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Macrofaunal assemblages from mud volcanoes in the Gulf of Cadiz: abundance, biodiversity and diversity partitioning across spatial scales

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Abstract

The Gulf of Cadiz is an extensive seepage area in the South Iberian Margin (NE Atlantic) encompassing over 40 mud volcanoes (MVs) at depths ranging from 200 to 4000 m. The area has a long geologic history and a central biogeographic location with a complex circulation assuring oceanographic connectivity with the Mediterranean Sea, Equatorial and Northern Atlantic regions. The geodynamics of the region promotes a notorious diversity in the seep regime despite the relatively low fluxes of hydrocarbon-rich gases. We analyse quantitative samples taken during the cruises TTR14, TTR15 and MSM01-03 in seven mud volcanoes grouped into Shallow MVs (Mercator: 350 m, Kidd: 500 m, Meknès: 700 m) and Deep MVs (Captain Arutyunov: 1300 m, Carlos Ribeiro: 2200 m, Bonjardim: 3000 m, Porto: 3900 m) and two additional reference sites (ca. 550 m). Macrofauna (retained by a 500 μm sieve) was identified to species level whenever possible. The samples yielded modest abundances (70–1567 ind. per 0.25 m²) but a number of species among the highest ever reported for cold seeps. Among the 366 recorded species, 22 were symbiont-hosting bivalves (Thyasiridae, Vesicomomyidae, Solemyidae) and tubeworms (Siboglinidae). The multivariate analyses indicated significant differences between Shallow and Deep MVs: The environmental conditions at the Shallow MVs makes them highly permeable to the penetration of background fauna leading to high diversity of the attendant assemblages (H' : 2.92–3.94; $ES_{(100)}$: 28.3–45.0; J' : 0.685–0.881). The Deep MVs showed, in general, contrasting features but were more heterogeneous (H' : 1.41–3.06; $ES_{(100)}$: 10.5–30.5; J' : 0.340–0.852) and often dominated by one or more siboglinid species. The rarefaction curves confirmed the differences in biodiversity of Deep and Shallow MVs as well as the approximation of the latter to the reference sites. The Bray–Curtis dissimilarity demonstrated the high β -diversity of the assemblages, especially in pairwise comparisons involving samples from the deeper MVs. Diversity partitioning assessed for species richness, Hurlbert's expected number of species and Shannon–Wiener index confirmed the high β -diversity across different spatial scales (within MVs, between

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MVs, between Deep and Shallow MVs). We suggest that historical and contemporary factors with differential synergies at different depths contribute to the high α -, β - and γ -diversity of the mud volcano faunal assemblages in the Gulf of Cadiz.

1 Introduction

5 After only a few decades since the discovery of cold seeps and their associated faunal assemblages (Paull et al., 1984; Suess et al., 1985; Juniper and Sibuet, 1987), it is now well established that these are immensely diverse and widely distributed reducing marine environments spanning from coastal to hadal depths in different geological settings of passive and active margins (e.g. Levin, 2005; Cordes et al., 2010b; Levin and Sibuet, 2012). Along the European Atlantic and Mediterranean margins there are three main regions characterized by the presence of seep-related structures (mud volcanoes, pockmarks, brine pools): the Nordic margin, the South Iberian region and the Eastern Mediterranean (Foucher et al., 2009; Vanreusel et al., 2009).

15 The South Iberian region, probably the one that has been less systematically investigated, in fact also encompasses the Moroccan margin and it is a complex of two distinct but tectonically and oceanographically related areas – the Alboran Sea (West Mediterranean) and the Gulf of Cadiz (NE Atlantic). Evidence for the presence of Jurassic oceanic crust in the central Gulf of Cadiz (Sallarès et al., 2011) indicates that the geologic history of the region is probably one of the most ancient in the NE Atlantic. The geodynamic setting is determined by two plate driving mechanisms: (i) subduction 20 (West Mediterranean) associated with the formation of an accretionary wedge (Gulf of Cadiz) which is thought to be inactive at present and (ii) oblique collision between Iberia and Nubia, presently active and causing thrusting and dextral wrenching along the Azores-Gibraltar Fault Zone (Zittelini et al., 2009). The present oceanographic setting is dominated by the exchanges through the Strait of Gibraltar (Ambar et al., 2002) 25 and the Gulf of Cadiz is characterized by the formation of meddies and recirculation of the Mediterranean outflow water. This intermediate layer between the surface and

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the Central North Atlantic water masses interacts with the seafloor at the upper slope (Quentel et al., 2011) while at the deeper western regions there is evidence for input of high-nutrient Antarctic Intermediate Water (Van Aken, 2000). These interactions are reflected in varying sea bottom current velocities and temperatures (from 4 °C at the deeper locations and up to 13 °C if under the influence of the Mediterranean outflow water).

It is now well established that mud volcanism and other processes associated with the escape of hydrocarbon-rich fluids in the Gulf of Cadiz sustain a broad diversity of chemosymbiotic organisms (Rodrigues et al., 2008, Hilário and Cunha, 2009; Hilário et al., 2010; Oliver et al., 2011; Rodrigues et al., submitted to this issue). This extensive seepage area encompasses over forty mud volcanoes that pierce the accretionary wedge at depths ranging from 200 to 4000 m. Flow rates are relatively low (0.05–15 cm yr⁻¹; Hensen et al., 2007; Vanneste et al., 2011) and major active mud volcanoes are located along strike-slip faults (Pinheiro et al., 2003) which constitute a feeder system, acting as main upward conduits for the deep-sourced overpressurized fluids and hydrocarbon gases. The deep hydrothermal alteration of basement rocks of the ocean-continent transition crust may be involved in the production of these fluids (Nuzzo et al., 2009). Although predominantly deep and thermogenic (Stadnitskaia et al., 2006; Hensen et al., 2007; Nuzzo et al., 2009) different sources of methane and transport processes determine a highly variable pore fluid geochemistry at different MVs (Niemann et al., 2006; Nuzzo et al., 2009; Scholz et al., 2009, 2010), and even at different locations of a single mud volcano (Sommer et al., 2009; Vanneste et al., 2011). Even at the deeper mud volcanoes the fluxes of methane are relatively low (up to 407 mmol m⁻² yr⁻¹, Niemann et al., 2006; 0–806 mmol m⁻² yr⁻¹, Vanneste et al., 2011) and its concentrations in surface sediments are generally < 0.001 mM (Niemann et al., 2006). The complete depletion of methane and sulphate within the subsurface matches elevated rates of sulphate-dependent anaerobic oxidation of methane (AOM); an efficient microbial filter and, in some cases, methanotrophic symbiont-hosting invertebrates consume a large proportion of the ascending methane, and directly and

