenever possible" mean? 90 % of the transects? What was the target altitude of the ROV, and was it the same at all transects? How did panning and tilting affect the field of view? (see also general comments).

A: See answer above

R1: 94-99: Generally, much more information on the sampling mode is necessary, including sampling strategy (e.g., straight line, deviations for interesting objects etc.), ROV speed, target altitude, field of view etc.

A: See answer above

R1: 104: What is "ID's"? I guess it should be IDs, but "ID" is not defined in the text. Same in line 111.

A: ID from identification. This was altered.

R1: 114: Is there a reason that specimens collected were obviously not used for proper identification?

A: Samples were used for proper identification whenever possible. Multidisciplinary research cruises such as SO239 based on larger research projects (JPIO) tend to have a multitude of institutes involved, with different or overlapping interests. Samples taken during the cruise were distributed and divided over different institutes, each working towards their own objectives. The organisms sampled for which we received identifications were incorporated as such, e.g. Porifera identifications as included here. Though as stated in ms L280-282, even when organisms were sampled and identified, they were hard to extrapolate across the video imagery. Same reasoning applied for the Ophiuroidea where many species were revealed based on the samples, though impossible to annotate or differentiate based on the imagery footage (Christodoulou et al 2019). Hence, no information on its abundance, distribution or even presence in other areas than the one sampled can be included. There is no use of having a name for one (sampled) species from one single location if you cannot recognise it elsewhere.

R1: 118: Which statistical testing? Did the authors use tests other than nMDS? If yes, they have to be described here in detail

A: We have added information on the Kendall species Associations test carried out. L136-138

R1: 127: Here and throughout the text: two significant digits are sufficient, for example 7.6 instead of 7.59 or 89 vs. 89.23. The two decimals pretend a non-existing precision of the data.

A: Ok

R1: 151/152: Aren't Acrocirridae polychaetes as well? ("... Acrocirridae were observed. . . as well."). Do you probably mean they were observed in high densities in some of the transects?

A: This was corrected.

R1: 189/190: This belongs into the discussion.

A: This is also mentioned in the discussion and it was mentioned here as well to recognise the limited sampling. This links back to the methodological issues as stated by the reviewer, which we fully recognise throughout the manuscript.

R1: 193: Insert ", respectively" after "Table A1"

A: Ok

R1: 195: "less linear" - how was this assessed? I cannot see any linear or non-linear relations in Fig. 4d, nor can I see any curves crossing.

A: Replaced linear by straightforward. Curves cross at smaller sample sizes (<100 individuals) for ROV13,08 and 10.

R1: 198: What does "small sample size" mean? I think the sample size in this study was always small.

A: Less than 100 individuals, this was added.

R1: 205: Should read "least overlap". Explain similarity between these findings and the results from the seamount: For both habitat types, the samples at APEI3 had least overlap with the other sites.

A: Ok

R1: 220ff: According to Fig. 3 (not Fig. 8!), the majority of ophiuroids on the nodule fields were unidentified

A: This was corrected to *"The majority of the very abundant Ophiuroidea observed at the CCZ seamounts were small and situated on hard substrata (morphospecies 5), while most of the Ophiuroidea at nodule fields (including morphospecies 6) were observed on the soft sediments. Morphospecies 6 was only rarely observed on the seamounts (Fig. 3)"* L253-256

R1: 244/245: This is not clear. Variation "along the video transects" was obviously not analysed and cannot be seen in Fig. 5. Probably the authors mean "between transects"?

A: This was corrected to *"among the video transects of both seamounts and nodule fields"*

R1: 251: Kendall's coefficient is not mentioned in methods section. See comment above.

A: This was added, see reply above

R1: 255-258: This makes no sense. If sampling depth differs between seamounts and NF, and nMDS distinguishes between seamount and NF groups, then the grouping must correspond also to depth sampled. Omit this paragraph (and Fig. 7b) and state in the discussion that differences between seamounts and NF could be a result of different depths sampled.

A: It was already stated in the discussion, but in our opinion it is a visual presentation of this statement, which is why we decided to keep it for now.

R1: 262: This is not quite clear. Rephrase: ". . . at different locations and additionally, for the seamounts, different depth ranges." Possible differences in substrate etc. should be mentioned here.

A: Change was carried out

R1: 269: Rephrase: ". . . since (mega)faunal communities could be very different even between adjacent seamounts . . .

A: Ok

R1: 270: Which parameters? Name examples for depth-dependent parameters which drive faunal composition

A: *"parameters that vary with depth, such as temperature, oxygen concentration, substratum type, food availability, and pressure"* This was added. L302-304

R1: 278: Why would "elevated topography (peaks)" favour Porifera and Anthipataria? Name possible mechanism(s). By the way: Seamounts are per definitionem elevated topography

A: *Peaks are more exposed and appear thus more advantageous for filter feeders such as Porifera and Antipatharia. This is mentioned in the text.* L311

R1: 288-295: Do the authors mean that faunal density is negatively correlated with nodule coverage? This is in contrast to Vanreusel et al 2016, who found higher abundances at higher nodule coverage. So obviously in this study, the driver for the differences in faunal density was not nodule coverage, but probably organic input.

A: We searched for a possible explanation as to why APEI3 stood out and found that this difference, besides their more northward location under more oligotrophic waters (mentioned in L328-332), corresponded to a difference in nodule coverage. The patterns by which nodule coverage or densities influence the ecological patterns are still poorly understood. The nodule coverage data as mentioned in our ms originate from Table S1-1 from Vanreusel et al. 2016, and indeed in the body of text these authors reported higher epifaunal densities in areas with dense nodule coverage, reporting >25 versus ≤10 in sessile individuals per 100 m² for nodule rich and nodule free areas respectively. Nevertheless, if we compare the nodule coverage from Table S1 to figure 3 (both from Vanreusel et al. 2016) same patterns as those described in our study are observed, namely: Higher nodule coverage in APEI3 and lower densities both for sessile and mobile fauna. It is possible that Vanreusel et al. made their statement by looking at the license areas only and not included the APEI in this comparison. We chose to keep our statement, but elaborated on other possible reasons that could influence (e.g. POC) the patterns as observed (L328-331).

R1: 296: Grammar: neither - nor

A: Ok

R1: 319: Clearly distinguish between own data and data from literature by rephrasing, e.g. "Vanreusel et al. (2016) found that ophiuroids. . ."

A: Ok

R1: 322 : The available data cannot show a gradient, therefore it should read: ">50% less at seamounts compared to nodule fields"

A: This was corrected

R1: 331: ". . . studied here."

A: Ok

R1: 338: How can an uneven distribution (of holothuroids) affect composition?

A: Unevenly distributed organisms can give different perceptions in sampling. Organisms with a wide distribution range can, when unevenly distributed, be present/absent in adjacent sampling localities, thus resulting in different faunal composition for these sampling locality.

R1: 350: This is an isolated statement here - what does it imply?

A: We moved it to the beginning of the paragraph which now starts off as follows: "Stalked organisms, such as Crinoidea (Echinodermata) and Hexactinellida (except for Amphidiscophora, Porifera) rely on hard substrata for their attachment and are considered being among the most vulnerable organisms when mining is concerned." L381-383

R1: 352: And what about nodule-covered areas - did they host these taxa in Vanreusel's or this study? This paragraph is a bit confusing and should be re-sorted, also clearly distinguishing between own results and those of others.

A: This paragraph was re-written (taking into account the comments of the other reviewer as well) to make the distinction between our results and those from literature more easily: L391-402

R1: 361: "communities"

A: Ok

R1: 362: "... were more abundant. ..." - compared to what?

A: "... more abundant than on seamounts". This was added.

R1: 370: "...they are known to ..." Citation?

A: This statement was based on our personal observations. We have rephrased it and changed the structure of the paragraph as to convey our point more clearly: L411-417

R1: 372: Does the reference (Baco 2007) apply to both statements? I suggest to rephrase, e.g. "The exception. ...common on seamounts, as also reported in other studies (e.g., Baco. 2007)." Baco 2007 is not in the reference list!

A: Baco 2007 refers to the Scleractinia being common on seamounts. We have clarified this and added the reference for Baco 2007 to the reference list. L415-417

R1: 373: Insert: "... Enteropneusta which in this study were found only on seamounts, were. ..."

A: Ok

Figures

R1: Fig. 1: What does the "A" in the upper left corner mean?

A: Figure 1 underwent some change taking into account the comments from the other reviewer.

R1: Fig. 2: I suggest to add morphospecies "names" (as given in Fig. 3) to the examples.

A: Morphospecies names were added to the caption.

R1: Fig. 3: This figure should be simplified. Most of the morphospecies were observed in very low numbers, and in these cases differences between NF and seamounts are difficult to see in the figure and rather not relevant. I suggest to include in this figure only morphospecies and higher taxa with a substantial mean abundance (e.g., >0.5 ind./100 m per habitat type); other morphospecies could be summarised or omitted. By contrast, Table A1 should be extended and present the results for all morphospecies, not only higher taxa (see below).

A: Figure 3 was withheld because it was considered a key figure to show how different the presence/absence/abundance of the fauna varies between the two ecosystems, but the 3 parts were separated more clearly with different breaks at the X-axis, thus enhancing readability and interpretability. We also reorganised figure 3 to correspond to the order of A Table 1.

Table A1 now includes the morphospecies densities as well.

R1: Fig. 4: Axis labels are incomplete (units are missing). What does "exact" on the y-axis in panels a and c mean? And what is sample size (units?) in panels b and d? I guess that not sample size was used for the rarefaction curves, but accumulated number of observations. Caption is incomplete: What do the shaded areas in panels a and c and horizontal and vertical lines in panels b and D mean?

A: Sample size is the number of individuals observed (or number of observations as you will). Information was added to the caption and axis were renamed and/or clarified. Horizontal lines of the lower panels were omitted because they did not provide significant information for the interpretation of the figure.

R1: Fig. 5: What does "values are relative" mean? - percent (of what?)? This has to be explained.

A: Values are relative due to different transect lengths and differences in richness. This was changed to "Values are indicative rather than absolute due to different transect lengths and differences in richness."

R1: Fig. 6: y-axis incomplete, should include quantity and unit.

A: I am not sure what this is about, since the Y-axis is complete and has ind/100m as title.

R1: Fig. 7: Omit panel b).

A: See comment above

Tables

R1: Tab. 1: Be consistent with units: here, #obs/100 m is given, whereas throughout the text and in figures and in Tab. A1 the unit for density is ind/100 m.

A: The number (#) of observations is more of a methodological way to describe it, since it was used prior to the results. This could be changed.

R1: See also comment to Fig. 3. I suggest to list data for all distinguished morphospecies here and sums for higher taxa. It is irritating that densities for higher taxa (e.g., Holothuroidea) are given, but they do not include the identified morphospecies within that taxon. Not identified taxa should be clearly indicated, e.g. Holothuroidea indet., and they should sum up with the distinguished morphospecies to total Holothuroidea, etc. I also suggest to include absolute number of observations in addition to densities in this table. This would facilitate the evaluation whether, e.g., differences are based on a substantial number of individuals, or rely on just one or two specimens. The order of taxa in the table is not clear

A: Morphospecies densities (ind/100m) have been added to the table. We opted for densities to correspond to the data used in the manuscript. Absolute numbers are easy to deduce since the length of the transects are given in Table 1. The order of taxa in the table was reorganised

Reviewer 2:

Major comments

R2: 1) The appendix fig1 is a very important figure to place all the observations into context. I would move it from an appendix to a regular figure for the paper or as new panels in existing Fig. 1.

A: We have incorporated it as an extra panel in Figure 1.

R2: 2) The data presentation used to compare nodule transects to seamount transects should be refined. Right now figure 3 portrays averages of densities for fine taxa which, based on the finding of very low overlap between transects, means little - averaging a large number for one transect with low or zero numbers in other transects. Rather the data should be presented at broader taxonomic categories (as in Appendix table 1) with average (and standard deviation) densities and # of morphospecies. This would also follow the language in the results section better. Data on each fine morphospecies (level of taxonomy in Fig3) could be presented an an appendix and by transect.

A: We decided to add morphospecies to Table A1 (also taking into account the comments made by the other reviewer), which thus give information per transect, help to elucidate Fig. 3 and add the desired level of taxonomy. The order of morphospecies along the y-axis of figure 3 was also reorganised to correspond to that of Table A1. Figure 3 was thus withheld because it was considered a key figure to show how different the presence/absence/abundance of the fauna varies between the two ecosystems, but the 3 parts were separated more clearly and different breaks at the X-axis, thus enhancing readability and interpretability. This information was added in the figure caption as well.

R2: 3) Once data is pooled at higher taxa levels, statistical comparisons could be drawn to compare the average # morphospecies and average density between nodule and seamount transects.

A: There were no significant differences for the densities (T-tests for samples with unequal variance) per taxon (taxa pooled and tested: Actiniaria, Alcyonacea, Ceriantharia, Corallimorpharia, Antipatharia, Pennatulacea, Scleractinia, Zoantharia, Bryozoa, Asteroidea, Crinoidea, Echinoidea, Holothuroidea, Ophiuroidea, Porifera, Tunicata, Enteropneusta, Gastropoda. The pooled data were visualised as a histogram with st. dev. and added as an extra figure to this reply to the reviewer. We added the figure to the manuscript as Fig. A1 in the appendix and referred to it in Section 3.2.

The number of morphospecies per higher pooled taxon group proved more difficult, since often we know there are >1 or >2 morphospecies, but not the exact numbers. Using the minimum known number would be and under-estimation, which is why we chose not to test these pooled observations.

This comment was also taking into account for the overlapping morphospecies in Figure 6, for which standard deviation bars were added.

R2: 4) The authors conclude that the ratio of hard/soft habitat may explain some of the faunal trends they observe. Can't this ratio be determined from the transects? If possible add this metric to help explain faunal communities along or between the transects.

A: We have added information on the amount of hard substrata under the form of 3 categories: (1) Predominant soft substrata (<40% hard substrata), (2) mix or transition (40-60% hard substrata) and (3) predominant hard substrata (>60% hard substrata), annotated over 10m distance units. Very few

significant relationships were revealed (only for Mann Borgese ROV15), though backscatter data is currently being analysed to model the geomorphology along the transects in more detail and help reveal more details of fauna/substratum relationships, but it is out of the scope of this article.

The following paragraph was added in section 3.1 (also taking into account the other reviewer's suggestions): L205-2017

"About 57% of all sessile fauna was associated with predominantly hard substrata, followed by 31% on the mixed substrata. For the mobile taxa the pattern was less pronounced with 41 and 42% associated with predominantly hard and mixed soft/hard substrata respectively. The amount of predominantly hard and soft substrata were negatively correlated though, not significantly. This was due to the elevated amount of mixed hard/soft substrata featuring equal amounts 40-60%. Over all seamount transects pooled together, no taxa were significantly correlated with the amount of hard substrata, nor with soft substrata. When looking at the individual transects, no significant correlations were found between taxa and substrata for ROV02 or ROV04 or ROV09, most likely due to the equal distribution of the amount of hard/soft/mix substrata. In this perspective, ROV15 stood out, as it was dominated by predominantly hard substrata (56/%): For this transect, Pennatulacea were significantly negatively correlated with the amount of hard substrata and Zoantharia/Octocorralia were significantly and positively correlated with hard substrata, as were Ophiuroidea, Asteroidea, Crinoidea and Mollusca."

Specific comments

R2: line 82 - Explain why the north or northwestern flanks of the seamounts were chosen for the transects.

A: These flanks were chosen based on the positioning of the vessel and the predominant surface current in order to avoid the umbilical of the ROV to drift/being transported towards the vessel. Predominant currents in the CCZ are South-East oriented, allowing for an ROV positioning "downstream" of the vessel's location while visiting the north-northwestern seamount flanks. This was added in L81-84

R2: line - 97-99 - Provide the range of altitude, speed that were kept constant.

A: Target altitude was <2m above seafloor and travel speed ~0.2m/s, though interrupted by sampling actions, instrument check-ups, exploration, object avoidance (in the case of the uphill seamount transects) etc. This was added in the body of text. L99-102

R2: line 208-209 - this statement appears true when examining the higher taxa pooled data in appendix table 1. however in the cited fig 3, its hard to actually make this comparison because averages at finer taxonomic categories are highly variable due to lack of fine taxa overlap between transects.

A: This is amended by adding the morphospecies information in Table A1 and reorganising figure 3 to correspond to Table A1.

R2: line 267 - Start the sentence with, "Amongst the seamount transects,..."

A: Ok

R2: line 269 - The point of this sentence is not clear as it opposes the trend you find. Clarify.

A: This sentence was to point out that it could rather be depth influencing their similarity than their adjacent location. This part was changed (also taking into account the comments from the other reviewer) L298-303

R2: lines 296-306 - Nice to see a paragraph which lays out what future transecting should look like. The paragraph mentioned that wider depth ranges should be included and the data and literature certainly support that. Might it also be wise to have transects that move along contours so there are many replicate observations at a given depth, instead of conducting uphill transects? Adding a sentence or two addressing across slope vs with slope transecting would be worthwhile.