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indirectly affect methane turnover (Sommer et al., 2009). These environmental setting variations and AOM activity are reflected by diverse microbial community compositions (Vanreusel et al., 2009), and most symbiont-hosting invertebrates in the Gulf of Cadiz are thiotrophic (Rodrigues et al. 2010, 2011a).

5 The widespread occurrence of massive authigenic carbonate chimneys and crusts typically at depths between 600 and 1200 m along the margins of Morocco and Spain suggests that this was a very active seepage area in the past (Wehrmann et al., 2011; Magalhães et al., 2012; Vanneste et al., 2012). In these carbonate provinces there are frequent observations of mounds, thickets and debris of mostly dead cold-water
10 scleractinean corals (Wienberg et al., 2009) and also extensive Buccinid and Mytilid graveyards (usually within the crater of the mud volcanoes).

The aim of this paper is to describe patterns in biodiversity, abundance and community structure of the benthic macrofaunal assemblages in seven mud volcanoes along a bathymetric gradient in the Gulf of Cadiz. We analyse quantitative samples taken
15 during the cruises TTR14 and TTR15 onboard the RV Prof. Logachev (Kenyon et al., 2006; Akhmetzhanov et al., 2007), and MSM01-03 onboard the RV Maria S Merian. These patterns are analysed and discussed in relation to the present knowledge on the environmental conditions of the study locations.

2 Material and methods

2.1 Study sites

20 The study sites are seven mud volcanoes (MVs) located along a bathymetric gradient ranging approximately from 350 m to 4000 m water depths (Fig. 1).

The shallowest sites, Mercator (350 m) and Kidd MVs (490 m), belong to the El Ar-raiche field at the upper slope of the Moroccan margin (Van Rensbergen et al., 2005).
25 Mercator MV is one of the two sites in the Gulf of Cadiz where active gas bubbling has been observed (the other one being Darwin MV; Vanreusel et al., 2009); the porewater

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is characterized by an extreme enrichment of Cl^- and SO_4 consistent with the dissolution of minerals (e.g. halite and gypsum), and Li concentration; radiogenic $^{87}\text{Sr}/^{86}\text{Sr}$ ratios are consistent with a terrigenous/continental deep source of the fluids (Scholz et al., 2010). The carbon isotopic signature of methane indicates high thermal maturity (Stadnitskaia et al., 2006; Hensen et al., 2007; Nuzzo et al., 2009). According to the scheme of dominant gas production and transport processes proposed by Nuzzo et al. (2009) Mercator MV is an example of admixture of thermogenic gases from different depths and influence of evaporate deposits. Meknes MV is located in the Moroccan carbonate province (700 m). The porewater is characterized by a moderate depletion of Cl^- (Hensen et al., 2007; Scholz et al., 2010). The carbon isotopic signature of methane is more depleted than in Mercator MV suggesting lower thermal maturity (Nuzzo et al., 2009).

Captain Arutyunov MV is located at ca. 1300 m water depth (just below the influence of the MOW) and seats on one of the major SWIM (SW Iberian Margin) lineaments, where the thickness of the accretionary prism is greatest (14 km; Thiebot and Gutscher, 2006). Both the thickness of the sediments and the low thermal gradient are thought to promote the production of large amounts of oil and hydrocarbon gases (Nuzzo et al., 2009 and references therein). The geochemistry is highly variable showing moderate Cl^- enrichment, and often extremely steep porewater gradients and high turnover rates coinciding with a shallow sulfate-methane transition zone (Niemann et al., 2006; Hensen et al., 2007; Sommer et al., 2009; Scholz et al., 2010). The Capt. Arutyunov MV is categorized as an example of extensive generation and transport of fluids, petroleum and hydrocarbon gases from deeply-buried sediments (Nuzzo et al., 2009).

The thickness of the accretionary prism decreases westwards where Carlos Ribeiro (ca. 2200 m) and Bonjardim (ca. 3000 m) are also aligned with major thrust faults. The geochemistry of these two MVs is very similar displaying the advection of strongly hydrocarbon-enriched fluids (Nuzzo et al., 2009; Vanneste et al., 2011). The carbon isotopic signature of methane indicates low thermal maturity (Stadnitskaia et al., 2006; Nuzzo et al., 2009). The fluids show a distinctly depleted chlorinity that results from

clay mineral dehydration promoted by positive heat flux anomalies (Nuzzo et al., 2009 and references therein; Scholz et al., 2010). Carlos Ribeiro and Bonjardim MVs are categorized as examples of active generation of C₂+ -enriched gases under tectonic control (Nuzzo et al., 2009).

Porto MV is located at the rim of the accretionary prism at ca. 3900 m; the values ⁸⁷Sr/⁸⁶Sr are the least radiogenic and may be attributed to admixing of a basement-derived fluid component carrying an isotopic signature from interaction with the basaltic (oceanic) crust (Scholz et al., 2009, 2010). A relevant characteristic distinguishing Porto from the other deep MVs is that the isotope signatures suggest partial recycling of thermogenic methane mediated by AOM-related methanogenic archaea in the shallow sediments; subsequently Nuzzo et al. (2009) indicate Porto MV as a possible example of deep acetoclastic methanogenesis.