A: This is a valid observation which was added in L340-342

R2: line 308 - The sentence should be slightly reworded based on the authors findings to "Seamounts were shown to share FEW fauna with surrounding habitats...."

A: Based on the literature, fauna from seamounts tend to occur in neighbouring habitats quite easily, but that is not the case here. We rephrased the sentence to clarify this. L345-347

R2: Line 316 - this topic sentence needs to be improved. Rather than simply reiterating the results section can this paragraph be rewritten and a topic sentence created that summarizes the functional differences between taxa on seamounts vs nodules?. See literature by Rowden et al that look at functional variation of taxa on seamounts and neighboring areas. E.g. Rowden et al 2010 Marine Ecology

A: We added in topic sentences for several paragraphs and cleaned up the remainder of the body of text as to not repeat the results. L353-409

On a side note, functional traits of seamount and nodule field fauna are being investigated in a broader framework, extending beyond the feeding group and including life history, mobility etc.

R2: Table 1 - Add "SM:" before Mann Borgese

A: This was added to the table.

R2: figure 3 - given that there is so little overlap in the morphospecies between each sampled transect. Figure 3 is a bit hard to interpret. Error bars would help. Its great that the taxonomic diversity is presented but this might be better in the appendix.

A: We changed the Y-axis of the different parts of the figure 3 graph to make them more easily interpretable. See reply above. We have uploaded an extra figure linked to this reply to the reviewers that was included in the appendix, pooling the densities into larger taxa and referred to in section 3.2.

R2: Instead, appendix table 1 which presents higher taxa and # morphospecies might be the better data to show in the main paper. Here densities at higher levels can be better compared. Averaging the abundances across the transects seems ill advised given the differences observed between each one.

A: We have added the morphospecies' densities to the appendix table, which also clarifies the taxonomy and figure 3.

R2: Figure 6 - It is not clear what data is being presented here. Are these only taxa present on both seamounts and nodules? Please clarify in the figure caption

A: Yes, indeed. We clarified the caption.

1 Are seamounts refuge areas for fauna from polymetallic 2 nodule fields?

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18 Running title: Seamounts as refuge areas for nodule fauna

19 Six keywords: megafauna, seamounts, nodule fields, image analysis, deep sea, mining

20 Abstract

21 Seamounts are abundant and prominent features on the deep-sea floor and intersperse with the
22 nodule fields of the Clarion-Clipperton Fracture Zone (CCZ). There is a particular interest in
23 characterising the fauna inhabiting seamounts in the CCZ because they are the only other ecosystem
24 in the region to provide hard substrata besides the abundant nodules on the soft sediment abyssal
25 plains. It has been hypothesised that seamounts could provide refuge for organisms during deep-sea
26 mining actions or that they could play a role in the (re-)colonisation of the disturbed nodule fields.
27 This hypothesis is tested by analysing video transects in both ecosystems, assessing megafauna
28 composition and abundance.

29 Nine video transects (ROV dives) from two different license areas and one Area of Particular
30 Environmental Interest in the eastern CCZ were analysed. Four of these transects were carried out as
31 exploratory dives on four different seamounts in order to gain first insights in megafauna
32 composition. The five other dives were carried out in the neighbouring nodule fields in the same
33 areas. Variation in community composition observed among and along the video transects was high,
34 with little morphospecies overlap on intra-ecosystem transects. Despite these observations of
35 considerable faunal variations within each ecosystem, differences between seamounts and nodule
36 fields prevailed, showing significantly different species associations characterising them, thus
37 questioning their use as a possible refuge area.

1. Introduction

Seamounts are abundant and prominent features on the deep-sea floor (Wessel et al., 2010). They are common in all the world's oceans, occurring in higher abundances around mid-ocean ridges, island-arc convergent areas, and above upwelling mantle plumes (Kitchingman et al., 2007). Seamounts are defined as isolated sub-surface topographic feature, usually of volcanic origin, of significant height above the seafloor (International Seabed Authority (ISA), 2019). They are generally isolated, typically cone shaped undersea mountains rising relatively steeply at least several hundred meters from the deep-sea floor. Seamounts comprise a unique deep-sea environment, characterised by substantially enhanced currents and a fauna that is dominated by suspension feeders, such as corals (Rogers, 2018). They represent hard substrata in the otherwise soft sediment deep sea and can thus be considered habitat islands (Beaulieu, 2001). Given the growing evidence that seamounts differ substantially across a range of spatial scales, the concept of seamounts as a single, relatively well-defined habitat type is outdated (Clark et al., 2012). Depth and substrate type are key elements in determining the composition and distribution of benthic fauna on seamounts, while location is likely the subsequent most important driver of faunal composition and distribution patterns (e.g. Tittensor et al., 2009). Connectivity varies substantially between seamounts, resulting in the presence of taxa with very localised to very wide distributions (Clark et al., 2010).

The Clarion-Clipperton Fracture Zone (CCZ), in the equatorial eastern Pacific Ocean, is most known for its extensive polymetallic nodule fields that will potentially be mined in the future. In this area, nodules represent the most common hard substrate on the soft-sediment abyssal plains, and many organisms rely on them for survival (Vanreusel et al., 2016). Removal of hard substrate through mining actions will impact all these organisms, which were estimated at about 50% of all megafaunal species in the CCZ (Amon et al., 2016). Nodule fields in the CCZ are interspersed by seamounts (Wedding et al., 2013), the only feature offering hard substrata besides the nodules. Based on this feature/characteristic, it has been hypothesised that seamounts could provide refuge for organisms during deep-sea mining activities or that seamounts could play a role in the (re-)colonisation of the disturbed nodule fields. Whether or not this is true may have important implications for management of the impacts of polymetallic nodule mining in the CCZ. However, knowledge on the biodiversity inhabiting seamounts in this region is currently lacking.

The objectives of the current study were twofold: (i) Provide first insights in seamount megafauna within the CCZ, (ii) Compare the benthic fauna inhabiting seamounts and nodule fields in the eastern CCZ. Since this is the first time the seamounts at the eastern CCZ were visited, a separate section is dedicated to describe these first insights.

2. Material and Methods

2.1. Study site and data

During the SO239 ECORESPONSE cruise in 2015 (Martinez Arbizu and Haeckel, 2015), four seamounts were visited for the first time within two different license areas and one area of particular environmental interest (APEI) within the Clarion-Clipperton Fracture zone (CCZ) (Table 1). Nodule fields within the same license areas were visited and sampled as well. Video imagery and faunal samples were collected by a Remotely Operated Vehicle (ROV Kiel 6000 (GEOMAR), equipped with a high definition Kongsberg OE14-500 camera).

Seamount transects were carried out uphill towards the summit resulting in a depth gradient along the transect, ~~whilst nodule transects featured rather stable depth ranges~~ (Table 1). The four seamount transects were characterised by different depth ranges and lengths and, due to the vessel's positioning and the predominant South-East surface currents, but were all situated-carried out downstream, on the north to north-western flanks of the seamounts (Table 1 and Fig. S1). The names of the seamounts used here, Rüppel and Senckenberg (BGR, German License area), Heip (GSR, Belgian License area) and Mann Borgese (APEI3), are the ones agreed upon by the scientist during the ECORESPONSE cruise (Martinez Arbizu and Haeckel, 2015), pending incorporation of these names in the GEBCO gazetteer. The seamounts differed in shape and size with Senckenberg and Heip being a sea-mountain range, while Rüppel and Mann Borgese were more isolated, stand-alone seamounts (Fig. S1). Nodule field dives were carried out on relatively flat surfaces (maximum depth range covered during a dive or transect was 30m difference, Table 1) and were referred to by the dive number and license area. The five nodule transects were all located between 4000-5000m depth and the transects differed in length between dives as well (Table 1). Within the same license area distance between different transects was 16 to 60km, while distance between license areas added up to several hundreds of kilometres (minimum ~700kms BGR – GSR, Fig. 1).

Investigated areas were restricted to the eastern part of the CCZ with APEI3 being the most north- and westward bound area. The optical resolution of the camera enabled reliable identification of organisms larger than 3 cm (Martinez Arbizu and Haeckel, 2015). The combination of exploration and opportunistic sampling restricted a systematic image collection. Target ROV travelling altitude was <2m and travelling- speed was ~0.2m/s and camera zoom which, along with the camera zoom, were kept constant whenever possible. Due to the explorative nature of the dives, while the pan and tilt of the ROV camera were not kept constant.

2.2. Video analysis and statistics

All videos were annotated to the lowest taxonomic level possible. The number of morphospecies, defined as morphologically different organisms within the lowest taxonomic group identified, were assessed. Identifications were double checked with scientists working in the same area as well as taxonomic experts and comprise different taxonomic levels (e.g. Genus, Family). Those ~~ID's~~ identifications restricted to higher taxon groups (Family, Class, etc.) and for which it was impossible to attribute a morphospecies, are referred to as taxa and are likely to morphologically differ between transects. Xenophyophores, living on the soft sediment deep-sea floor, were less prominently present at seamounts than at nodule fields and were not quantified. Fish (Actinopterygii), ~~crustaceans~~ Crustacea (Nematocarinidae, Aristidae, Peracarida) and Polychaeta were quantified but left out of the comparing analysis due to their lack of representativity and possible attraction due to ROV lights. The same was done for jellyfish and other doubtful identification ~~ID's~~ that could not be confidently assigned to a higher taxonomic group (Table A1). A subset of the nodule field transects from BGR, GSR and APEI3 was presented by Vanreusel et al. (2016), corresponding to 44% of what was studied here and limited organism identification to a higher taxonomic level (Order (e.g. Alcyonacea) or Class (e.g. Ophiuroidea)). In our study, the entire transects (100%) were annotated to morphospecies level, allowing a detailed comparison between seamounts and nodule fields.

Three categories of substratum types were distinguished: (1) Predominant soft substrata (<40% hard substrata), (2) mix or transition (between 40 and 60% hard substrata) and (3) predominant hard substrata (>60% hard substrata), were annotated per 10m distance units based on the video footage and tested for correlations with taxonomic abundances.

ROV transects on the seamounts were carried out as exploratory dives. Sampling strategy both at seamounts and nodule fields combined video and sampling or specimen collection. Due to varying altitude of the ROV and the use of camera pan, tilt and zoom, it was not possible to use surface coverage as a standardisation measure. We used video transect length instead. For the transect length calculation for each dive, we omitted all parts of the video footage in which the ROV was at an altitude of >10m, or sections where the ROV was not visualising the seafloor (e.g. during transiting or inspecting ROV parts or instruments). Visualisation of ancient disturbance tracks were omitted as well, as these fell out of the scope of the article. Faunal densities were calculated as the number of observations per 100m, in order to compensate for time spent collecting samples and differing transect lengths. Statistical testing was carried out in R (R core team, 2018) and the Non-metric multidimensional scaling analysis (NMDS) was based on Bray-Curtis dissimilarity and used the vegan package (Oksanen et al., 2018). The Kendall's coefficient of concordance (W) was calculated to identify significantly associated groups of species, based on correlations and permutations (Legendre, 2005).

3. Results

About 80% of all taxa observed across the two adjacent ecosystems, could be identified to a morphospecies level. A total of 252 taxa were observed across the two adjacent ecosystems, of which 207 (or 82%) could be identified to a morphospecies level. At a first view, morphospecies revealed to be quite different between seamounts and nodule fields (Fig. 2). While the number of faunal observations at the seamount transects were within similar ranges (33.1-40.734-42 ind./100m), those at the nodule transects featured both highest and lowest values (7.596.3-89.2367.5 ind/100m) (Table 1). The lowest number of faunal observations were done at the two APEI3 nodule transects (ROV13 and 14) and highest at the GSR nodule transect ROV08ROV10. What follows is a first description of eastern CCZ seamount mega fauna (3.1.) and a detailed comparison with the neighbouring nodule fields (3.2.)

3.1. Insights in CCZ seamount mega fauna

The most abundant and diverse (most morphospecies) taxa at the seamount transects comprised Echinodermata (Asteroidea, Crinoidea, Holothuroidea and Ophiuroidea), Anthozoa (Actiniaria, Alcyonacea, Pennatulacea, Scleractinia) and Porifera (Hexactinellida) (Table A1, Fig. 3). Keeping in mind the limitation of the video sampling, differences among the benthic seamount community composition are described here.

The transect at Mann Borgese (APEI3) was characterised by high densities of Antipatharia, more specifically Antipathidae (4918.451 ind./100m), and solitary Scleractinia (8.447.9 ind./100m) (Table A1, Fig. 3). Antipathidae observations were mostly grouped at the end of the video transect, i.e. at the summit. Densities of both Antipatharia and Scleractinia were much lower on the other seamount transects (<0.21 ind./100m) with Scleractinia being absent from Heip and Senckenberg transects. Alcyonacea corals were observed on all seamount transects. Isididae were found at Senckenberg and Heip transects, and one individual from Chrysogorgiidae was observed at the latter as well. Varying

163 numbers of Primnoidae were observed on all transects (Table A1). High abundances of Pennatulacea
164 were observed at Senckenberg (3.648 ind./100m), representing about 2628% of sessile fauna
165 annotations for this transect.

166 Enteropneusta were only observed on Rüppel and Senckenberg transects in the BGR area,
167 represented by two different morphospecies, namely *Yoda* morphospecies (Torquaratoridae) at
168 Rüppel and *Saxipendium* morphospecies (Harrimaniidae) at Senckenberg.

169 Highest Polychaeta densities were observed at Heip transect in the GSR area, which was mainly due
170 to high densities of free-swimming Acrocirridae (4.2 ind./100m vs. 0.256 and 0.32 ind./100m in BGR
171 and 0.11 ind./100m in APEI3 area). Free-swimming Acrocirridae were observed in very high densities
172 as well (Table A1). Aphroditidae polychaetes were only present at the BGR transects (3.0.2
173 ind./100m (corresponding to 3 individuals along the transect) at Rüppel and 0.04 ind/100m (or 1
174 individual along the transect) at Senckenberg) (Table A1).

175 Porifera densities were highest at the Heip transect (3.7.5 ind./100m), followed by Rüppel (3.52.72
176 ind./100m), Senckenberg (1.91.92 ind./100m) and lastly Mann Borgese (1.80.68 ind./100m). Six
177 Porifera families were annotated featuring >7 to >10 morphospecies (Fig. 3, Table A1). Cladorhizidae
178 (two individuals) were only observed on Heip transect, and one *Poliopogon* sp. (Pheronematidae)
179 was observed at Mann Borgese transect. Rossellidae gen. sp. nov. was present on three seamount
180 transects, exception being Mann Borgese.

181 Overall Echinodermata densities were highest at Senckenberg seamount (1717.64 ind./100m),
182 followed by Rüppel (1210.24 ind./100m) (Table A1, Fig. 3), both adding up to 6047% of all image
183 annotations for these transects. The number of morphospecies for all echinoderm taxa (Asteroidea,
184 Echinoidea, Holothuroidea and Crinoidea) was also highest at these 2 seamounts in the BGR area
185 (Fig. 3). For comparison, echinoderms at Heip (11.68.10 ind./100m) and Mann Borgese transects
186 (3.08.3 ind./100m) were responsible for 4932% and 8.82% of observations respectively. Crinoidea
187 densities were highest at Senckenberg (4.322 ind./100m), while Holothuroidea were most abundant
188 at Rüppel (5.24.4 ind./100m). The holothuroid families of Elpidiidae and Laetmogonidae were only
189 observed at Senckenberg and Rüppel (BGR). Psychropotid and synallactid holothuroids were
190 observed on all seamounts, represented by different morphospecies. Deimatid holothuroids were
191 not observed on Mann Borgese, but were present in the three other seamount transects, again with
192 different morphospecies and densities. Velatid Asteroidea were only observed at Senckenberg and
193 Rüppel (BGR), while Brisingida and Paxillosida were observed on all four seamounts.
194 Aspidodiadematid echinoids were absent from the Heip transect and urchinid echinoids were
195 absent from the Mann Borgese transect.

196 A species accumulation curve (Fig. 4a) confirmed the limitations of the restricted and exploratory
197 nature of the sampling as no asymptote was reached. The rarefaction curves (Fig. 4b) showed that
198 the transects with the most faunal observations, which corresponded here to the longer transects,
199 were more diverse. However, at smaller sample sizes curves did not cross, thus maintaining the
200 differences observed at higher sample sizes with the Senckenberg transect (ROV04) as most diverse
201 followed by Rüppel (ROV02) (both BGR). The video transect carried out at Mann Borgese (ROV15,
202 APEI3) was the least diverse.

A comparison of all morphospecies observed along the 4 transects was presented in a Venn diagram (Fig. 5a). Each seamount transect was characterised by a highest number of unique morphospecies, only observed on the transect in question and not elsewhere. Only ~~two~~three morphospecies were present in all seamount transects, namely *Ceriantharia msp. 2*, a small red galatheid crab and a foliose sponge. Highest number of overlapping morphospecies (~~#1416~~) was observed between Rüppel and Senckenberg, both in the BGR area (Fig. 5a). Mann Borgese showed the smallest degree of overlap with the other transects (Fig. 5a).

About 57% of all sessile fauna was associated with predominantly hard substrata, followed by 31% on the mixed substrata. For the mobile taxa, the pattern was less pronounced with 41 and 42% associated with predominantly hard and mixed hard/soft substrata respectively. The amount of predominantly hard and soft substrata was negatively correlated, though not significantly. This was due to the equal amounts (40-60%) of mixed hard/soft substrata. Over all seamount transects pooled together, no taxa were significantly correlated with the amount of hard substrata, nor with soft substrata. When looking at the individual transects, no significant correlations were found between taxa and substrata for ROV02 or ROV04 or ROV09, most likely due to the equal distribution of the amount of hard/soft/mix substrata. In this perspective, ROV15 stood out, as it was dominated by predominantly hard substrata (56%). For this transect, Pennatulacea were significantly negatively correlated with the amount of hard substrata and Zoantharia/Octocorralia were significantly and positively correlated with hard substrata, as were Ophiuroidea, Asteroidea, Crinoidea and Mollusca.

Due to the limited sample size, the representativity of the observed biological patterns remains to be corroborated by a more elaborate sampling strategy.

3.2. Comparison of seamount and nodule field faunal composition and variation

The faunal composition and richness (number of morphospecies in higher taxonomic groups) of the nodule transects can be consulted in Fig. 3 and Table A1, respectively. In concordance with the seamount transect, the species accumulation curve of the nodule transects did not reach an asymptote either (Fig. 4c). The rarefaction curves showed that the relations among transects were less ~~linear~~straightforward for the nodule transects versus the seamount ones and did cross at smaller sample sizes (<100 individuals, Fig. 4d). ROV13 and ROV14 transects (both APEI3) were the longest in distance travelled (Table 1) but featured less faunal observations. At small sample sizes, the richness at ROV13 and 14 was highest. ROV08 and ROV10 (both GSR) showed parallel curves with ROV08 being more diverse (Fig. 4d).

A venn diagram showing the morphospecies overlap among the nodule transects showed a total of ~~5~~6 species re-occurring on all 5 transects (Fig. 5b). These were: Munnopsidae msp. 1 (Isopoda, Crustacea), Actiniaria msp.7 (Cnidaria), Ophiuroidea msp. 6 (Echinodermata), *Holascus* sp. and *Hyalonema* sp. (Hexactinellida, Porifera). There was a high number of unique morphospecies for each transect, though not as high as for the seamount transects (Fig. 5). ROV13 and 14 (both APEI3) showed ~~littlest-least~~ overlap with the other transects, which is similar to what was observed at the seamounts.

Observations and quantifications of morphospecies confirmed the high degree of dissimilarity between the two neighbouring ecosystems. Porifera, Ophiuroidea (Echinodermata), Actiniaria and Alcyonacea (Cnidaria) were more abundant at nodule fields (Fig. 3). These taxonomic groups were

also most diverse on nodule fields (i.e. highest number of morphospecies), exception being the Alcyonacea which featured more morphospecies on the seamounts (12 to 8 morphospecies for seamounts and nodule fields respectively) (Fig. 3). Of all Porifera, Cladhorizidae were more diverse at nodule fields than at seamounts (7 to 1 morphospecies, respectively).

There were only 21 morphospecies (10%) that were observed both on seamounts and nodule fields (Fig. 6). While this subset of morphospecies occurred in both ecosystems, they did so in very different densities, i.e. very abundant in one ecosystem and very low in abundance in the other: examples are Galatheididae small red msp. (Decapoda, Crustacea), *Synallactes* white msp. (Holothuroidea), Ophiuroidea msp. 5 and 6, Comatulida msp. 1 (Crinoidea), *Hyalonema* sp. and *Hyalostylus* sp. (both Hexactinellida, Porifera) (Fig. 6).

Three Ophiuroidea morphospecies were present at both seamounts and nodule fields (Fig. 2, 83 and 6). The majority of the very abundant Ophiuroidea ophiuroids observed at the CCZ seamounts were small and situated on hard substrata (morphospecies 5-), ~~While while the most-most of the Ophiuroidea at nodule fields (abundant morphospecies at nodule fields (including~~ morphospecies 6) ~~were as mostly~~ observed on the soft sediments ~~of the nodule transects. This m~~ Morphospecies 6 was only rarely observed on the seamounts (Fig. 3). Another easily recognisable morphospecies was found on Porifera, corals and animal stalks and was more abundant at seamounts than at nodule fields (morphospecies 4) (Fig. 2 and 3).

Crinoidea, Asteroidea (both Echinodermata) and Antipatharia (Cnidaria) were more abundant on the seamounts (Fig. A1). This coincided with a higher diversity for Asteroidea and Antipatharia on the seamounts as well. Crinoidea diversity was similar (5 to 4 morphospecies comparing seamounts to nodule fields). Holothuroidea occurred in similar densities in both ecosystems (Fig. A1, though they were characterised by different morphospecies (Fig. 3). Overall densities of Echinoidea were ~~highest comparable between seamounts and~~ at nodule fields, though ~~for the nodule fields this -this was~~ mostly due to one very abundant morphospecies, namely Aspidodiadematidae msp 1, ~~which was absent at the seamounts~~ (Fig. 3). Besides this ~~one very abundant morphospecies, which was only present at nodule fields, Echinoidea echinoids showed higher densities at seamounts and~~ were more diverse ~~at seamounts~~ (11 morphospecies vs. 5 at nodule fields).

There was no morphospecies overlap for Tunicata, Antipatharia, and Actiniaria. Alcyonacea, Ceriantharia, Corallimorphidae and Crinoidea only shared 1 morphospecies between seamounts and nodule fields, namely *Callozostxon* cf. *bayeri*, Ceriantharia msp. 2, *Corallimorphus* msp. 2 and Comatulida msp. 1 respectively (Fig. 6).

There were no observations of Enteropneusta, Scleractinia and Zoantharia (Cnidaria), Aphroditidae (Polychaeta) or holothuroid Deimatidae at the nodule fields transects (Table A1, Fig. A1). While Actinopterygii were left out of the analysis, it should be noted that fish observations were more abundant and diverse at the seamounts than on the nodule fields (Table A1).

There was quite some faunal variation observed among the video transects ~~carried out in the different license areas of~~, both for seamounts and nodule fields (see fig. 5). The (dis)similarities were analysed by a nMDS analysis, which grouped the 9 different video transects based on their taxonomic composition. Despite the large intra-ecosystem variation, they pooled in two distinct groups separating the nodule fields from the seamounts (Fig. 7a). Within each group, BSR and GSR

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287 transects were more similar to one another both for seamounts and nodule fields, whilst APEI3
288 transects stood out more.

289 The Kendall's coefficient of concordance (W, Legendre, 2005) corroborated the existence of two
290 significantly different species associations, whose composition corresponded to the fauna
291 characterising the nodule fields ($W=0.20$, $p<0.001$, after 999 permutations) and the seamounts
292 ($W=0.30$, $p<0.001$, after 999 permutations).

293 Depth was fitted as a vector on top of the nMDS plot (Fig. 7b) and showed that the discrepancy in
294 faunal composition between the two ecosystems also corresponded to a difference in depth, with
295 the nodule transects all being situated below the 4000m isobath and the seamount transects ranging
296 from 1650 to >3500m (Fig. 7b).

297 4. Discussion

298 4.1. Intra-ecosystem faunal variation

299 Community composition varied markedly within seamounts and nodule fields. The limited sampling
300 ($n=9$ transects), ~~across at different localities-locations and additionally, for the seamounts, different~~
301 ~~depth ranges and for the seamounts different depth gradients~~, precluded any general conclusions on
302 quantifications of biodiversity *per se*. However, taking this into account, it was also the first time
303 seamounts were visited in the area, thus granting first insights in the fauna inhabiting these
304 seamounts and allowing a first comparison with nodule faunal composition.

305 ~~The two BGR seamount transects were most similar in faunal composition, followed by the Heip~~
306 ~~seamount transect (GSR). These seamount video transects were characterised by more similar depth~~
307 ~~ranges, and the two BGR transects were also geographically closest to each other. For seamounts,~~
308 ~~distance separating them might be a less determining factor than depth since (mega)faunal~~
309 ~~communities can be very different even between adjacent seamounts (Schlacher et al., 2014;~~
310 ~~Boschen et al., 2015). Overall, parameters that vary with depth, such as temperature, oxygen~~
311 ~~concentration, substratum type, food availability, and pressure are considered major drivers of~~
312 ~~species composition on seamounts (Clark et al., 2010; McClain et al., 2010). The quantification of the~~
313 ~~amount of hard and soft substrata was not distinctive enough to explain differences observed here.~~
314 ~~The difference in depth could also explain the higher dissimilarity with Mann Borgese (APEI3) who~~
315 ~~featured the shallowest transect and summit, which was dominated by Antipatharia. The two BGR~~
316 ~~seamount video transects were geographically closest to each other and were most similar in faunal~~
317 ~~composition but also in depth. For seamounts, distance separating them might be a less determining~~
318 ~~factor since adjacent seamounts were shown to be very different in inhabiting fauna (Schlacher et~~
319 ~~al., 2014; Boschen et al., 2015). Overall, parameters that vary with depth are considered major~~
320 ~~drivers of species composition on seamounts (Clark et al., 2010; McClain et al., 2010). Depth could~~
321 ~~be explanatory for the higher degree of similarity of these two BGR seamount transects and, to a~~
322 ~~lesser extent, the Heip seamount transect (GSR). Similarly, it could explain the higher dissimilarity~~
323 ~~with Mann Borgese (APEI3) who featured the shallowest transect and summit, which was dominated~~
324 ~~by Antipatharia. Antipatharians were previously reported to be more dominant towards peaks as~~
325 ~~compared to mid-slopes at corresponding depths (Genin et al., 1986). Based on their filter-feeding~~
326 ~~strategy, Porifera (except carnivorous Cladorhizidae), were also thought to benefit from elevated~~
327 ~~topography (peaks) or exposed substrata in analogy to corals (Genin et al., 1986; Clark et al., 2010),~~
328 ~~though no such pattern was apparent here. Porifera are notoriously difficult to identify based on~~
329 ~~imagery. Although the sampled individuals allowed some identifications to genus or species level~~

330 (Kersken et al., 2018a and b), identifications remained hard to extrapolate across the different video
331 transects. Generally, as in our study, seamount summits have been more intensively sampled
332 (Stocks, 2009) although the little work done at seamount bases and deep slopes indicated that these
333 areas support distinct assemblages (Baco, 2007).

334 Among the nodule transects a considerable amount of variation in faunal composition was observed
335 (this study, Vanreusel et al., 2016). The two APEI3 nodule transects (ROV13 and 14) stood out in
336 faunal composition, diversity and in low number of faunal observations.~~Among the nodule transects~~
337 ~~a considerable amount of variation in faunal composition was observed (this study, Vanreusel et al.,~~
338 ~~2016). The two APEI3 nodule transects (ROV13 and 14) stood out, both in the low number of faunal~~
339 ~~observations, faunal composition and diversity.~~ They were also the only two transects situated
340 below the 4500m isobaths. But rather than depth, the nodule coverage ~~was thought to be more of~~
341 ~~may be considered an important~~ driving factor, since the density of nodule megafauna was shown
342 to vary with nodule size and density/coverage (Stoyanova, 2012; Vanreusel et al., 2016, Simon-Llédó
343 et al., 2019). Here as well, the APEI3 transects were characterised by a high nodule coverage (~40-
344 88%, Vanreusel et al., 2016), whereas the BGR and GSR nodule transects (ROV3 and ROV 8 + 10,
345 respectively) had a nodule coverage <30% and were also more similar in faunal composition
346 (Vanreusel et al., 2016). Other factors that could be at play are ~~the~~ more oligotrophic surface
347 waters of the northern CCZ ~~were proposed to be~~ which could be the cause of the overall lower faunal
348 densities at APEI3 nodule fields (Vanreusel et al., 2016). Volz et al. (2018) corroborated this, with the
349 location of the APEI3 site in the proximity of the carbon-starved North Pacific gyre being
350 characterised by a reduced POC-flux quantified to being 22-46% lower than the GSR and BGR areas
351 respectively.

352 The species accumulation curves showed that no asymptote was reached ~~not-neither~~ at seamounts,
353 nor at nodule fields. Consequently, longer transect lengths might be necessary to representatively
354 quantify and assess megafauna density and diversity (Simon-Llédó et al., 2019). In addition, for a first
355 in-depth description and assessment of seamount fauna composition, one video transect is
356 insufficient to describe the diversity and shifts in faunal assemblages of the surveyed seamounts.
357 Rather, an ampler imaging strategy should be developed, with a minimum transect length exceeding
358 1000ms (Simon-Llédó et al., 2019) and replicate transects carried out on different faces of the
359 seamount, on slopes with varying degree of exposure to currents and different substrate types.
360 Wider depth ranges should be taken into account as well. Alternatively, across slope transects,
361 following depth contours should be considered as these could provide observation replicates for a
362 given depth. Despite its limitations, this study grants first insights in the seamount inhabiting
363 megafauna of the eastern CCZ and an important first comparison with nodule fauna.

364 4.2. Faunal (dis)similarities between seamounts and nodule fields

365 In other areas, seamounts were shown to share fauna with surrounding habitats (Clark et al., 2010)
366 and could thus potentially serve as source populations for neighbouring environments (McClain et
367 al., 2009)~~Seamounts were shown to share fauna with surrounding habitats (Clark et al., 2010) and~~
368 ~~potentially serve as source populations for neighbouring environments (McClain et al., 2009).~~ While
369 generally few species seemed restricted to seamounts only (Clark et al., 2010), in this study,
370 morphospecies revealed to be quite different between seamounts and nodule fields with little
371 overlap between both. Despite the high degree of variation observed among all the video transects,
372 these grouped into two distinctly separate clusters, separating nodule from seamount transects. The

373 few overlapping morphospecies did occur in different densities in each ecosystem, implying a
374 different role or importance in the ecological community and its functioning.

375 Overall, nodule fields showed higher faunal densities than seamounts. Shifts in density patterns
376 between nodule fields and seamounts were more evident in a number of taxa, where the variety of
377 morphospecies and feeding strategy within each group was likely to be at play. An example of this
378 are the Echinodermata, which include Asteroidea (predators and filter feeders (Brisingida)),
379 Crinoidea (filter feeders), Echinoidea (deposit feeders), Holothuroidea (deposit feeders) and
380 Ophiuroidea (omnivores).

381 Taxa contributing to the differences between the two ecosystems are discussed here. Ophiuroids
382 were more abundant on the nodule fields. Asteroids-Asteroidea were more abundant on seamounts
383 and both Echinoidea and Asteroidea were more diverse in this ecosystem as well, and echinoids (with
384 exception of one very abundant morphospecies at the nodule fields) were both more abundant and
385 diverse on the seamounts. Ophiuroidea were most abundant on the nodule fields (ratio 7 to 1 when
386 compared to seamounts). Both ophiuroid and echinoids were shown to be present in both nodule-
387 rich and -free areas, though their densities decreased more than 50% comparing the former to the
388 latter (Vanreusel et al., 2016). Despite the abundance of hard substrata at seamounts, this was true
389 for the ophiuroid densities observed here (>50% decrease from nodule fields to seamounts), but not
390 for the echinoids, where differences in overall density between ecosystems were less pronounced.
391 Ophiuroids did not show high levels of richness or endemism on seamounts (O'Hara, 2007). Same
392 ophiuroid morphospecies were present at seamounts and nodule fields but in very different
393 abundances and they were observed on different substrata types showed preference for different
394 substrata (at nodule fields on soft sediment (morphospecies 6), at seamounts on hard substrata
395 (morphospecies 5)), which appeared to correspond to indicates different lifestyles, feeding behaviour
396 and corresponding dietary specialisations (Persons and Gage, 1984). Previously it was already
397 demonstrated that Ophiuroidea did not show high levels of richness or endemism on seamounts
398 (O'Hara, 2007). At nodule fields Ophiuroidea Ophiuroids were often observed in association with
399 xenophyophores at nodule fields (Amon et al., 2016, this study) and a similar observation was done
400 at east Pacific seamounts off Mexico (Levin et al., 1986), though no such associations were observed
401 on the seamounts studied here.

402 Holothuroidea densities were thought to possibly decrease when less soft sediment was available
403 since they feed mainly on the upper layers of the soft-bottom sediment (Bluhm and Gebruk, 1999).
404 No significant link was established between holothuroid densities and the amount of hard substrata
405 in this study, but their community composition varied distinctly between nodule fields and
406 seamounts with more families being observed at the latter. Additionally, at the seamounts, many
407 holothurians were observed on top of rocks, possibly reflecting different feeding strategies and
408 explaining the observations of different morphospecies. Geographical variations, different bottom
409 topography, differences in nodule coverages and sizes and/or an uneven distribution of holothurians
410 on the sea floor were thought to play a role in holothuroid community composition (Bluhm and
411 Gebruk, 1999). On the other hand, variability in deep-sea holothuroid abundance was proposed to
412 depend primarily on depth and distance from continents (see Billet, 1991 for a review).