2.2 Collection and processing of samples

In most cold seep studies different habitats, such as tubeworm fields, mussel beds and bacterial mats, are identified and sampled discretely. In the Gulf of Cadiz the chemosymbiotic species are usually inconspicuous or burrowed and in most cases it is practically impossible to use visual cues for targeted sampling. Herein we present data from samples collected in locations chosen following video surveys of the study sites, but using non TV-assisted gear. Whenever applicable, observations regarding the dominant biological facies of each sample are given in Table 1. During the cruise TTR 14 the macrofaunal assemblages in Kidd MV were sampled at the crater and at the flank; an additional sample was taken at some distance away of the MV for reference. The same procedure was carried out at Mercator MV during the cruise TTR15. During the cruise MSM01-03, six mud volcanoes were sampled totalling 10 individual samples that were taken mostly from the crater area (Fig. 1, Table 1). In all cases a USNEL box corer (area = 0.25 m²) was used and the sample processing was initiated onboard by sub-sampling the different depth layers of the sediment. The material was sliced at 0–1, 1–3, 3–5, 5–10, 10–20 cm following the standard techniques adopted as

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general practice in HERMES and HERMIONE cruises. The supernatant water on box cores was passed through a 500 μm sieve to retain any small fauna resuspended during sampling and core retrieval on deck. The sediment layers were carefully washed with seawater through 2 mm, 1 mm and 500 μm sieves. The sieved material was fixed immediately in 96 % ethanol. In the lab, the material was sorted under a stereo microscope into major taxa and kept in 96 % ethanol, for taxonomic identification to species level whenever possible. Many taxa were not yet matched with a species name but they were all sorted into putative species and ascribed a codename consistent throughout the samples. Therefore the estimated species richness can be considered accurate, although minor changes may be made in the future following more thorough taxonomic revisions of the material. Species abundances were determined for each sediment layer in each sample, but for the purposes of this paper sediment layers' values were not used and densities were calculated as the number of individuals in relation to the area sampled and expressed as individuals per m^2 (ind m^{-2}). Special attention was given to the count of Siboglinidae tubeworms (since the tube of many of them is brittle and the number of tube fragments generally does not correspond to the number of animals). All tube fragments were dissected and only those containing the forepart of the animal were considered.

2.3 Data analysis

Species richness, Shannon–Wiener diversity index (H'), Pielou (1969) evenness index (J') and Hurlbert (1971) expected species richness ($ES_{(n)}$), were calculated using the community analysis PRIMER v6 software (Clarke and Gorley, 2006). This software was also used for multivariate analyses. The abundance data were first organized into a sample vs species matrix and non-metric multidimensional scaling (nMDS) ordination was performed using the Bray–Curtis similarity measure, after square root transformation. An analysis of similarities by randomization/permutation tests (ANOSIM) was performed on the MDS results of samples to assess for differences in the assemblages putatively related to the bathymetric variation in environmental conditions.

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Because the lack of replication would not allow for testing individual MVs these were grouped into shallow and deep MVs. Based on the characteristics of the study area, the 1200 m water depth was used as the transition between two very different sets of oceanographic and geochemical conditions. The samples were grouped into the following three categories: R – reference sites in the vicinity of Mercator and Kidd; SMVs – shallow MVs including Mercator, Kidd and Meknès; DMVs – deep MVs including Captain Arutyunov, Carlos Ribeiro, Bonjardim and Porto. SIMPER analysis (Similarity Percentages – species contributions) was performed to assess the percentage contributions of each species to the similarity within and dissimilarity between groups of samples.

For the analysis of β -diversity we used a dissimilarity matrix based on Bray-Curtis measure after square-root transformation. In order to describe different contributions for the overall β -diversity, the pairwise comparison values (pwc) were grouped into seven categories: (i) WSMV – pwc between samples within the same MV in the Shallow field; (ii) BSMV – pwc between samples from different MVs in the Shallow field; (iii) WDMV – pwc between samples within the same MV in the Deep field; (iv) BDMV – pwc between samples from different MVs in the Deep field; (v) BS&DMV – pwc between shallow and deep MVs; (vi) BR&SMV – pwc between reference sites and shallow MVs; (vii) BR&DMV – pwc between reference sites and deep MVs.

Diversity partitioning was assessed for species richness (S), Hurlbert's expected number of species ($ES_{(100)}$) and Shannon-Wiener index (H'). The total diversity ($\gamma = \alpha + \beta$) is partitioned into the average diversity within samples (α) and among samples (β) and therefore β -diversity can be estimated by $\beta = \gamma - \alpha$ (Wagner et al., 2000; Magurran, 2004). To extend the partition across multiple scales (β_1 : within MV, β_2 : between MVs and β_3 : between MV fields), the smallest sample unit for level 1 are individual boxcore samples, while for the upper levels sampling units are formed by pooling together the appropriate groups of nested samples. The diversity components are calculated as $\beta_m = \gamma - \alpha_m$ at the highest level and $\beta_j = \alpha_{i+1} - \alpha_i$ for each lower level. The

additive partition of diversity is

$$\gamma = \alpha_1 + \sum_{i=1}^m \beta_i j_m \quad (1)$$

The total diversity can therefore be expressed as the percentage contributions of diversity in each hierarchical level (Crist et al., 2003). Partitioning was carried out ascribing the same weight for each sample as they all represent the same area sampled (alternatively samples can be weighted by abundance; see Crist et al., 2003). Values of α_i were therefore calculated as a weighted average (according to the number of samples pooled).

3 Results

A total of 7036 specimens were ascribed to 366 putative species from which 111 (30.0%) were singletons. Annelida (52.2% of total abundance; 136 species) and Arthropoda (33.5%; 152 species) were the most abundant and the most species-rich Phyla. Mollusca and Echinodermata, although common, showed much lower values of abundance and species richness (8.4%; 56 species and 3.6%, 12 species, respectively). These four groups were represented in all samples except St. 218 in Cap. Arutyunov MV (containing shallow gas hydrates), where crustaceans and echinoderms were absent. Although present in some samples, sponges and cnidarians were not included in this study.

3.1 Symbiont-hosting species

The sixteen samples studied yielded a total of twenty symbiont-hosting species belonging to the bivalve families Thyasiridae (*Thyasira vulcolutre*), Vesicomidae (*Isorropodon megadesmus*; *Isorropodon perplexum*) and Solemyidae (*Acharax gadirae*; *Solemya elarraichensis*) and the polychaete family Siboglinidae (*Lamelisabella denticulata*;

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Spirobrachia tripeira; *Siboglinum* cf. *poseidoni*; and eight other *Siboglinum* species, and four species of *Polybrachia* ascribed according to Hilário et al., 2010). Seven chemosymbiotic species occurred in the shallow MVs (0–2 species per sample) and 16 in the deep MVs (up to five different species per sample; Fig. 2a). Additionally the mixotroph bivalves *Axinulus croulinensis* and *Thyasira granulosa* were also present in the shallow MVs but because they are facultative heterotrophs they are included in the section below. Details on the distribution of chemosymbiotic species are given by Oliver et al. (2011) and Rodrigues et al. (2012). The abundance of chemosymbiotic species was highly variable (Fig. 3a) but, in average, lower in the Shallow MVs than in the Deep MVs (Average \pm SE; Siboglinidade: 221.3 ± 112.1 and 1578.4 ± 939.0 ind m^{-2} ; Bivalves: 1.3 ± 1.3 and 135.2 ± 127.2 ind m^{-2} , respectively). Chemosymbiotic species were often the dominant taxa in the Deep MVs, accounting for up to 79.5 % of the total abundance. The highest number of individuals occurred in the two samples taken at Cap. Arutyunov MV: St. 218 with the highest abundance of bivalves and the second highest abundance of siboglinids (161 and 567 ind. per $0.25 m^2$) and St. 180 with the highest abundance of siboglinids (1240 ind per $0.25 m^2$). A humbler maximal siboglinid count in the Shallow field was recorded in St. 242 at Mercator MV (261 ind per $0.25 m^2$). Note that siboglinid counts refer only to occupied tubes and animals found out of the tubes; empty tubes can easily reach densities one order of magnitude higher.

3.2 Heterotrophic species

As for the remaining fauna, and in comparison with the two reference sites (51.5 ± 9.5 species per sample and 660 ± 60.0 ind m^{-2}), the Shallow MVs showed increased species richness and abundance (68.7 ± 4.4 species per sample and 1387.6 ± 860.0 ind m^{-2}) while the Deep MVs showed decreased species richness and only slightly higher abundance (29.4 ± 7.9 species per sample and 860.6 ± 339.7 ind m^{-2}). The highest number of species and individuals of heterotrophic species was found in St. 241, an area with solitary corals (*Caryophyllia* sp.) at Mercator MV (98 species and 665 ind.), the lowest species richness in St. 218 in association with dissociating gas

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hydrates at Capt. Arutyunov MV (16 species), and the lowest abundance in St. 133, an area covered by shell ash at Bonjardim MV (52 ind.) (Figs. 2b and 3b).

The composition of the MV assemblages at the level of major taxonomic groups (chemosymbiotic species excluded) does not reveal clear trends in relation to the Reference sites. In terms of abundance (Fig. 3b), the relative contribution of polychaetes is increased in Mercator, Cap. Arutyunov and Bonjardim MVs, and decreased in Meknès and Porto MVs; overall, crustaceans dominate most MV assemblages except for St. 218 at Cap. Arutyunov MV (0%), and are particularly abundant in Meknès (in average the amphipods account for over 40% of the abundance), Carlos Ribeiro (isopods: 45%) and Porto MVs (isopods: 39%); the contribution of molluscs is decreased in Kidd and Meknès MVs and greatly increased in Capt. Arutyunov MV; finally, the abundance of ophiuroids in Mercator MV and sipunculids in Kidd MV are also relevant. In terms of species richness (Fig. 2b) the relative contribution of *Scolecida* (*Polychaeta*) is often impoverished in the MV assemblages (e.g. Meknès, Carlos Ribeiro, Bonjardim, Porto MVs) while the opposite occurs with the gastropods (Mercator, Kidd, Capt. Arutyunov MVs) and crustaceans (Kidd, Meknès, Carlos Ribeiro and Porto MVs); decapods (e.g. the small crabs *Cymonomus granulatus*, *Ebalia nux*, *Monodaeus couchi*) were found exclusively in Mercator and Meknès MVs (7 and 2 species, respectively).

3.3 Comparison between Shallow and Deep MVs

The MDS plot (Fig. 4) based on the whole dataset (chemosymbiotic and heterotrophic species) reflects well the variability in the composition of the assemblages. It shows a high dispersion of the heterogeneous Deep MVs samples along the first axis, and the Shallow MVs samples closer together and clustering per MV, but spreading along the second axis in between the reference sites and the Deep MVs. The results of the ANOSIM test support a significant overall difference between sites (Global R: 0.618; 0.1% significance level) as well as for the pairwise comparisons between Shallow MVs and Reference sites (R: 0.517; Significance level: 3.6%; all 55 possible permutations used) and between Shallow and Deep MVs (0.674; 0.2% 999 permutations). The

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comparison between Deep MVs and Reference sites retrieved a non-significant value (0.400; 14.3 %) owing to the high heterogeneity of Deep MVs samples combined with the low number of possible permutations (21).

In common, the assemblages from the Shallow MVs have higher biodiversity and evenness (H' : 2.92–3.94; $ES_{(100)}$: 28.3–45.0; J' : 0.685–0.881; k_5 : 28.2–60.5 %). The assemblages from the Deep MVs showed, in general, contrasting features but were more heterogeneous (H' : 1.41–3.06; $ES_{(100)}$: 10.5–30.5; J' : 0.340–0.852; k_5 : 49.7–89.7 %; Table 2); the assemblage from the siboglinid fields in Porto MV, with the highest evenness and biodiversity among the Deep MVs was particularly distinct.