413 Holothuroid composition varied distinctly between nodule fields and seamounts with more families
414 being observed at the latter. Many holothurians feed on the upper layers of the soft-bottom

sediment (Bluhm and Gebruk, 1999), suggesting that their numbers would decrease when there is less sediment available. However, at the seamounts, many holothurians were observed on top of rocks, possibly reflecting different feeding strategies and explaining the observations of different morphospecies. Geographical variations, different bottom topography, differences in nodule coverages and sizes and/or an uneven distribution of holothurians on the sea floor were thought to play a role in holothuroid community composition (Bluhm and Gebruk, 1999). On the other hand, variability in deep-sea holothuroid abundance was proposed to depend primarily on depth and distance from continents (see Billet, 1991 for a review).

Stalked organisms, such as Crinoidea (Echinodermata) and Hexactinellida (except for Amphidiscophora, Porifera) rely on hard substrata for their attachment and are considered being among the most vulnerable organisms when mining is concerned. Crinoidea were proportionally more abundant on seamounts, possibly because hard substrata were less limiting than in the nodule fields. Porifera proportions-densities (stalked and non-stalked) varied among all analysed transects, revealing no particular trends in abundance. However, the species composition of deep-sea glass sponge communities from seamounts and polymetallic nodule fields was distinctly different. Polymetallic nodule field communities were predominated by widely-distributed genera such as *Caulophacus* and *Hyalonema*, whereas seamount communities seemed to have a rather unique composition represented by genera like *Saccocalyx*. Stalked organisms are considered being among the most vulnerable organisms when mining is concerned.

Corals were generally considered to be more abundant on seamounts than adjacent areas, due to their ability to feed on a variety of planktonic or detritus sources suspended in the water column (Rowden et al., 2010). In this study, the Alcyonacea densities were lower on the seamounts than on the nodule transects. The majority of Alcyonacea morphospecies of the seamounts did not occur on the nodule fields and vice versa, with exception of *Callozostron cf. bayeri* which was also present at the nodule fields but in very low densities (1/10 of those observed at seamounts). The Antipatharia were most abundant at the Mann Borgese seamount (APEI3) compared to all other transects. The depth difference of more than 3000m between this particular seamount and the nodule fields could explain the abundance in Antipatharia which were shown to be more abundant at lower depths (Genin et al., 1986). Additional presence of Pennatulacea at seamounts, a taxon that was virtually absent from the nodule field transects and that appeared more linked to predominant soft substrata at seamounts, resulted in completely distinct coral communities for each ecosystem.

While alcyonacean and antipatharian corals were virtually absent from nodule-free areas (Vanreusel et al., 2016), this was not the case for the seamounts (although Alcyonacea densities were lower than on nodules). Depth difference added up to more than 3000m between Mann Borgese seamount (APEI3) and the nodule transects, which could explain the difference in Antipatharia which were more abundant at lower depths (Genin et al., 1986). The antipatharian and alcyonacean morphospecies that were abundant on the seamounts did not occur on the nodule fields and vice versa, with exception of *Callozostron cf. bayeri* which was present at the nodule fields but in very low densities (1/10 of those observed at seamounts). Additional presence of Pennatulacea, which were virtually absent from the nodule field transects, resulted in a completely distinct coral community for both ecosystems.

Actiniaria were ~~more abundant on nodule fields. It was~~ denominated the second most common group at CCZ nodule fields, after the xenophyophores (~~Kamenskaya et al., 2015~~). ~~(Kamenskaya et al., 2015) and, in our study, were also more abundant on nodule fields than on seamounts.~~ Depending on the species and feeding strategy, the ratio hard/soft substrata and their preference for either one could play a role. Since morphospecies were distinct between seamounts and nodule fields, their role in the respective communities are likely to differ as well. Combinations of deposit feeding and predatory behaviour in Actiniaria have been observed, as well as burrowing activity, preference for attachment to hard substrata and exposure to currents (Durden et al., 2015a; Lampitt and Paterson, 1987; Riemann-Zürneck, 1998), all factors that could influence the differences in morphospecies observed.

Some taxa were only observed on the seamounts in this study, while they ~~occurred~~ were also known to occur on nodule fields elsewhere, be it in low densities. For instance, Enteropneusta, which in this study were found only on seamounts, were observed previously at CCZ nodule fields though observations were rather rare (Tilot, 2006). They appeared more abundant at the nodule fields of the Deep Peru Basin (DISCOL area), though a wide range in abundances was displayed there as well (Bluhm, 2001). The exception were the Scleractinia, which were ~~absent at nodule fields but~~ quite common on seamounts, as also reported in other studies (e.g. ~~this study~~, Baco, 2007, Rowden et al., 2010), but, distinctly absent at nodule fields. Contrastingly, Enteropneusta were observed previously at CCZ nodule fields though observations were rather rare (Tilot, 2006). They appeared more abundant at the nodule fields of the Deep Peru Basin (DISCOL area), though a wide range in abundances was displayed there as well (Bluhm, 2001).

Explanation for the discrepancies ~~observed here~~ in faunal composition and the low degree of morphospecies overlap between seamount and nodule fields, as observed here, can be multiple. For one, nodules may not be considered a plain hard substratum, with their metal composition, microbial colonisation and the nodule/sediment interface influencing the epi- and associated megafaunal composition. The possibility of a specific deep-sea faunal community that tolerates or benefits from manganese substrata has been previously proposed (Mullineaux, 1988). The comparison between seamounts and nodule fields as two neighbouring hard-substrata ecosystems also entailed a comparison between depth gradients and possible thresholds (>4000m for nodule fields and 1500>x<4000m for seamounts). Related to this is the steepness of the seamount slope and its current exposure playing a role in the faunal colonisation (Genin et al., 1986; Rappaport et al., 1997). Other studies showed that habitat heterogeneity increased megafaunal diversity at seamounts (Raymore, 1982) and elsewhere, such as abyssal plains (Lapointe and Bourget, 1999; Durden et al., 2015b, Leitner et al., 2017, Simon-Llédo et al., 2019). Within this perspective the smaller-scale substratum heterogeneity transcending the ratio hard/soft substrata or amount of hard substrata ~~available~~ could play a role as well.

5. Conclusions

Based on our current knowledge; seamounts appear inadequate as refuge areas to help maintain nodule biodiversity. In order to conclusively exclude seamount habitats as a refuge for nodule fauna, a more comprehensive sampling should be carried out. The sampling strategy wielded in this study lacked replicates, uniformity and was limited in sample size. Seamount bases should be taken into consideration as well as they can be characterised by distinctly different assemblages than the summits and they ~~occur at a feature~~ depth ranges more similar to nodule fields.

499 While their role as refuge area for nodule field fauna is currently debatable, the possible uniqueness
500 of the seamount habitat and its inhabiting fauna implies that seamounts need to be included in
501 management plans for the conservation of the biodiversity and ecosystems of the CCZ.

502 Author Contributions

503 DC, PAR, SPR, DK analysed the images. DC analysed the data. PMA, PAR, AC conceptualised and
504 carried out the sampling. All authors contributed to the redaction of the manuscript.

505 Data Availability

506 Data sets are made available through OSIS-Kiel data portal, BIIGLE and PANGAEA.

507 Competing interest

508 The authors declare that they have no conflict of interest

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525

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 627

628 Tables

629 Table 1: Overview table on details of imagery transects analysed in the Clarion-Clipperton license
 630 areas. Video duration includes time spent sampling. Number of observations include undetermined
 631 organisms. Transect lengths do not include parts visualising ancient disturbance tracks or parts when
 632 the seafloor was not visualised or visible.

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Station/Dive	License Area	Seamount (SM) or Nodule field (NF)	Depth (m)	Video duration	Transect length	# obs/ dive	# obs /100m
SO239_29_ROV02	BGR	SM: Rüppell	3000-2500	7h47	1250m	<u>414429</u>	<u>33-134.</u> <u>3</u>
SO239_41_ROV03	BGR	NF	4080-4110	6h32	1590m	<u>102393</u> <u>2</u>	<u>64-358.</u> <u>6</u>
SO239_54_ROV04	BGR	SM: Senckenberg	3350-2850	8h45	2500m	<u>853890</u>	<u>34-135.</u> <u>6</u>
SO239_131_ROV08	GSR	NF	4470-4480	7h35	710m	<u>486445</u>	<u>68-562.</u> <u>8</u>

SO239_135_ROV09	GSR	SM: Heip	3900-3550	7h35	1000m	365359	36.535. <u>9</u>
SO239_141_ROV10	GSR	NF	4455-4480	7h35	520m	464351	89.267. <u>5</u>
SO239_189_ROV13	APEI 3	NF	4890-4930	9h01	1790m	136113	7.66.3
SO239_200_ROV14	APEI 3	NF	4650-4670	9h19	1490m	184179	12.212. <u>0</u>
SO239_212_ROV15	APEI 3	SM: Mann Borgese	1850-1650	6h25	900m	366378	40.742. <u>0</u>

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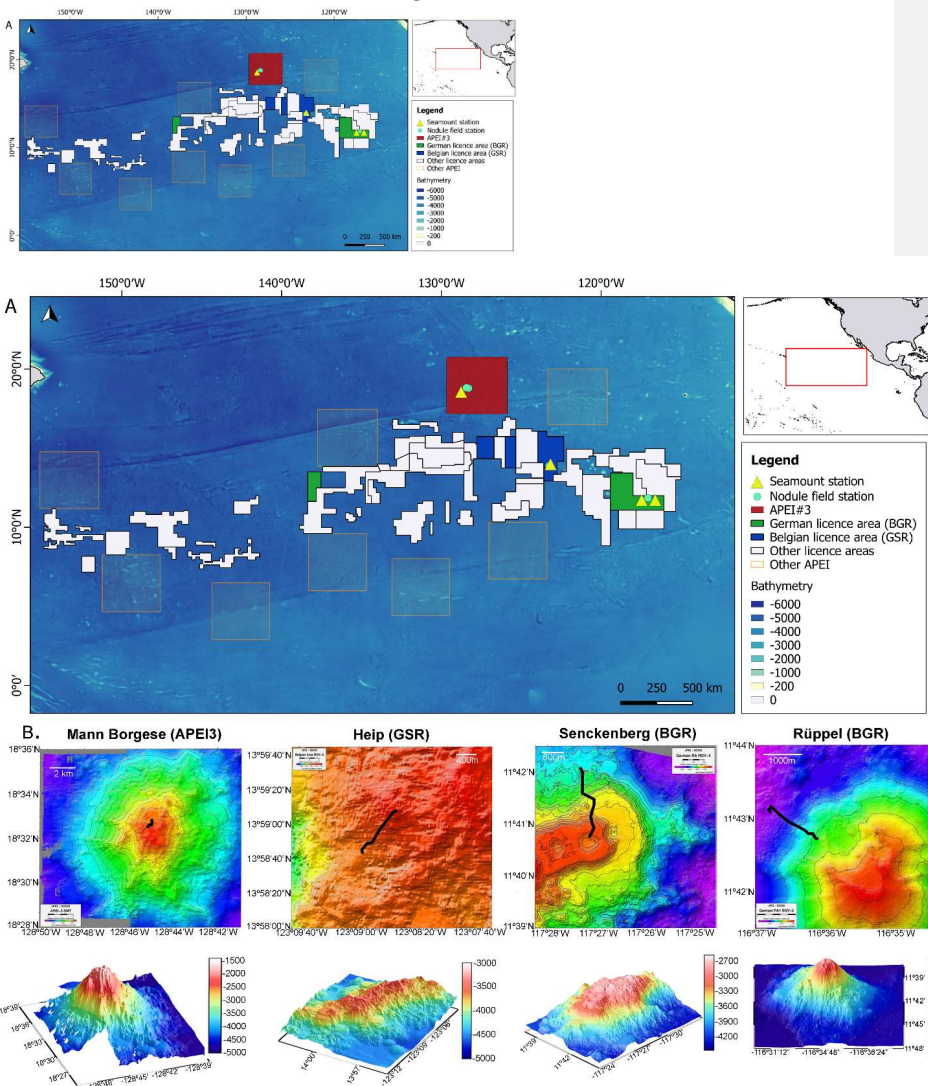
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Figures

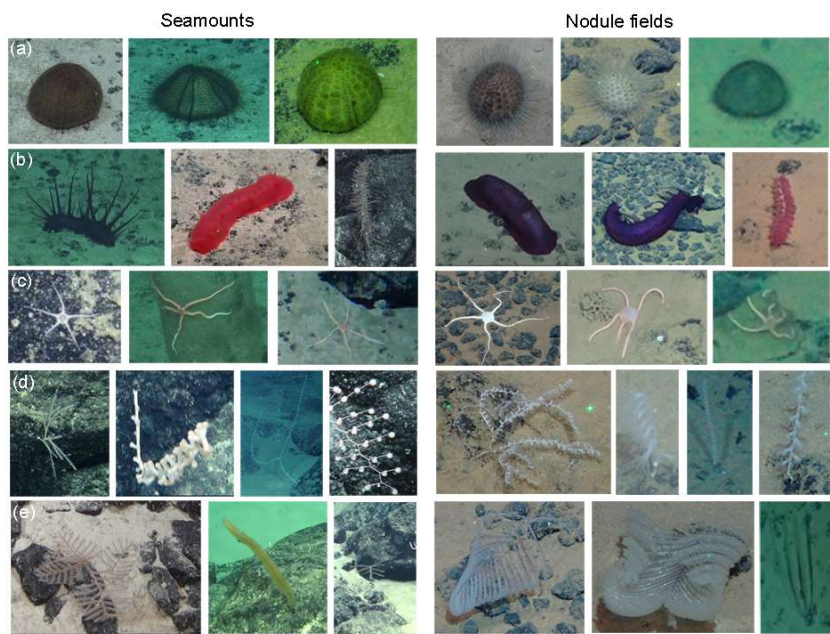
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651 Fig. 1. (A). Location of the Clarion-Clipperton Fracture zone in the equatorial eastern Pacific Ocean
652 featuring the contract areas from the International Seabed Authority (ISA) and the positions of the
653 sampled areas (seamounts and nodule fields). Information on transect length and depth gradients
654 can be found in Table 1. (B). Location of the seamount transects carried out towards the summit on
655 the north–north-western flank and seamount profiles. Rüppel (BGR, ROV02) and Mann Borgese
656 (APEI3, ROV15) are single seamounts, while Senckenberg (BGR, ROV04) and Heip (GSR, ROV09) are