The polychaetes *Paradoneis lyra*, *Lauberiopsis cabiochi*, the tanaids *Fageapseudes retusifrons*, and *Atlantapseudes nigrichela*, and the ophiuroids *Amphiura filiformis* and *Amphipholis squamata* were frequently found among the dominant species in the Shallow MVs while the Deep MVs were often dominated by different chemosymbiotic species. The dominant species are generally retrieved by the SIMPER analysis as major contributors for the similarity and dissimilarity of the groups of samples (Supplement). The major contributors for the average similarity of the Reference sites are detritivores such as *Scolecida polychaetes* (*Aricidea simonae*, Capitellidae sp005, *Ophelina* cf. *cylindricaudata*, *Paradoneis lyra*), several crustaceans (*Ampelisca* sp022, *Harpinia* sp046, *Leucon* sp001, *Sphyrapus malleolus*), the ophiuroid *A. squamata* and the bivalve *Microgloma tumidula*. In the Shallow MVs major contributors are again detritivores but also include carnivore/omnivore polychaetes (e.g. *Glycera lapidum*, *Paradiopatra hispanica*), and several crustaceans with a more opportunistic diet (caprellid amphipods, apseudomorph tanaids). In the Deep MVs the detritivore contributors are mostly asellote isopods and tanaidomorph tanaids; the polychaete contributors are mostly Aciculata species (carnivores/omnivores) and siboglinids (chemosymbiotic). The siboglinids are also major contributors for the Shallow/Deep and Reference/Deep average dissimilarities. The Reference/Shallow average dissimilarity is explained mostly by crustaceans (mainly amphipods). Overall, SIMPER results are determined by the high variability in the composition of the assemblages. The average

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similarities for the three groups Reference, Shallow and Deep are very low (20.1, 27.6 and 12.8%, respectively) because there are many singletons and few common species among samples. For the same reason, the average dissimilarities for the comparisons between groups are very high (R/S: 81.4%; R/D: 92.8%; S/D: 87.8) and are accounted for by a large number of species with very low percentage contributions.

3.4 Biodiversity analyses

The rarefaction curves (Fig. 5) all show relatively steep slopes (partly explained by the large number of singletons) and are far from reaching asymptotic values revealing that the sampling effort is insufficient to assess the full biodiversity either at individual MVs or at larger scales. These curves also demonstrate the contrasting biodiversity of the Deep and Shallow MVs as well as the approximation of the latter to the reference sites, both at the scale of the individual MVs (Fig. 5a) and at the regional scale (Fig. 5b). The β -diversity analysis (Fig. 6) confirms the high variability of the faunal assemblages: the lowest dissimilarity obtained for the pairwise comparison between two samples of the Meknès MV was rather high (49.3%). Complementarity increases progressively for pairwise comparisons involving Shallow MVs – from comparisons within the same MV to between Shallow and Deep MVs (WSMV < BSMV < BR&SMV < BS&DMV). Comparisons involving Deep MVs showed very high dissimilarity values and widely overlapping ranges in the different categories (WDMV, BDMV, BS&DMV, BR&DMV). St. 218 at Capt. Arutyunov MV was particularly distinct – pairwise comparison involving this station showed consistently the highest dissimilarity values (95.5–100% between St. 218 and stations from other MVs; 82.6% between St. 218 and St. 180 also in Capt. Arutyunov).

The partitioning of diversity (Fig. 7) shows that only 15.4% of the total species richness is due to α -diversity (the average number of species per sample is 56.5 for a grand total of 366) and that most of the β -diversity is due to differences in species composition between MV fields (β_3 : 43.2%) and between individual MVs (β_2 : 27.6%) – this partition can be explained by the high number of singletons, low number of species shared

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by two or more MVs and high rate of species replacement along the depth gradient. When the Shannon–Wiener index is calculated the abundance is taken into account. The dominant species have more weight for explaining the partition of diversity – in this case the few samples showing high dominance (e.g. St. 180 and St. 218 in Capt. Arutyunov) do not affect much the average sample α -diversity (2.99; 70.3% of the total H' diversity) but they are sufficiently important to explain the relatively high contributions of β_1 and β_3 (11.2 and 11.7%, respectively). These levels are the most impacted because the dominant species are not the same in the samples from the same MV (level β_1) and because of the disproportionate effect of the abundance (and dominance) of the Capt. Arutyunov samples in the estimates of H' for the Deep MV field. Rarefaction ($ES_{(100)}$ estimates) somehow balances the weights of singletons and dominant species showing a partition of diversity with still high but intermediate contribution of α -diversity (60.3%) and a major contribution of level 2 (between MVs; β_2 : 22.3%) for β -diversity. Note that in the partition of diversity the expected contribution of α -diversity is higher for S than for $ES_{(100)}$ or H' (Crist et al., 2003) – the number of species is cumulative with increasing scales (always higher for larger scales) while in the case of $ES_{(100)}$ or H' individual samples can yield values that are close or even higher than the estimates for pooled samples.

4 Discussion

4.1 Faunal assemblages in the Gulf of Cadiz and other cold seeps

The number of species reported here either globally (366 species) for the Gulf of Cadiz (GoC) or for each individual MV (21–76 species in the Deep MVs; 97–188 in the Shallow MVs) is certainly among the highest ever reported for cold seeps. There are very few cold seep studies with species level taxonomic resolution to enable comparisons of species richness or other biodiversity indices (H' , $ES_{(100)}$) between different regions and the usual discrepancies in methodologies further complicate such comparisons.

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Levin et al. (2010) provide counts of up to 55 and 88 species from clam beds in Hydrate Ridge and Eel River (NE Pacific), respectively, at depths comparable to the Shallow MVs in GoC. Eleven locations at depths between 1005 and 2750 m in the Gulf of Mexico yielded a total of 88 taxa and $ES_{(100)}$ values ranging 7.7–15.4 (Cordes et al., 2010a). Other reported $ES_{(100)}$ estimates for cold seeps vary between 8 and 36 (Levin, 2005; Levin et al., 2010). The values observed in the Deep MVs of the GoC (10.5–30.5) are within this range but the ones in the shallow MVs were consistently higher (mostly over 36 and up to 45).

On the other hand, the average densities of the MV assemblages (1610 and 2466 indm^{-2} for the Shallow and Deep MVs, respectively) in the GoC are far from the reported maximal values for other cold seeps: 93 000 indm^{-2} from *Oligobrachia haakonmosbiensis* fields in the Håkon Mosby MV (Decker et al., 2012) and over 20 000 indm^{-2} from *Calyptogena* beds in Hydrate Ridge (Levin, 2005), mytilid beds in Regab (Menot et al., 2010) and reduced sediments from Amsterdam MV in the East Mediterranean (Ritt et al., 2012). However, they are in line with reported values from thyasirid beds in the North Sea (Dando et al., 1991), *Acharax* beds at Hydrate Ridge (Sahling et al., 2002) and several habitats from Napoli MV in the East Mediterranean (Ritt et al., 2012).

High variability of abundances between different habitats has been reported in most of these regions but the comparisons between different studies and regions are often confounded by methodological aspects (e.g. sampling gear, mesh size), and by the different bathymetric and biogeographic settings of different study areas. Although there is apparently no clear relationship to the geochemistry of the fluids (Levin, 2005), the huge variability in the densities reported for cold seeps must result from the influence of symbiont-hosting foundation species and the availability of colonisers determined by depth and biogeography operating on the numerous combinations of geological drivers, fluid flux, and biogeochemical processes occurring in these ecosystems.

Methanotrophic symbiont-hosting species (e.g. *Siboglinum poseidoni*) may significantly modify sediment geochemistry by promoting sulfate reduction which has a dual

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effect on the macrofaunal assemblages: by increasing sulfide porewater concentration while diversifying niches for microbial communities, they increase toxicity for the infauna but also diversify sources of food derived from microbial activity (Levin et al., 2003; Sommer et al., 2009). The predicted outcome is a macrofaunal assemblage highly dominated by sulfide tolerant heterotrophic species (Levin and Mendoza, 2007) and /or thiotrophic species which is in agreement with the high abundance and low evenness values observed in siboglinid-dominated assemblages from the GoC. For instance, localized optimal conditions for the establishment of *Siboglinum* spp. fields (Sommer et al., 2009) lead to maximal densities and lowest evenness occurring in Capt. Arutyunov MV. In Porto MV, the siboglinid fields are characterised by varying densities of the larger-sized species *Lamelisabella denticulata* and *Spirobrachia tripeira*. Their tubes protruding from the sediment, often as clumps, host a diverse epifauna, and the macrofaunal assemblage here showed high evenness and $ES_{(100)}$ values. The morphology and physiology of frenulate tubeworms, capable of overcoming low fluid flow conditions (Sommer et al., 2009) enables them to form dense populations under a very wide range of geochemical conditions and endure high temporal and spatial variability. Apparently, they may also yield very different attendant macrofaunal assemblages.

In the deep-sea, benthic macrofaunal assemblages are highly dependent on allochthonous organic matter (Rex et al., 2006; Smith et al., 2008). Chemosynthesis-derived organic matter and free-living chemolithotrophic bacteria and archaea available at cold seeps constitute a relevant addition to the nutrition of heterotrophic organisms (Carney, 1994). As in many other locations (e.g. review by Levin, 2005; Levin et al., 2010; Menot et al., 2010), in the GoC we observed increased abundance at seeps in comparison to reference sites at comparable depths. There were no reference samples for the deeper sites which yielded generally lower abundance than the shallow MVs. Diversified food sources may explain the increased abundance (and biomass) in seeps relative to reference sites and an increasing dependence on chemosynthetic food sources at deeper seeps as the availability of photosynthetically derived material decreases (Sahling et al., 2003; Levin and Sibuet, 2012). This pattern is very clear in

the GoC by the increasing prevalence of chemosymbiotic species in the assemblages of the Deep MVs both in terms on species richness and abundance and by the occurrence of mixotrophs exclusively in the Shallow MVs. Also interesting are the changes in the trophic structure of the attendant heterotrophic assemblages in cold seeps. In general, most deep-sea organisms are deposit feeders (Jumars and Wheatcroft, 1989) but we observed higher trophic diversity in the GoC MV assemblages. The available bacterial biomass and dissolved or particulate organic matter favours increased contributions of grazers such as gastropods putatively feeding on free-living bacteria (e.g. in Mercator, Kidd and Capt. Arutyunov MVs). Other opportunistic feeders such as corophioid amphipods and apseudomorph tanaids (mostly in Shallow MVs) use their appendages to collect detritus from the sediment but also may use a secondary filter feeding strategy (Larsen, 2005). Because the high density assemblages in the MVs may support higher trophic levels, omnivores and carnivores or scavengers such as Aciculata polychaetes (both in Shallow and Deep MVs) and small crabs (Shallow MVs) are also well-represented by relatively common species (e.g. *Glycera lapidum*, *Paradiopatra hispanica*, *Monodaeus couchi*). Grazers and non-lethal predators are customary in many seeps (e.g. review by Sibuet and Olu, 1998; Olu-Le Roy et al., 2004) but the limited foraging options due to defensive and/or environmental toxicity in cold seeps and a relative lack of deep-sea candidates for evolutionary specialized predation makes this option selectively disadvantageous (Carney, 1994).

4.2 Depth-related variability in faunal assemblages

Depth-related changes in the abundance, biodiversity and composition of cold seep assemblages are still poorly known (Levin and Sibuet, 2012). Besides the work by Sahling et al. (2003) in the Sea of Okhotsk covering a depth range from 160 to 1600 m and the work by Cordes et al. (2010a) in the Gulf of Mexico comparing the assemblages from eleven locations between 1005 and 2750 m with shallower sites, there are few studies attempting to infer depth-related effects on cold seep assemblages. Also, detailed accounts of seep macrofauna focus mainly on shelf or upper slope sediments and there

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is few detailed information on the macrofauna of seeps deeper than 3000 m (Levin and Mendoza, 2007). With over 40 known mud volcanoes spanning shelf and abyssal depths, the Gulf of Cadiz provides an ideal setting for such studies. However, the lack of visual cues (conspicuous symbiont-host megafauna) for targeted sampling of different habitat types and the few opportunities for quantitative collections limited the options for replicate sampling and constrained our working hypothesis. Therefore, we focused our analysis on the differences between two groups of MVs, Shallow and Deep, separated by a transition depth of 1200 m coinciding with relevant changes in the oceanographic and geologic settings. Among the most striking environmental features are: (i) the influence of the Mediterranean Outflow Water impinging higher seawater temperatures and lower oxygen to the benthic assemblages of shallower MVs; (ii) the transition from continental-oceanic crust and thickness of the accretionary prism which are determinant for the dominant fluid sources and transport mechanisms; (iii) the SWIM faults acting as main upward conduits for the fluids at the Deep MV field; (iv) the occurrence of solid-phase methane hydrates at the Deep MVs.