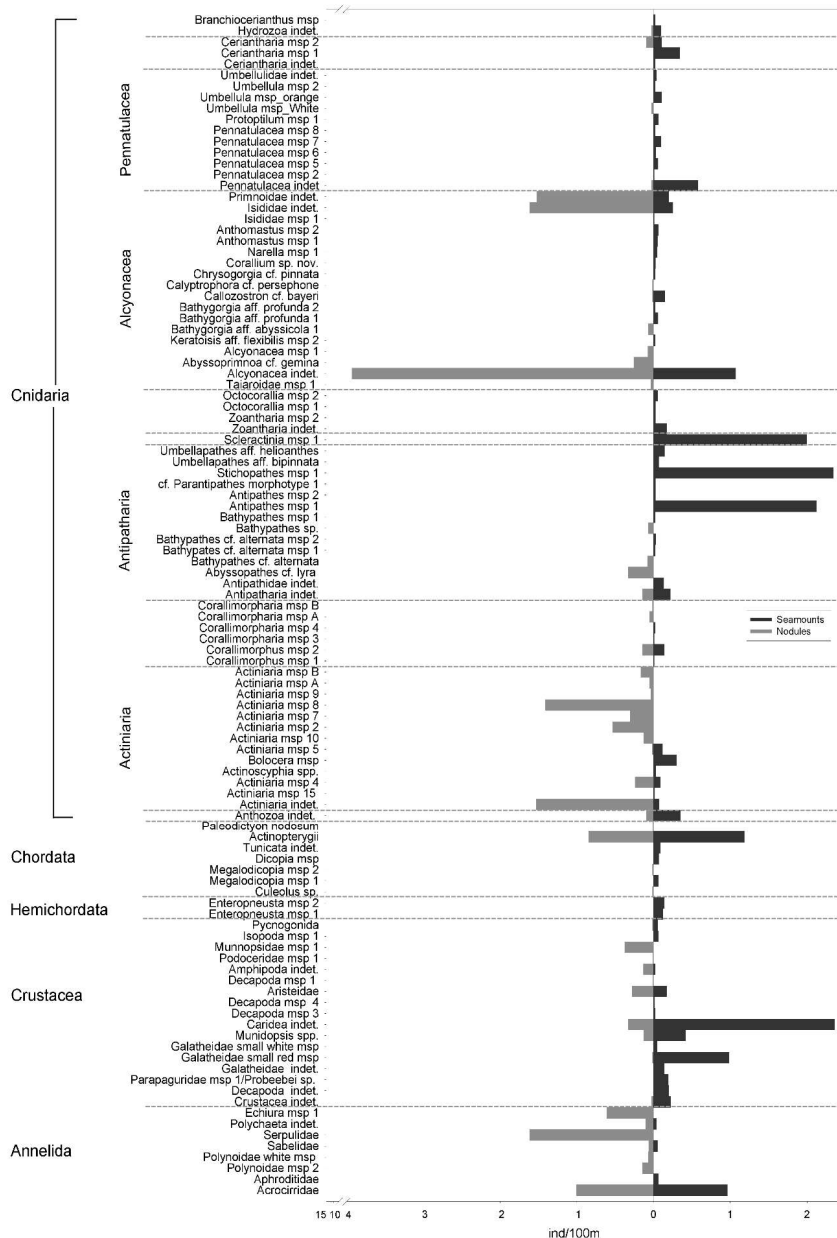


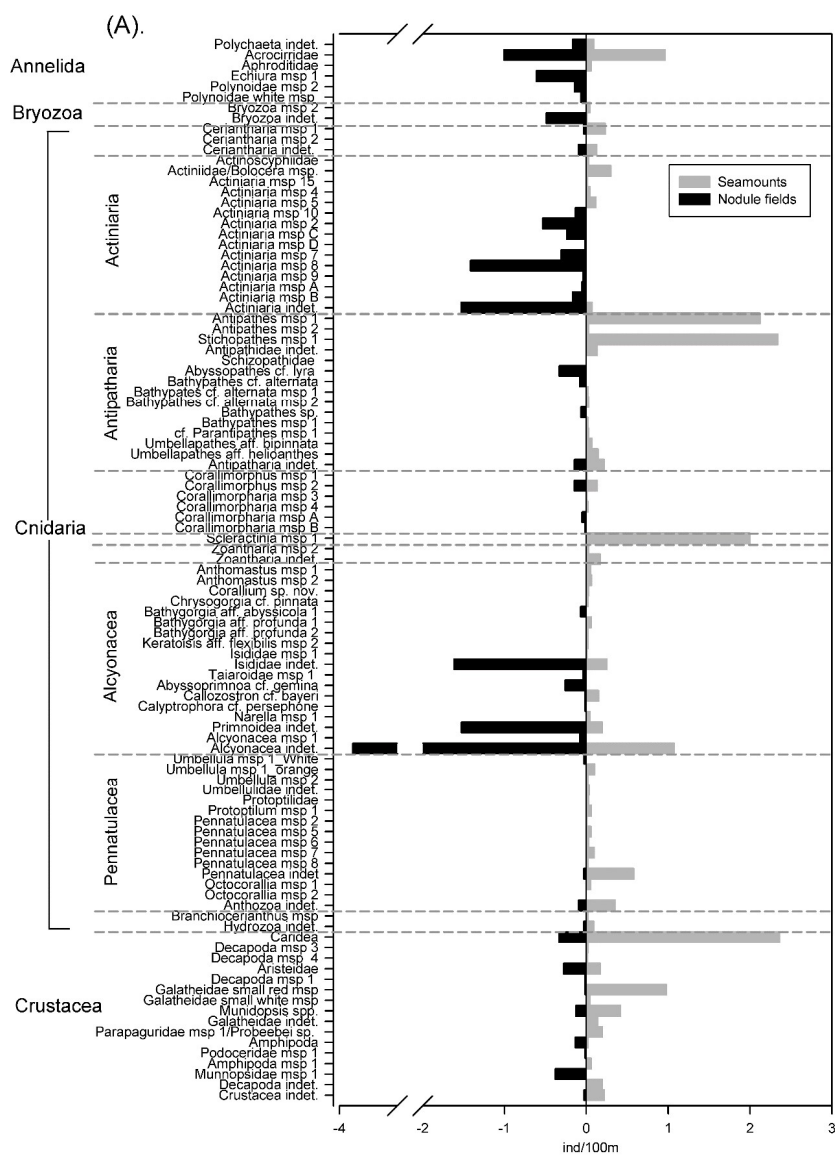
657 sea-mountain ranges.
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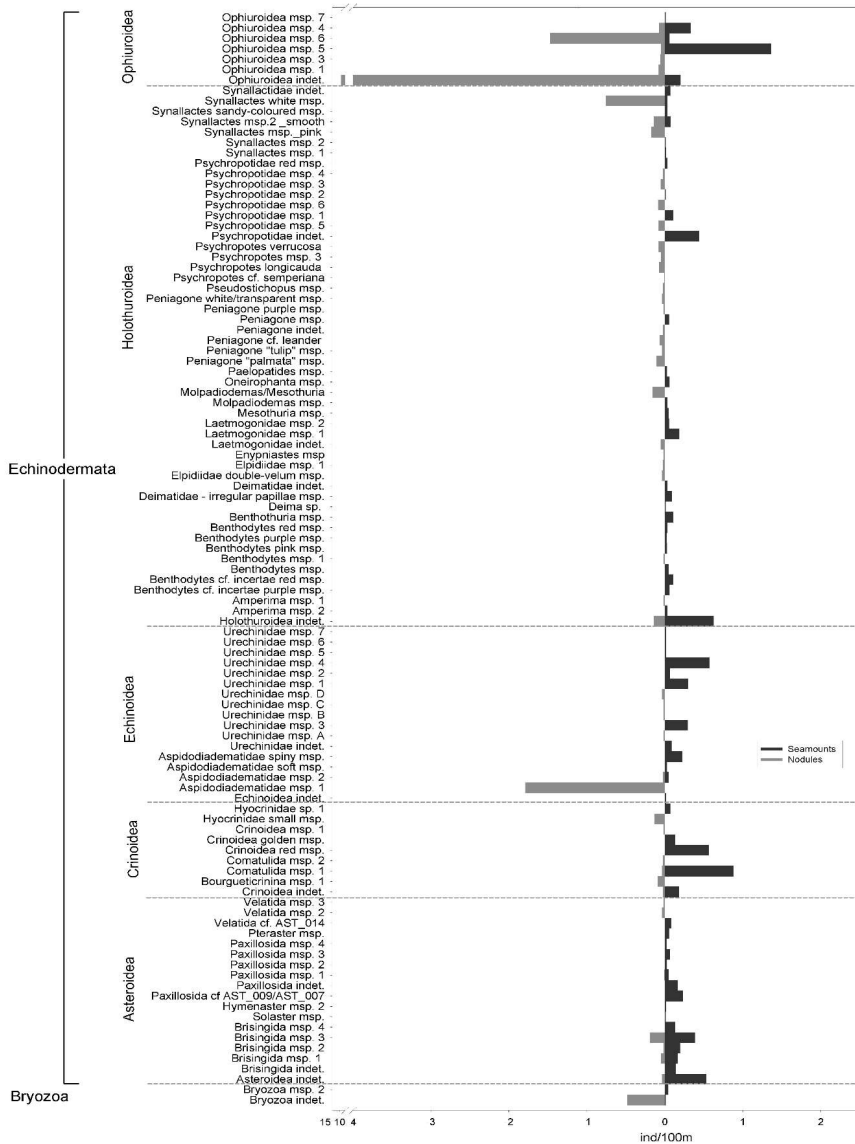


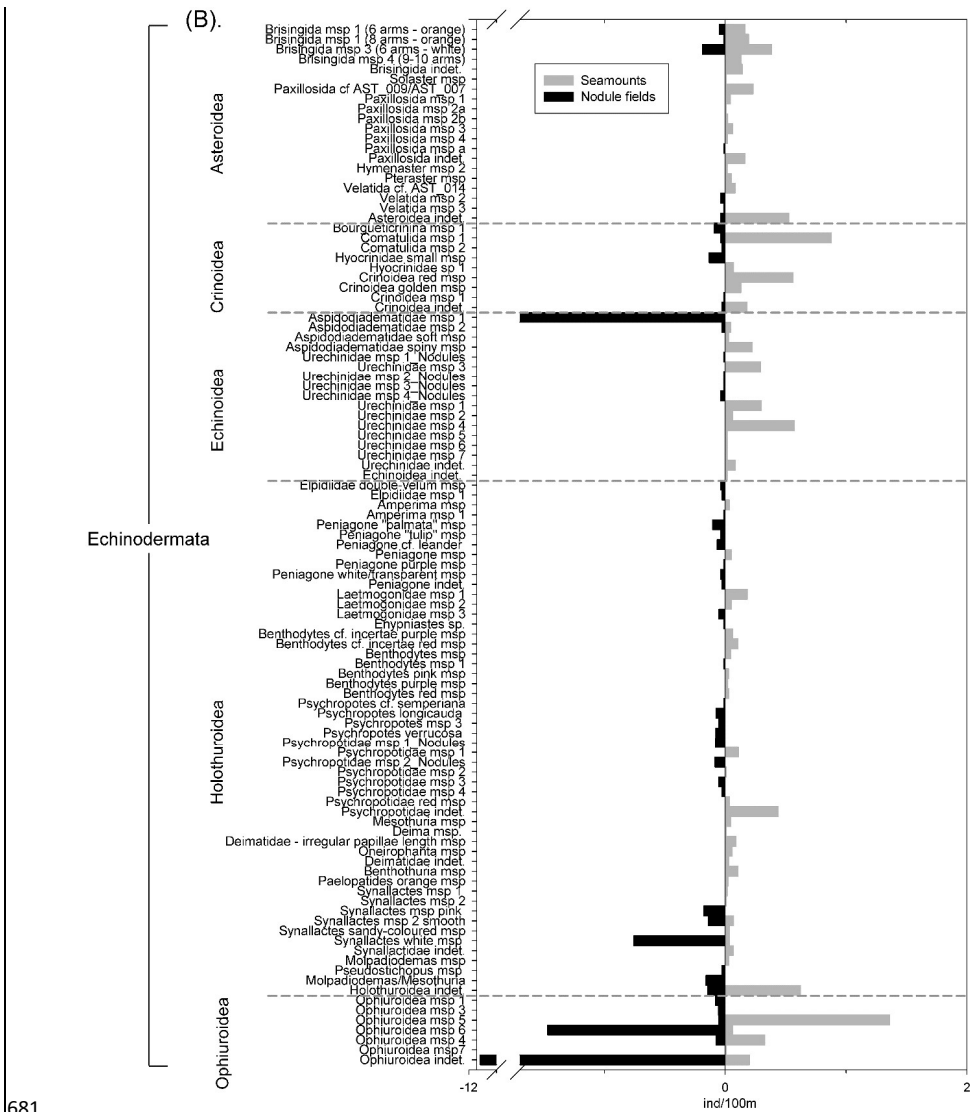
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661 Fig. 2. Some examples of different morphospecies at seamounts and nodule fields in the CCZ.
662 Selected taxa were (a) Echinoidea (from left to right, *Urechinidae* msp 4 (URC 019), *Urechinidae* msp
663 2 (URC 013), *Urechinidae* msp 3 (URC 009), *Urechinidae* msp. A (URC 020), *Urechinidae* msp. B
664 (URC 021), *Urechinidae* msp. C (URC 005), (b) Holothuroidea (from left to right, *Psychropotidae*
665 msp 1 (HOL 088-), *Benthodytes* red msp. (HOL 101), *Deimatidae* - irregular papillae msp. (HOL 070),
666 *Psychropotes verrucosa* (HOL 045), *Laetmogonidae* (HOL 030), *Synallactes* msp 2 pink (HOL 008)(c)
667 Ophiuroidea (from left to right, *Ophiuroidea* msp. 5 (OPH 003), *Ophiuroidea* msp. 4 (OPH 005),
668 *Ophiuroidea* msp. 6 (OPH 006), *Ophiuroidea* msp. 6 (OPH 006), *Ophiuroidea* (OPH 012),
669 *Ophiuroidea* msp. 4 (OPH 005)), (d) Alcyonacea (from left to right, *Callozostron* cf. *bayeri* (ALC 009),
670 *Bathygorgia* aff. *profunda* 2 (ALC 005), *Keratoisis* aff. *flexibilis* msp 2 (ALC 029), *Chrysoqorgia* cf.
671 *pinnata*-, *Abyssoprinoa* cf. *gemina* (ALC 008), *Bathygorgia* aff. *profunda*, 1, *Calyptrophora* cf.
672 *persephone* (ALC 007), *Bathygorgia* aff. *abyssicola* 1 (ALC 003), (e) Antipatharia (*Umbellapathes* aff.
673 *helioanthes* (ANT 018), cf. *Parantipathes* morphotype 1 (ANT 017), *Bathypates* cf. *alternata* msp 1
674 (ANT 010), *Bathypates* cf. *alternata* (ANT 006), *Abyssopathes* cf. *lyra* -(ANT 022), *Bathypates* sp.
675 (ANT 003)). Codes refer to an ongoing collaboration in creating one species catalogue for the CCZ
676 and align all morphospecies of different research groups. Copyright: SO239, ROV Kiel 6000, GEOMAR
677 Helmholtz Centre for Ocean Research Kiel

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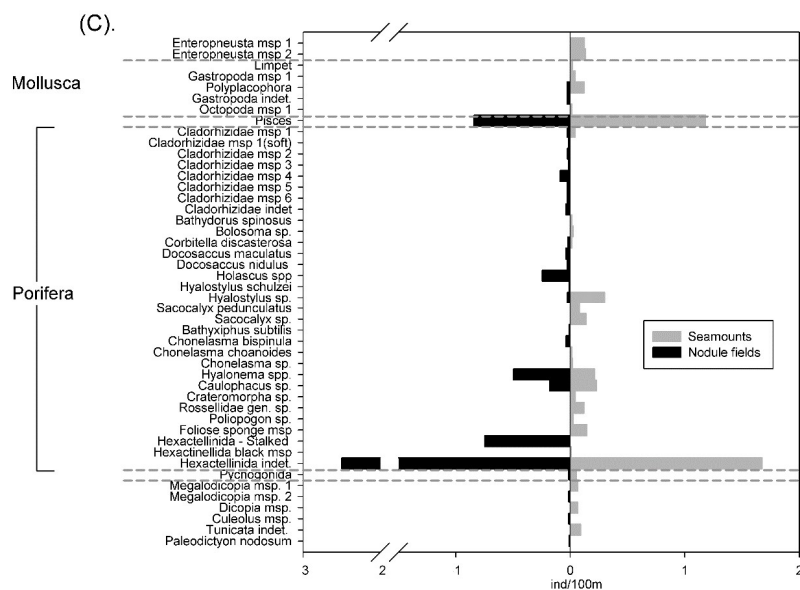
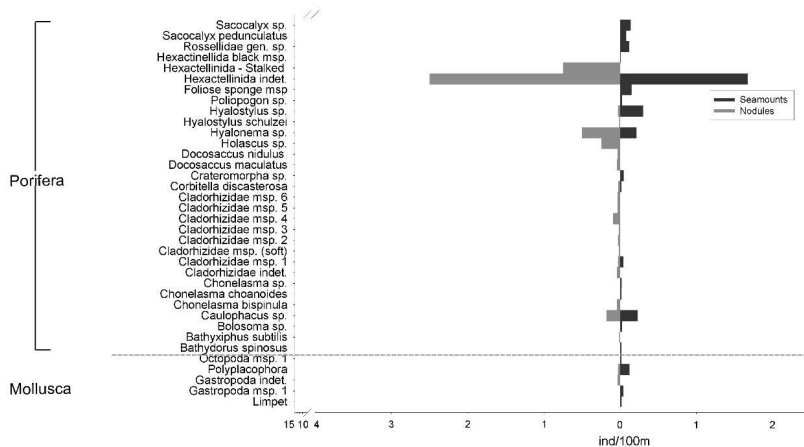
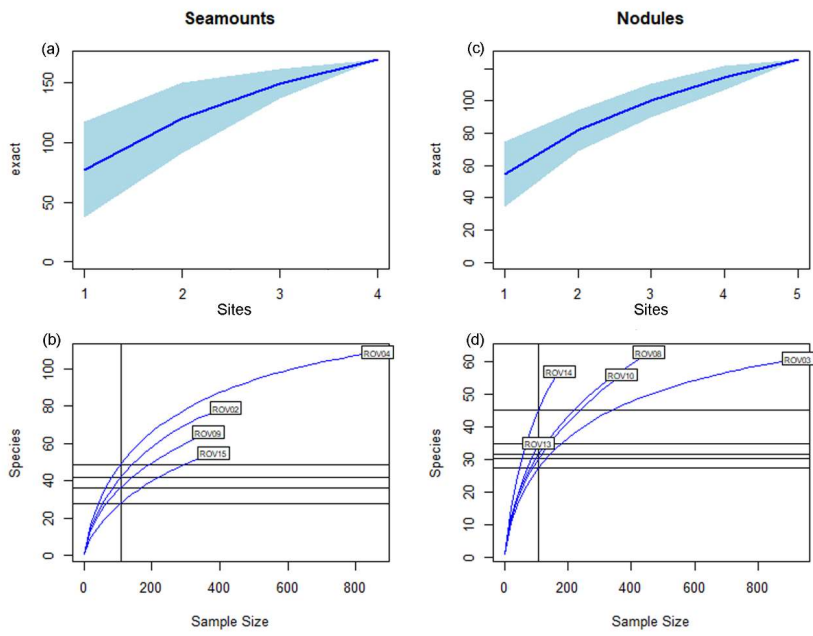


Fig. 3. Back-to-back histogram comparing average densities of morphospecies and taxa (ind/100m) for seamount (#4) and nodule field (#5) video transects. (a) Annelida, Bryozoa, Cnidaria and Crustacea, (B) Echinodermata and (C) Mollusca, Porifera, Hemichordata and Chordata (Tunicata).