The interpretation of the MDS plot suggests that some of the Shallow MV assemblages are more closely related to the Reference assemblages and the dispersion of Shallow and Deep MV samples along the vertical and horizontal axis, respectively indicate that the prevalent factors of change in each field may differ.

Shallow MVs show a milder seepage regime with low chemical toxicity. The higher seawater temperatures and the proximity to the coast and to the euphotic zone add to a greater input of photosynthetically-derived organic matter. These environmental conditions make the Shallow MVs highly permeable to the penetration of background fauna leading to high abundance and diversity of the attendant faunal assemblages. The occurrence of authigenic carbonates (Magalhães et al., 2012) and fossil cold water corals (Wienberg et al., 2009) increases habitat heterogeneity and diversify the suite of potential habitats for invertebrates (Cordes et al., 2010b). The trophic diversity is also enhanced (as mentioned above) and the chemosymbiotic species are thiotrophic or mixotrophic showing modest abundance and species richness.

degree of novelty in the fauna samples, high variability in the composition and structure of the assemblages likely driven by habitat heterogeneity, and subsequent high β -diversity. In different regions, the main differences between seep assemblages have been consistently related more closely to habitat type and or foundation symbiont – hosting species than to geographical distance or depth (Levin and Mendoza, 2007; Cordes et al., 2010a; Decker et al., 2012; Ritt et al., 2012) and most of the β -diversity has been allocated to habitat-related differences between the faunal assemblages. In the GoC, maybe because habitat types cannot be clearly discriminated, the dissimilarity is lowest for within Shallow MV comparisons and increases with increasing spatial scale. Again the Deep MVs show a different pattern with consistently high dissimilarities at all spatial scales. The partition of diversity also showed the relevance of β -diversity at the level of between MV fields (β_3 level) for the three diversity indicators analysed. β_3 level represented even the highest contribution to β -diversity in the partition of species richness which we attributed to the high rate of species replacement along the depth gradient.

4.3 A hotspot of biodiversity

The 366 species reported here for the GoC are only a fraction of the over 1000 taxa (unpublished data) estimated from quantitative and non-quantitative samples collected up to now in over 18 MV and other locations in adjacent habitats (carbonate chimneys and concretions, dead scleractinean cold water corals). Comprehensive studies on species distributions of major taxa have been published or are in preparation (Rodrigues et al., 2011b; Génio et al., submitted to this issue). A significant effort has been allocated to taxonomic work for species level identification including the description of new heterotrophic and chemosymbiotic species (Dworshack and Cunha, 2007; Błażewicz-Paszkowycz, 2011a, 2011b; Hilário and Cunha, 2009; Oliver et al., 2011; Rodrigues et al., 2008, 2011b), and molecular approaches (e.g. Génio et al., 2008) leading to the discovery of a high diversity of taxa with low genetic divergence (e.g. Hilário et al., 2010; Moura et al., 2011, 2012). In fact, the 22 chemosymbiotic species

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(including the two mixotrophic species) referred herein belong to a total pool of 32 species which derive at least part of their nutrition from chemolithotrophic production of microbial symbionts (Rodrigues et al., 2012). These numbers are remarkable and, up to now, have no match in any other cold seep region of the world.

5 Macrofaunal assemblages are known to respond in different ways and at different scales to structural complexity and environmental variability arising from habitat heterogeneity – abiotic sources of heterogeneity such as variability in fluid flow, geochemistry, and substrate type, which give rise to different sets of microbial communities, microbial symbiont-hosting foundation species, and attendant heterotrophic assemblages; and
10 biotic sources of heterogeneity derived from the biogenic habitats subsequently created by these species (Cordes et al., 2010b). As discussed in the previous sections, these sources of habitat heterogeneity are certainly relevant and may explain a high degree of β -diversity that has also been reported from many other cold seep regions. The specificity of the environmental conditions and putative isolation of cold seep habi-
15 tats also promote a varying degree of novelty in the faunal assemblages intrinsically contributing to α - and γ -diversity.

What are then the outstanding characteristics of the GoC?

- (a) The geodynamic setting of the region promotes a notorious diversity in the seep regime and because each MV in the GoC is characterized by a particular set of
20 environmental conditions they act as keystone physiographic structures contributing to a stepwise accumulation of species along the depth gradient.
- (b) The location of GoC is central to various biogeographic regions in the Atlantic and Mediterranean and this setting promotes the overlap of many species distributions extending their range into the GoC domain.
- 25 (c) These biogeographic regions are currently interconnected by different water masses under a regional oceanographic regime with high seasonal and interannual variability likely to provide numerous options for colonization of the seabed in the GoC.

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(d) The geologic history of the region is long (180–160 Ma ago) and there is evidence of a much more widespread and active seepage regime in the past which may have created a pool of evolutionary candidates for specialization to different seepage conditions.

(e) Paleoclimatic changes in the oceanographic regime and the decay of past seepage activity may have augmented the distance between active seeps within the GoC while present-day oceanographic retention mechanisms are also likely to promote genetic isolation and divergence especially of those species with lower dispersal capabilities.

These factors may have different synergies at different depth ranges and overall contribute to the high α -, β - and γ -diversity observed in the Gulf of Cadiz.