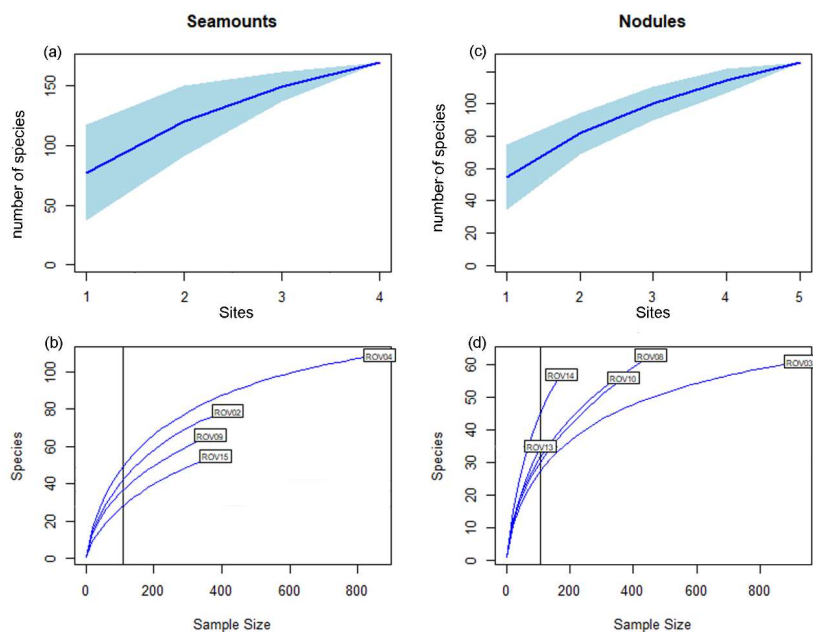


Fig. 4. Species accumulation (upper panel, a and c) and rarefaction curves (lower panel, b and d) for the seamount (#4) and nodule field (#5) transects. Seamount dives: ROV02= Rüppel (BGR), ROV04=Senckenberg (BGR), ROV09=Heip (GSR), ROV15=Mann Borgese (APEI3) in the lower left panel (b). Nodule field dives: ROV03 was carried out in the BGR area, ROV08 and 10 in the GSR area and ROV13 and 14 in the APEI3, presented in the lower right panel (d). Sample size is the number of individuals. Vertical line in the lower panel shows sample size=100.

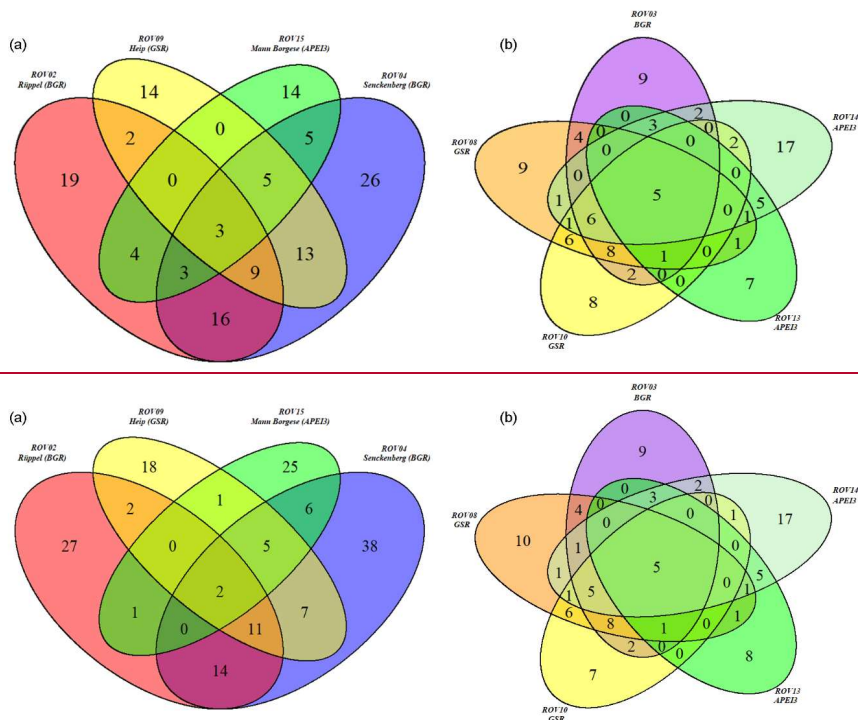


Fig. 5. A Venn diagram showing the unique and shared morphospecies among seamount video transects. Values are **relative-indicative rather than absolute** due to different transect lengths and differences in richness. Left panel (a) features seamount transects and the right panel features the 5 nodule field transects. Colour codes were adapted among panels, with APEI3 nodule transects in green, related to Mann Borgese seamount transect. BGR (ROV03) transect was purple in correspondence to BGR seamount transects (red=Rüppel and blue=Senckenberg). GSR transects (ROV08 and 09) were shades of yellow.

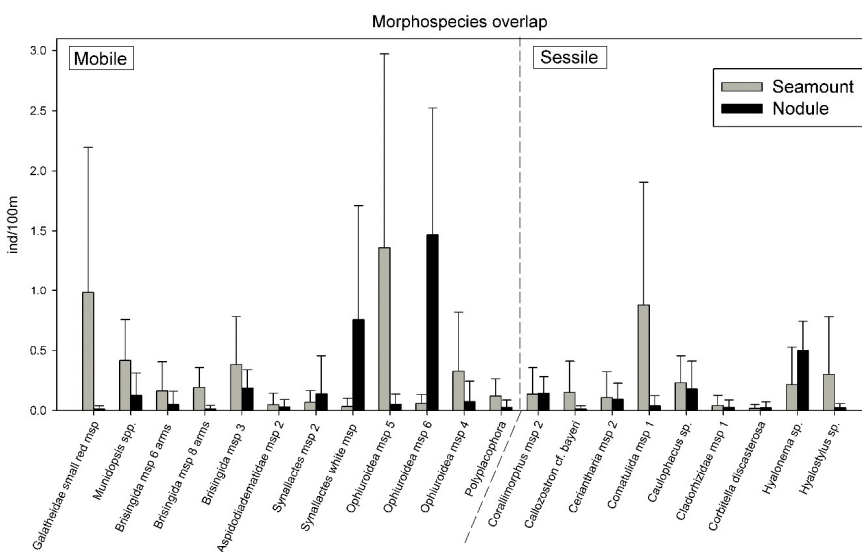
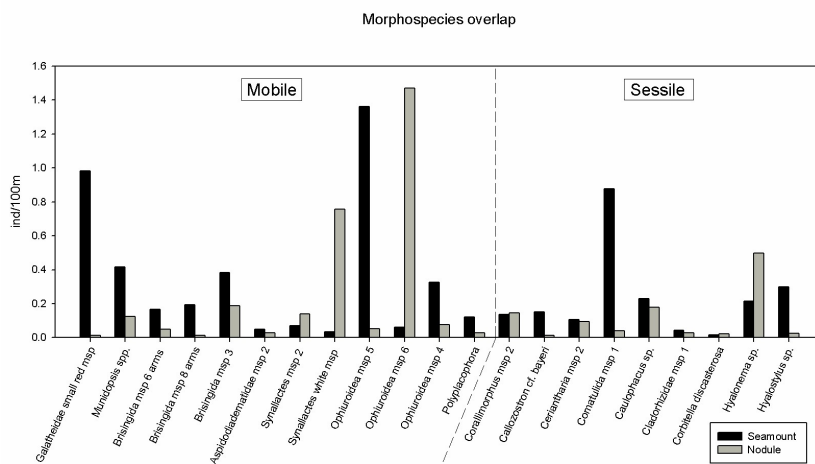


Fig. 6. Morphospecies present in both seamounts and nodule field transects overlap and their average density (ind/100m) and standard deviation per ecosystem. between seamounts and nodule fields

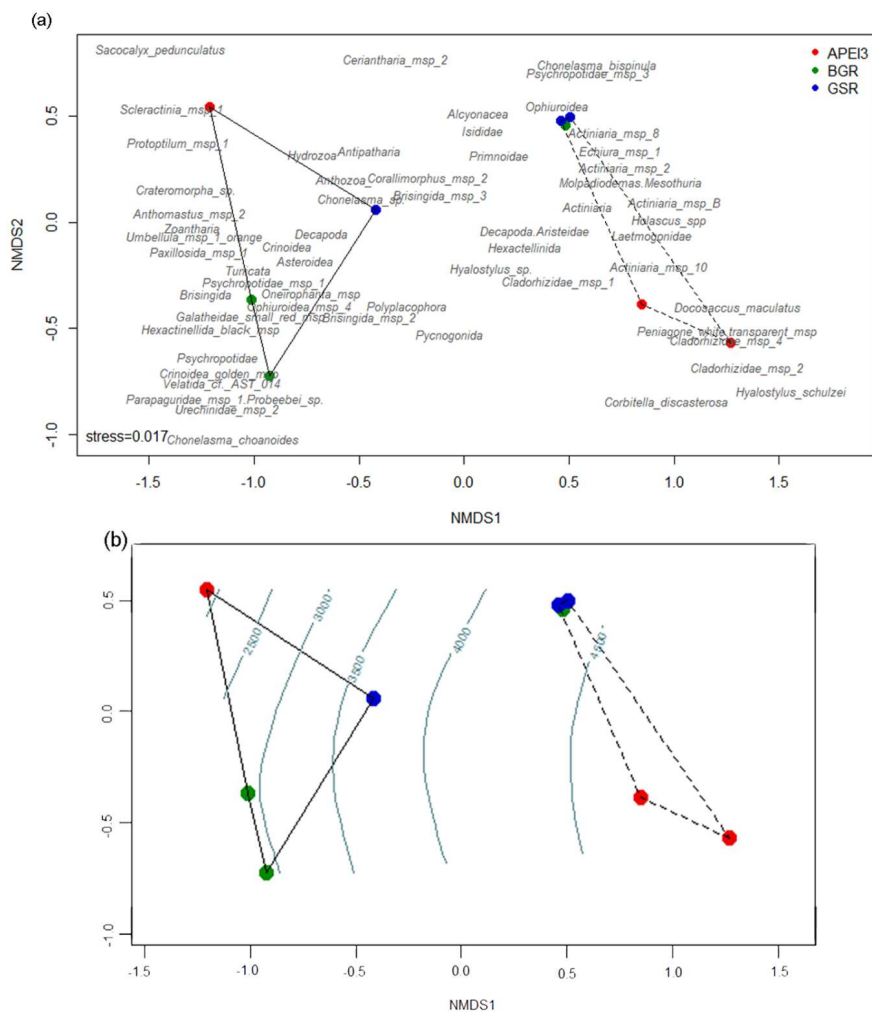


Fig. 7. nMDS-plot with faunal densities and Bray-Curtis distances. Left-Upper panel (a) presents the grouping of the video transects based on their faunal composition and right-lower panel (b) features the same plot but with depth as a vector fitting. Dotted lines group the nodule transects while the full line groups the seamount transects.

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Appendix

733 Table A1. Overview of all ~~taxa~~ densities (ind./100m) ~~and number of morphospecies (msp.)~~ observed
734 within each video transect. ~~Msp. numbers represent often minimum numbers and are indicative.~~

735 Higher taxa are in bold. * indicates taxa left out of the statistical analyses due to lack of
736 representativity. Indets were organisms impossible to attribute to a lower taxonomic group.

737 ROV02=Rüppel, ROV04=Senkcnberg, ROV09=Heip, ROV15=Mann Borgese

	SEAMOUNTS								NODULE FIELDS									
	ROV02 (BGR)		ROV04 (BGR)		ROV09 (GSR)		ROV15 (APEI3)		ROV03 (BGR)		ROV08 (GSR)		ROV10 (GSR)		ROV13 (APEI3)		ROV14	
	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp	ind/100m	# msp
-																		
Cnidaria			0.04	1	0.1	1	0.22	>1										
Actiniaria	0.32	3	0.32	3			0.44	2	4.40	8	6.90	6	9.42	8	0.22	3	1.14	6
-Actinoscyphiidae			0.12	1				-										
-Actiniidae/Bolocera msp.	1.20	1	0.20	1				-										
Alcyonacea			0.16	>1	1.7	>1	2.78	>2	9.06	>2	6.62	>1	4.04	>1	0.11	1	0.20	>2
-Alcyoniidae	0.24	1	0.04	1			0.11	1										
-Chrysogorgiidae					0.1	1		-										
-Isididae			0.20	2	1.2	>4		-	0.13	>1	5.77	>2	2.50	>2				
-Taiaroidea								-					0.19	1				
-Primnoidae	0.80	2	0.64	2	0.2	1	0.11	1	2.83	>3	4.08	>2	2.12	>2				
Coralliidae							0.11	1										
Ceriantharia	0.40	1	0.04	1	0.5	2	1.11	2			0.28	1	0.19	1				
Corallimorpharia/Corallimorphidae			0.64	4				-	0.38	3	0.28	1	0.38	2				
Antipatharia			0.08	1	0.2	1	0.67	>1	0.25	>1	0.28	>1	0.19	>1				
-Antipathidae							19.11	2										
-Schizopathidae			0.92	3	0.2	2	0.11	1	0.69	2	0.85	3	0.77	2			0.07	1
Pennatulacea	0.16	>1	3.04	>6			0.11	1										
-Umbellulidae			0.56	>2			0.11	1									0.13	1
-Protoptilidae			0.04	1			0.22	1										
Scleractinia	0.16	1					8.11	1										
Zoantharia			0.48	1			0.44	1										
Anthozoa	0.16	2	0.08	2	0.60	2	0.67	1	0.13	>1	0.14	>1	0.19	>1				
Bryozoa	-	-	0.04	1	0.2	1	-	-	0.44	>1	1.55	>1	0.19	>1	0.11	>1	0.13	>1
Echinodermata	0.16		0.16				-											
Asteroidea	0.32	>1	0.28	>1	1.3	>1	0.11	1	0.19	1.00								
-Brisingida	0.64	5	1.44	7	2.0	4	0.44	2	0.44	2	0.42	1	0.19	1			0.20	2
-Paxillosida	0.16	2	1.56	8	0.8	3	0.11	1							0.06	1		

	–Velatida	0.64	3	0.24	2			-				0.19	1			0.07	1
	Crinoidea	2.80	4	4.32	6	0.2	2	0.56	≥2	0.69	2	0.42	2			0.34	3
	Echinoidea							-									
	–Aspidodiadematidae	0.16	1	0.20	1			0.89	2	3.96	1	2.82	2				
	–Urechinidae	1.76	5	2.40	5	2.0	4	-		0.06	1			0.06	1	0.27	3
	Holothuroidea	1.60		0.84		0.3		0.22	-	0.38	2	0.56	3			0.13	1
	–Deimatidae	0.08	1	0.36	3	0.4	3	-									
	–Elpidiidae	0.32	2	0.08	1			-									
	–Lactmogonidae	0.56	2	0.48	1			-				0.19	1			0.07	1
	–Mesothuriidae			0.12	1			-									
	–Molpadiodemidae			0.12	1			-									
	–Psychropotidae	1.60	>3	1.64	>6	0.4	4	0.22	2	0.50	4	0.99	5	0.06	1	0.07	1
	–Synallactidae	1.04	3	0.16	3			0.44	1	2.45	2	1.69	3			0.07	1
	Ophiuroidea	0.48	2	3.40	5	4.4	3	0.44	1	20.44	4	18.45	4	25.96	3	1.34	3
	Crustacea*	0.07	-	0.31	-	0.51	-	-	-	-	-	-	-	-	-	0.13	-
	Decapoda	4.42	4	3.15	5	4.41	3	0.75	2	0.25	2	0.99	2	1.73	2	0.47	3
	–Galatheidae	4.48	3	1.28	5	1.0	2	0.56	1	0.06	1						
	–Parapaguridae	0.64	>1	0.24	2			-									
	–Peracarida	-	-	0.08	1	0.25	2	-	-	0.63	2	0.70	2	0.19	1	0.28	3
	Enteropneusta	-	-							-	-	-	-	-	-	-	-
	<i>Yoda</i> msp.	0.64	1	-	-	-	-	-	-								
	<i>Saxipendum</i> msp.	-	-	0.52	1	-	-	-	-	-	-	-	-	-	-	-	-
	Mollusca																
	Gastropoda			0.08	2	0.3	2					0.14	1				
	Polyplocophora	0.32	1	0.52	1			0.22	1							0.13	1
	Cephalopoda	0.08	1	-	-	-	-	-	-								
	Pisces*	2.96	9	1.44	10	0.6	5	0.33	3	1.57	4	0.42	2	1.54	3	0.34	2
	Annelida/Polychaeta *	0.16	1	0.12	>2	-	-	0.11	1	0.13	-	-	-	0.38	-	-	-
	Acrocirridae	0.16	1	0.16	1	4.2	1			0.57	1	0.14	1	0.58	1	1.79	1
	Aphroditidae	0.24	1	0.04	1												

Echiura					-	0.57	1	1.13	1	1.15	1	0.20	1	
Polynoidea	-	-	-	-	-	-	-	0.28	2	0.58	1	0.13	1	
Porifera	0.08	1	0.04	1	0.6	>1	-							
Hexactellinida	1.76	>2	1.16	>3	3.7	>2	1.00	>2	5.79	>2	5.21	>2	8.27	>2
–Euplectellidae	0.47	3	0.20	2	1.4	2	0.44	2	1	1	1	3	0	1
–Euretidae	0.08	1			0.1	1	-				0.19	1	0.06	1
–Hyalonematidae			0.08	>1	0.8	>2	0.11	1	0.38	>1	0.70	>1	0.77	>1
–Rosselidae	0.32	1	0.44	3	0.7	2	0.22	2	0.57	1	0.14	1	0.19	1
–Pheronematidae							0.11	1					0.00	0
Cladorhizidae					0.2	1	-		0	1	0	2		
Pycnogonida	0.16	1	-	-	0.1	1	-	-	-	-	-	-	-	-
Tunicata	0.08	1	-	-	0.1	1	-	-						
Megalodicopia msp.	0.16	1	0.04	1	0.1	1	-	-					0.07	1
Guleolus msp.							-	-					0.07	1
Dicopia msp.	0.4	1	-	-	-	-	-	-	-	-	-	-	-	-
Paleodictyon nodosum*							-					0.06	1	
Indets	0.88	>1	0.28	>1	-	-	0.22	>1	-	-	-	-	-	-

	SEAMOUNTS				NODULE FIELDS				
	ROV2	ROV4	ROV9	ROV15	ROV3	ROV8	ROV10	ROV13	ROV14
	ind/100m	ind/100m	ind/100m	ind/100m	ind/100m	ind/100m	ind/100m	ind/100m	ind/100m
Annelida*									
Polychaeta indet. * (No Serpulidae)	0.14	0.12	-	0.11	0.31	-	0.38	0.06	0.07
Acrocirridae	0.14	0.16	3.56	-	0.57	0.14	0.58	1.79	1.95
Aphroditidae	0.20	0.04	-	-		-		-	-
Echiura msp 1		-	-	-	0.57	1.13	1.15	-	0.20
Polynoidea		-	-	-		-		-	-
Polynoidae msp 2		-	-	-		0.14	0.58	-	-
Polynoidae white msp		-	-	-		0.14		0.06	0.13

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★ <u>Bryozoa</u>										Formatted: Font: 11 pt
★ <u>Bryozoa msp 2</u>		-	<u>0.17</u>	-	-	-	-	-	-	Formatted Table
★ <u>Bryozoa indet.</u>		<u>0.038</u>	-	-	<u>0.44</u>	<u>1.55</u>	<u>0.19</u>	<u>0.11</u>	<u>0.13</u>	Formatted: Font: 11 pt
★ <u>Cnidaria</u>										Formatted: Font: 11 pt
★ <u>Anthozoa</u>										Formatted: Font: 11 pt
★ <u>Ceriantharia</u>										Formatted: Font: 11 pt
★ <u>Ceriantharia msp 1</u>	<u>0.34</u>	<u>0.04</u>	<u>0.34</u>	<u>0.22</u>	-	-	-	-	-	Formatted Table
<u>Ceriantharia msp 2</u>	-	-	-	<u>0.43</u>	-	<u>0.28</u>	<u>0.19</u>	-	-	Formatted: Font: 11 pt
<u>Ceriantharia indet.</u>	-	-	<u>0.08</u>		-			-	-	Formatted: Font: 11 pt
★ <u>Hexacorrallia</u>										Formatted: Font: 11 pt
★ <u>Actiniaria</u>										Formatted: Font: 11 pt
★ <u>Actinoscyphiidae</u>		<u>0.12</u>	-	-	-	-	-	-	-	Formatted Table
★ <u>Actiniidae/Bolocera msp.</u>	<u>1.02</u>	<u>0.19</u>	-	-	-	-	-	-	-	Formatted: Font: 11 pt
<u>Actiniaria msp 15</u>	<u>0.07</u>	-	-	-	-	-	-	-	-	Formatted: Font: 11 pt
<u>Actiniaria msp 4</u>	-	<u>0.08</u>	-	<u>0.11</u>	-	-	-	-	-	Formatted: Font: 11 pt
<u>Actiniaria msp 5</u>	<u>0.07</u>	<u>0.08</u>	-	<u>0.32</u>	-	-	-	-	-	Formatted: Font: 11 pt
<u>Actiniaria msp 10</u>	-	-	-	-	<u>0.31</u>	-	-	-	<u>0.34</u>	Formatted: Font: 11 pt
<u>Actiniaria msp 2</u>	-	-	-	-	<u>1.07</u>	<u>1.13</u>	<u>0.19</u>	-	<u>0.27</u>	
<u>Actiniaria msp C</u>	-	-	-	-	<u>0.38</u>	<u>0.42</u>	<u>0.38</u>	-	-	
<u>Actiniaria msp D</u>	-	-	-	-	<u>0.06</u>	-	-	-	-	
<u>Actiniaria msp 7</u>	-	-	-	-	<u>0.63</u>	<u>0.14</u>	<u>0.58</u>	<u>0.06</u>	<u>0.13</u>	
<u>Actiniaria msp 8</u>	-	-	-	-	<u>0.13</u>	<u>3.66</u>	<u>3.08</u>	-	<u>0.20</u>	
<u>Actiniaria msp 9</u>	-	-	-	-	-	-	<u>0.19</u>	-	-	
<u>Actiniaria msp A</u>	-	-	-	-	-	-	<u>0.19</u>	-	<u>0.07</u>	
<u>Actiniaria msp B</u>	-	-	-	-	<u>0.25</u>	<u>0.14</u>	<u>0.38</u>	<u>0.06</u>	-	
<u>Actiniaria indet.</u>	<u>0.14</u>	<u>0.15</u>	-	-	<u>1.57</u>	<u>1.41</u>	<u>4.42</u>	<u>0.11</u>	<u>0.13</u>	
★ <u>Antipatharia</u>										Formatted: Font: 11 pt
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Antipathidae											Formatted: Font: 11 pt
Antipathes msp 1	-	-	-	8.49	-	-	-	-	-		
Antipathes msp 2	-	-	-	0.11	-	-	-	-	-		
Stichopathes msp 1	-	-	-	9.35	-	-	-	-	-		
Antipathidae indet.	-	-	-	0.54	-	-	-	-	-		
Schizopathidae											Formatted: Font: 11 pt
Abyssopathes cf. lyra	-	-	-	-	0.50	0.56	0.58	-	-		Formatted Table
Bathypathes cf. alternata	-	-	-	-	-	0.14	0.19	-	0.07		
Bathypates cf. alternata msp 1	-	-	0.08	-	-	-	-	-	-		
Bathypathes cf. alternata msp 2	-	0.12	-	-	-	-	-	-	-		
Bathypathes sp.	-	-	-	-	0.19	0.14	-	-	-		
Bathypathes msp 1	-	-	0.08	-	-	-	-	-	-		
cf. Parantipathes msp 1	-	-	0.11	-	-	-	-	-	-		
Umbellapathes aff. bipinnata	-	0.19	0.08	-	-	-	-	-	-		
Umbellapathes aff. helioanthes	-	0.58	-	-	-	-	-	-	-		
Antipatharia indet.	0.07	0.08	0.08	0.65	0.25	0.28	0.19	-	-		
Corallimorpharia/Corallimorphidae	-	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Corallimorphus msp 1	-	0.04	0.00	-	-	-	-	-	-		
Corallimorphus msp 2	-	0.46	0.08	-	0.25	0.28	0.19	-	-		
Corallimorpharia msp 3	-	0.04	-	-	-	-	-	-	-		
Corallimorpharia msp 4	-	-	0.08	-	-	-	-	-	-		
Corallimorpharia msp A	-	-	-	-	0.06	-	0.19	-	-		
Corallimorpharia msp B	-	-	-	-	0.06	-	-	-	-		
Scleractinia											Formatted: Font: 11 pt
Scleractinia msp 1	0.14	-	-	7.85	-	-	-	-	-		Formatted Table
Zoantharia											Formatted: Font: 11 pt
Zoantharia msp 2	-	-	-	0.11	-	-	-	-	-		Formatted Table
Zoantharia indet.	-	0.46	-	0.22	-	-	-	-	-		Formatted: Font: 11 pt
Octocorralia											Formatted Table

Alcyonacea											Formatted: Font: 11 pt
Alcyoniidae											Formatted: Font: 11 pt
Anthomastus msp 1	0.20	-	-	-	-	-	-	-	-		
Anthomastus msp 2	0.00	0.15	-	0.11	-	-	-	-	-		
Coralliidae											Formatted: Font: 11 pt
Corallium sp. nov.	-	-	-	0.11	-	-	-	-	-		Formatted Table
Chrysogorgiidae											Formatted: Font: 11 pt
Chrysogorgia cf. pinnata	-	-	0.08	-	-	-	-	-	-		Formatted Table
Isididae											Formatted: Font: 11 pt
Bathygorgia aff. abyssicola 1	-	-	-	-	-	0.14	0.19	-	-		Formatted Table
Bathygorgia aff. profunda 1	-	0.15	0.08	-	-	-	-	-	-		
Bathygorgia aff. profunda 2	-	-	0.08	-	-	-	-	-	-		
Keratoisis aff. flexibilis msp 2	-	-	0.08	-	-	-	-	-	-		
Isididae msp 1	-	0.04	-	-	-	-	-	-	-		
Isididae indet.	0.14	-	0.76	0.11	0.13	5.63	2.31	-	-		
Taiaroidea											Formatted: Font: 11 pt
Taiaroidae msp 1	-	-	-	-	-	-	0.19	-	-		Formatted Table
Primnoidae											Formatted: Font: 11 pt
Abyssoprimnoa cf. gemina	-	-	-	-	-	0.70	0.58	-	-		Formatted Table
Callozostron cf. bayeri	0.07	0.54	-	-	0.06	-	-	-	-		
Calyptrophora cf. persephone	-	-	-	-	0.06	-	-	-	-		
Narella msp 1	-	0.08	-	0.11	-	-	-	-	-		
Primnoidea indet.	0.61	-	0.17	-	2.70	3.38	1.54	-	-		
Alcyonacea msp 1	-	-	-	-	0.13	-	-	0.11	0.13		
Alcyonacea indet.	-	0.15	1.44	2.69	8.93	6.62	4.04	-	0.07		Formatted: Font: 11 pt
Pennatulacea											Formatted: Font: 11 pt
Umbellulidae											Formatted Table
Umbellula msp 1 White	-	-	-	-	-	-	-	-	0.13		
Umbellula msp 1 orange	-	0.31	-	0.11	-	-	-	-	-		Formatted: Font: 11 pt

<u>Umbellula msp 2</u>	-	<u>0.08</u>	-	-	-	-	-	-	-
<u>Umbellulidae indet.</u>	-	<u>0.15</u>	-	-	-	-	-	-	-
<u>Protoptilidae</u>	-	-	-	<u>0.11</u>	-	-	-	-	-
<u>Protoptilum msp 1</u>	-	<u>0.04</u>	-	<u>0.22</u>	-	-	-	-	-
<u>Pennatulacea msp 2</u>	-	<u>0.04</u>	-	-	-	-	-	-	-
<u>Pennatulacea msp 5</u>	-	<u>0.23</u>	-	-	-	-	-	-	-
<u>Pennatulacea msp 6</u>	-	<u>0.12</u>	-	-	-	-	-	-	-
<u>Pennatulacea msp 7</u>	-	<u>0.38</u>	-	-	-	-	-	-	-
<u>Pennatulacea msp 8</u>	-	<u>0.08</u>	-	-	-	-	-	-	-
<u>Pennatulacea indet.</u>	<u>0.14</u>	<u>2.08</u>	-	<u>0.11</u>	-	<u>0.14</u>	-	-	-
<u>Octocorallia msp 1</u>	-	-	-	<u>0.22</u>	-	-	-	-	-
<u>Octocorallia msp 2</u>	-	-	-	-	-	-	-	-	-
<u>Anthozoa indet.</u>	<u>0.14</u>	<u>0.12</u>	<u>0.51</u>	<u>0.65</u>	<u>0.13</u>	<u>0.14</u>	<u>0.19</u>	-	-
<u>Hydrozoa</u>									
<u>Branchiocerianthus msp</u>	-	<u>0.08</u>	-	-	-	-	-	-	-
<u>Hydrozoa indet.</u>	-	<u>0.08</u>	<u>0.08</u>	<u>0.22</u>	-	<u>0.14</u>	-	-	-
<u>Crustacea*</u>									
<u>Decapoda</u>									
<u>Caridea</u>	<u>3.47</u>	<u>2.54</u>	<u>3.22</u>	<u>0.22</u>	<u>0.19</u>	-	<u>1.15</u>	<u>0.11</u>	<u>0.20</u>
<u>Decapoda msp 3</u>	-	<u>0.08</u>	-	-	-	-	-	-	-
<u>Decapoda msp 4</u>	<u>0.07</u>	-	-	-	-	-	-	-	-
<u>Decapoda/Aristeidae</u>	<u>0.07</u>	<u>0.08</u>	-	<u>0.54</u>	<u>0.06</u>	<u>0.56</u>	<u>0.58</u>	<u>0.11</u>	<u>0.07</u>
<u>Decapoda msp 1</u>	-	-	-	-	-	-	-	<u>0.06</u>	-
<u>Galatheidae</u>									
<u>Galatheidae small red msp</u>	<u>2.79</u>	<u>0.54</u>	<u>0.17</u>	<u>0.43</u>	<u>0.06</u>	-	-	-	-
<u>Galatheidae small white msp</u>	<u>0.07</u>	<u>0.12</u>	-	-	-	-	-	-	-
<u>Munidopsis spp.</u>	<u>0.82</u>	<u>0.35</u>	<u>0.51</u>	-	-	<u>0.42</u>	-	-	<u>0.20</u>
<u>Galatheidae indet.</u>	<u>0.14</u>	<u>0.15</u>	<u>0.17</u>	<u>0.11</u>	-	-	-	-	-

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▲ <u>Parapaguridae</u>											Formatted: Font: 11 pt
▲ <u>Parapaguridae msp 1/Probeebei sp.</u>	<u>0.54</u>	<u>0.23</u>	-	-	-	-	-	-	-	-	Formatted Table
▲ <u>Peracarida</u>											Formatted: Font: 11 pt
▲ <u>Amphipoda</u>	-	-	<u>0.08</u>	-	<u>0.06</u>	<u>0.28</u>	-	<u>0.06</u>	<u>0.27</u>		Formatted Table
▲ <u>Podoceridae msp 1</u>	-	-	-	-	-	-	-	<u>0.06</u>	-		
▲ <u>Amphipoda msp 1</u>	-	<u>0.08</u>	<u>0.17</u>	-	-	-	-	-	-		
▲ <u>Isopoda</u>											Formatted Table
▲ <u>Munnopsidae msp 1</u>	-	-	-	-	<u>0.57</u>	<u>0.42</u>	<u>0.19</u>	<u>0.17</u>	<u>0.54</u>		
▲ <u>Decapoda indet.</u>	-	<u>0.12</u>	<u>0.68</u>	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Crustacea indet.</u>	<u>0.07</u>	<u>0.31</u>	<u>0.51</u>	-	-	-	-	-	<u>0.13</u>		Formatted: Font: 11 pt
▲ <u>Echinodermata</u>											Formatted: Font: 11 pt
▲ <u>Asteroidea</u>											Formatted Table
▲ <u>Brisingida</u>											Formatted: Font: 11 pt
▲ <u>Brisingida msp 1 (6 arms - orange)</u>	-	<u>0.15</u>	<u>0.51</u>	-	<u>0.25</u>	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Brisingida msp 1 (8 arms - orange)</u>	<u>0.14</u>	<u>0.38</u>	<u>0.25</u>	-	-	-	-	-	<u>0.07</u>		Formatted: Font: 11 pt
▲ <u>Brisingida msp 3 (6 arms - white)</u>	-	<u>0.38</u>	<u>0.93</u>	<u>0.22</u>	<u>0.19</u>	<u>0.42</u>	<u>0.19</u>	-	<u>0.13</u>		Formatted: Font: 11 pt
▲ <u>Brisingida msp 4 (9-10 arms)</u>	<u>0.14</u>	<u>0.38</u>	-	-	-	-	-	-	-		
▲ <u>Brisingida indet.</u>	<u>0.27</u>	<u>0.08</u>	-	<u>0.22</u>	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida</u>											Formatted: Font: 11 pt
▲ <u>Solaster msp</u>	-	<u>0.04</u>	-	-	-	-	-	-	-		Formatted Table
▲ <u>Paxillosida cf AST_009/AST_007</u>	-	<u>0.50</u>	<u>0.42</u>	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 1</u>	<u>0.07</u>	-	-	<u>0.11</u>	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 2a</u>	-	<u>0.04</u>	-	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 2b</u>	-	-	<u>0.08</u>	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 3</u>	-	<u>0.08</u>	<u>0.17</u>	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 4</u>	-	<u>0.08</u>	-	-	-	-	-	-	-		Formatted: Font: 11 pt
▲ <u>Paxillosida msp 1</u>	-	-	-	-	-	-	-	<u>0.06</u>	-		Formatted: Font: 11 pt
▲ <u>Paxillosida indet.</u>	-	<u>0.65</u>	-	-	-	-	-	-	-		Formatted: Font: 11 pt

◀ Velatida											Formatted: Font: 11 pt
◀ Pterasteridae											Formatted Table
◀ Hymenaster msp 2	<u>0.07</u>	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
◀ Pteraster msp	<u>0.20</u>	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
◀ Velatida cf. AST_014	<u>0.14</u>	<u>0.19</u>	-	-	-	-	-	-	-		Formatted: Font: 11 pt
◀ Velatida msp 2	-	-	-	-	-	-	<u>0.19</u>	-	-		Formatted: Font: 11 pt
◀ Velatida msp 3	-	-	-	-	-	-	-	-	<u>0.07</u>		Formatted: Font: 11 pt
◀ Asteroidea indet.	<u>0.48</u>	<u>0.42</u>	<u>1.10</u>	<u>0.11</u>	<u>0.19</u>	-	-	-	-		Formatted: Font: 11 pt
◀ Crinoidea											Formatted: Font: 11 pt
◀ Comatulida											Formatted: Font: 11 pt
◀ Bourgueticrinina msp 1	-	-	-	-	<u>0.31</u>	-	-	-	<u>0.13</u>		Formatted: Font: 11 pt
◀ Comatulida msp 1	<u>1.97</u>	<u>1.54</u>	-	-	-	-	<u>0.19</u>	-	-		Formatted: Font: 11 pt
◀ Comatulida msp 2	-	-	-	-	-	-	-	-	<u>0.13</u>		Formatted Table
◀ Hyocrinida											Formatted: Font: 11 pt
◀ Hyocrinidae small msp	-	-	-	-	<u>0.38</u>	<u>0.28</u>	-	-	-		Formatted: Font: 11 pt
◀ Hyocrinidae sp 1	-	<u>0.19</u>	<u>0.08</u>	<u>0.00</u>	-	-	-	-	-		Formatted: Font: 11 pt
◀ Crinoidea red msp	<u>0.20</u>	<u>1.62</u>	-	<u>0.43</u>	-	-	-	-	-		Formatted Table
◀ Crinoidea golden msp	<u>0.14</u>	<u>0.38</u>	-	-	-	-	-	-	-		
◀ Crinoidea msp 1	-	-	-	-	-	-	-	-	<u>0.07</u>		
◀ Crinoidea indet.	<u>0.07</u>	<u>0.46</u>	<u>0.08</u>	<u>0.11</u>	-	<u>0.14</u>	-	-	-		Formatted: Font: 11 pt
◀ Echinoidea											Formatted: Font: 11 pt
◀ Aspidodiadematidae											Formatted Table
◀ Aspidodiadematidae msp 1	-	-	-	-	<u>3.96</u>	<u>2.68</u>	<u>2.31</u>	-	-		Formatted: Font: 11 pt
◀ Aspidodiadematidae msp 2	-	<u>0.19</u>	-	-	-	<u>0.14</u>	-	-	-		
◀ Aspidodiadematidae soft msp	-	-	-	<u>0.11</u>	-	-	-	-	-		
◀ Aspidodiadematidae spiny msp	<u>0.14</u>	-	-	<u>0.75</u>	-	-	-	-	-		
◀ Urechinidae											Formatted: Font: 11 pt
◀ Urechinidae msp 1 Nodules	-	-	-	-	-	-	-	-	<u>0.07</u>		Formatted Table
◀ Urechinidae msp 3	<u>0.20</u>	<u>0.04</u>	<u>0.93</u>	-	-	-	-	-	-		

Urechinidae msp 2 Nodules	-	-	-	-	-	-	-	-	0.07		Formatted: Font: 11 pt
Urechinidae msp 3 Nodules	-	-	-	-	0.06	-	-	-	-		Formatted: Font: 11 pt
Urechinidae msp 4 Nodules	-	-	-	-	-	-	-	0.06	0.13		Formatted: Font: 11 pt
Urechinidae msp 1	0.20	0.73	0.25	-	-	-	-	-	-		Formatted Table
Urechinidae msp 2	0.20	0.04	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Urechinidae msp 4	0.48	1.38	0.42	-	-	-	-	-	-		Formatted: Font: 11 pt
Urechinidae msp 5	0.07	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Urechinidae msp 6	0.07	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Urechinidae msp 7	0.07	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Urechinidae indet.	0.14	0.12	0.08	-	-	-	-	-	-		Formatted: Font: 11 pt
Echinoidea indet.	0.07	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Holothuroidea											Formatted: Font: 11 pt
Elasipodida											Formatted: Font: 11 pt
Elpidiidae											Formatted: Font: 11 pt
Elpidiidae double-velum msp	-	-	-	-	-	-	0.19	-	-		Formatted: Font: 11 pt
Elpidiidae msp 1	-	-	-	-	-	-	-	0.06	0.07		Formatted: Font: 11 pt
Amperima msp	0.14	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Amperima msp 1	-	-	-	-	0.06	-	-	-	-		Formatted: Font: 11 pt
Peniagone "palmata" msp	-	-	-	-	-	0.14	0.38	-	-		Formatted: Font: 11 pt
Peniagone "tulip" msp	-	-	-	-	-	-	0.19	-	-		Formatted: Font: 11 pt
Peniagone cf. leander	-	-	-	-	-	0.14	0.19	-	-		Formatted: Font: 11 pt
Peniagone msp	0.14	0.08	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Peniagone purple msp	-	-	-	-	-	-	-	-	0.07		Formatted: Font: 11 pt
Peniagone white/transparent msp	-	-	-	-	0.06	-	-	0.06	0.07		Formatted: Font: 11 pt
Peniagone indet.	-	-	-	-	0.13	-	-	-	-		Formatted: Font: 11 pt
Laetmogonidae											Formatted Table
Laetmogonidae msp 1	0.27	0.46	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Laetmogonidae msp 2	0.20	-	-	-	-	-	-	-	-		Formatted: Font: 11 pt
Laetmogonidae msp 3	-	-	-	-	-	-	0.19	-	0.07		Formatted: Font: 11 pt

⬅ <u>Pelagothuriidae</u>											
⬅ <u>Enypniastes sp.</u>	-	-	-	-	-	-	-	-	<u>0.07</u>		
⬅ <u>Psychropotidae</u>											
⬅ <u>Benthodytes cf. incertae purple msp</u>	-	<u>0.15</u>	<u>0.08</u>	-	-	-	-	-	-		
⬅ <u>Benthodytes cf. incertae red msp</u>	-	<u>0.42</u>	-	-	-	-	-	-	-		
⬅ <u>Benthodytes msp</u>	-	<u>0.19</u>	-	-	-	-	-	-	-		
⬅ <u>Benthodytes msp 1</u>	-	-	-	-	-	-	-	-	<u>0.07</u>		
⬅ <u>Benthodytes pink msp</u>	-	-	-	<u>0.11</u>	-	-	-	-	-		
⬅ <u>Benthodytes purple msp</u>	-	-	<u>0.08</u>	-	-	-	-	-	-		
⬅ <u>Benthodytes red msp</u>	-	<u>0.04</u>	<u>0.08</u>	-	-	-	-	-	-		
⬅ <u>Psychropotes cf. semperiana</u>	-	-	-	-	-	-	-	<u>0.06</u>	-		
⬅ <u>Psychropotes longicauda</u>	-	-	-	-	-	-	<u>0.38</u>	-	-		
⬅ <u>Psychropotes msp 3</u>	-	-	-	-	<u>0.06</u>	-	<u>0.19</u>	-	-		
⬅ <u>Psychropotes verrucosa</u>	-	-	-	-	<u>0.25</u>	<u>0.14</u>	-	-	-		
⬅ <u>Psychropotidae msp 1 Nodules</u>	-	-	-	-	<u>0.06</u>	<u>0.14</u>	<u>0.19</u>	-	-		
⬅ <u>Psychropotidae msp 1</u>	-	<u>0.35</u>	<u>0.08</u>	-	-	-	-	-	-		
⬅ <u>Psychropotidae msp 2 Nodules</u>	-	-	-	-	-	<u>0.42</u>	-	-	-		
⬅ <u>Psychropotidae msp 2</u>	-	<u>0.04</u>	-	-	-	-	-	-	-		
⬅ <u>Psychropotidae msp 3</u>	-	-	-	-	<u>0.13</u>	<u>0.14</u>	-	-	-		
⬅ <u>Psychropotidae msp 4</u>	-	-	-	-	-	<u>0.14</u>	-	-	-		
⬅ <u>Psychropotidae red msp</u>	<u>0.14</u>	-	-	-	-	-	-	-	-		
⬅ <u>Psychropotidae indet.</u>	<u>1.22</u>	<u>0.42</u>	-	<u>0.11</u>	-	-	-	-	-		
⬅ <u>Holothuriida</u>											
⬅ <u>Mesothuriidae</u>											
⬅ <u>Mesothuria msp</u>	<u>0.07</u>	<u>0.12</u>	-	-	-	-	-	-	-		
⬅ <u>Synallactida</u>											
⬅ <u>Deimatidae</u>											
⬅ <u>Deima msp.</u>	-	<u>0.04</u>	-	-	-	-	-	-	-		
⬅ <u>Deimatidae - irregular papillae length msp</u>	-	<u>0.27</u>	<u>0.08</u>	-	-	-	-	-	-		

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<u>Enteropneusta msp 1 cf. Yoda</u>	-	<u>0.50</u>	-	-	-	-	-	-	-
<u>Enteropneusta msp 2 cf. Saxipendum msp.</u>	<u>0.54</u>	-	-	-	-	-	-	-	-
▲ <u>Mollusca</u>									
▲ <u>Gastropoda</u>									
<u>Limpet</u>	-	-	<u>0.08</u>	-	-	-	-	-	-
<u>Gastropoda msp 1</u>	-	-	<u>0.17</u>	-	-	-	-	-	-
<u>Polyplacophora</u>	<u>0.27</u>	-	-	<u>0.22</u>	-	-	-	-	<u>0.13</u>
<u>Gastropoda indet.</u>	-	-	-	-	-	<u>0.14</u>	-	-	-
▲ <u>Cephalopoda</u>									
<u>Octopoda msp 1</u>	<u>0.07</u>	-	-	-	-	-	-	-	-
▲ <u>Pisces*</u>	<u>2.52</u>	<u>1.38</u>	<u>0.51</u>	<u>0.32</u>	<u>1.57</u>	<u>0.42</u>	<u>1.54</u>	<u>0.34</u>	<u>0.34</u>
▲ <u>Porifera</u>									
▲ <u>Demospongiae</u>									
<u>Cladorhizidae</u>									
<u>Cladorhizidae msp 1</u>		-	<u>0.17</u>	-	-	-	-	-	<u>0.13</u>
<u>Cladorhizidae msp 1(soft)</u>	-	-	-	-	-	-	-	-	<u>0.07</u>
<u>Cladorhizidae msp 2</u>	-	-	-	-	-	-	-	<u>0.06</u>	<u>0.07</u>
<u>Cladorhizidae msp 3</u>	-	-	-	-	-	-	-	-	<u>0.07</u>
<u>Cladorhizidae msp 4</u>	-	-	-	-	<u>0.06</u>	-	-	<u>0.11</u>	<u>0.27</u>
<u>Cladorhizidae msp 5</u>	-	-	-	-	-	<u>0.14</u>	-	-	-
<u>Cladorhizidae msp 6</u>	-	-	-	-	-	<u>0.14</u>	-	-	-
<u>Cladorhizidae indet</u>	-	-	-	-	-	-	-	<u>0.06</u>	<u>0.13</u>
▲ <u>Hexactellinida</u>									
<u>Euplectellidae</u>									
<u>Bathydorus spinosus</u>	<u>0.07</u>	-	-	-	-	-	-	-	-
<u>Bolosoma sp.</u>	-	-	-	<u>0.11</u>	-	-	-	-	-

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<u>Octacnemidae</u>									
◀ <u>Megalodicopia msp. 1</u>	<u>0.14</u>	<u>0.04</u>	<u>0.08</u>	-	-	-	-	-	-
◀ <u>Megalodicopia msp. 2</u>	-	-	-	-	-	-	-	<u>0.07</u>	
◀ <u>Dicopia msp.</u>	<u>0.27</u>	-	-	-	-	-	-	-	
<u>Pyuridae</u>									
◀ <u>Culeolus msp.</u>	-	-	-	-	-	-	-	-	<u>0.07</u>
<u>Tunicata indet.</u>	<u>0.14</u>	<u>0.04</u>	<u>0.08</u>	<u>0.11</u>	-	-	-	-	-
▶									
◀ <u>Paleodictyon nodosum</u>		-	-	-	-	-	-	<u>0.06</u>	-

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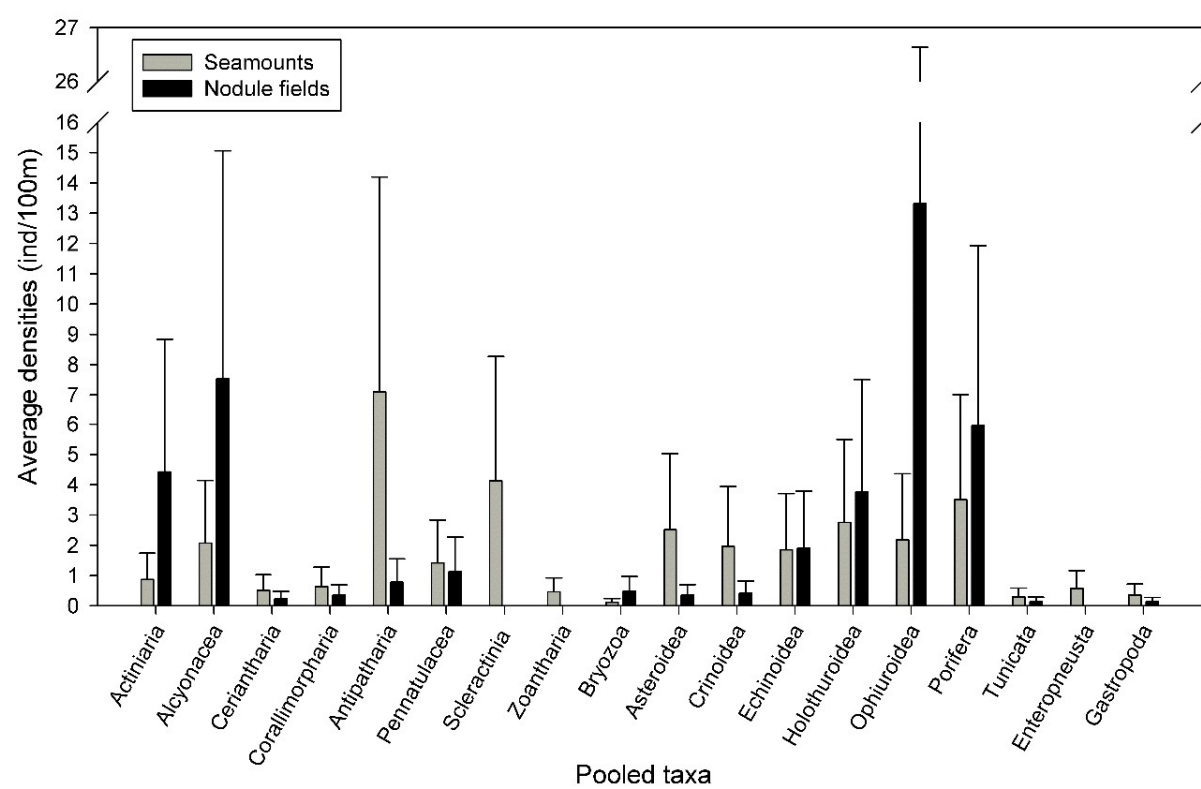
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Fig. A1. Average densities at higher taxa level per ecosystem and standard deviation.

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