Supplementary material related to this article is available online at:

<http://www.biogeosciences-discuss.net/9/18331/2012/>

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Table 1. Metadata for the sampling stations in the Gulf of Cadiz. Station code: the label is composed by the cruise name and station number as used in Pangaea database (www.pangaea.de); for simplification in figures and text the stations are referred only by the three last digits.

| Location | Station code | Date | Lat. (N) | Long. (W) | Depth (m) | Observations |
|--------------------|--------------|------------|-------------|-------------|-----------|--|
| Reference | TTR15_AT577 | 2005.07.26 | 35° 17.305' | 06° 39.672' | 485 | Hemipelagic sediments |
| Reference | TTR14_AT559 | 2004.08.08 | 35° 24.777' | 06° 43.782' | 552 | Hemipelagic sediments |
| Mercator MV | TTR15_AT575 | 2005.07.26 | 35° 17.905' | 06° 38.715' | 355 | Crater, mud breccia |
| Mercator MV | TTR15_AT576 | 2005.07.26 | 35° 17.657' | 06° 39.129' | 428 | Flank |
| Mercator MV | MSM1/3.241 | 2006.05.06 | 35° 17.918' | 06° 38.687' | 353 | Crater, mud breccia, <i>Caryophylla</i> facies |
| Mercator MV | MSM1/3.242 | 2006.05.06 | 35° 17.870' | 06° 38.717' | 353 | Crater, mud breccia, <i>Hyalonecia</i> facies |
| Kidd MV | TTR14_AT560 | 2004.08.08 | 35° 25.306' | 06° 43.976' | 498 | Crater, mud breccia |
| Kidd MV | TTR14_AT561 | 2004.08.08 | 35° 25.602' | 06° 44.099' | 526 | Flank |
| Meknès MV | MSM1/3.319 | 2006.05.14 | 34° 59.100' | 07° 04.439' | 695 | Crater, stiff mud breccia |
| Meknès MV | MSM1/3.321 | 2006.05.14 | 34° 58.796' | 07° 04.394' | 732 | Seep periphery, carbonate and coral rubble |
| Meknès MV | MSM1/3.335 | 2006.05.15 | 34° 59.035' | 07° 04.552' | 703 | Crater rim, coral rubble |
| Capt. Arutyunov MV | MSM1/3.180 | 2006.04.27 | 35° 39.740' | 07° 19.960' | 1323 | Crater rim, buccinid facies |
| Capt. Arutyunov MV | MSM1/3.218 | 2006.04.30 | 35° 39.700' | 07° 20.012' | 1318 | Crater, mud breccia with gas hydrates |
| Carlos Ribeiro MV | MSM1/3.157 | 2006.04.23 | 35° 47.270' | 08° 25.360' | 2200 | Crater, mud breccia |
| Bonjardim MV | MSM1/3.133 | 2006.04.19 | 35° 27.821' | 09° 00.128' | 3049 | Crater, mud breccia, shell ash and frenulate tubes |
| Porto MV | MSM1/3.145 | 2006.04.21 | 35° 33.699' | 09° 30.437' | 3860 | Crater, mud breccia, <i>Lamelisabella</i> / <i>Spirobrachia</i> fields |

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Table 2. Abundance and biodiversity data for macrofaunal samples. n : number of deployments pooled; A : area sampled; N : number of individuals; D : density; SE: standard error; S : species richness; $ES_{(100)}$: Hulbert's expected number of species per 100 individuals; H' : Shannon-Wiener diversity (ln base); J' : Pielou's evenness; k_1 : percentage contribution of the first dominant species; R: Reference sites; Mr: Mercator MV; K: Kidd MV; Mk: Menès MV; CA: Captain Arutyunov MV; CR: Carlos Ribeiro MV; B: Bonjardim MV; P: Porto MV; SMV: Shallow MVs samples pooled; DMV: Deep MVs samples pooled; All: all samples pooled. Pooled values of S , $ES_{(100)}$ and H' were used for diversity partition estimates.

| Site | n | A (m ²) | N | $D \pm SE$ (ind m ⁻²) | S | $ES_{(100)}$ | H' (ln) | J' | k_1 (%) | Dominant species |
|------------|------------|--------------------------|-------------|--------------------------------------|------------|--------------|--------------|--------------|--------------|-----------------------------------|
| R | 577 | 0.25 | 152 | | 45 | 36.0 | 3.02 | 0.794 | 29.6 | Diastylidae sp003 |
| R | 559 | 0.25 | 178 | | 61 | 45.5 | 3.71 | 0.901 | 9.6 | <i>Microgloma tumida</i> |
| R | 2 | 0.50 | | 660 ± 52.0 | 93 | 50.5 | 3.96 | | | |
| Mr | 575 | 0.25 | 448 | | 76 | 37.5 | 3.45 | 0.796 | 15.0 | <i>Paradoneis lyra</i> |
| Mr | 576 | 0.25 | 307 | | 73 | 41.2 | 3.63 | 0.847 | 12.7 | <i>Fageapseudes retusifrons</i> |
| Mr | 241 | 0.25 | 720 | | 100 | 45.0 | 3.94 | 0.856 | 7.5 | <i>Paradoneis lyra</i> |
| Mr | 242 | 0.25 | 529 | | 71 | 31.5 | 2.92 | 0.685 | 28.9 | <i>Siboglinum splb</i> |
| Mr | 4 | 1.00 | | | 188 | 48.2 | 4.14 | | | |
| K | 560 | 0.25 | 200 | | 62 | 43.6 | 3.64 | 0.881 | 10.5 | <i>Atlantapseudes nigrichella</i> |
| K | 561 | 0.25 | 217 | | 64 | 42.8 | 3.63 | 0.873 | 11.5 | <i>Sipuncula sp002</i> |
| K | 2 | 0.50 | | | 97 | 48.1 | 3.96 | | | |
| Mk | 319 | 0.25 | 400 | | 51 | 28.3 | 2.98 | 0.758 | 20.0 | <i>Harpinia sp020</i> |
| Mk | 321 | 0.25 | 478 | | 71 | 36.8 | 3.49 | 0.819 | 13.6 | <i>Siboglinum splf</i> |
| Mk | 335 | 0.25 | 324 | | 62 | 37.2 | 3.49 | 0.845 | 10.5 | <i>Lembos sp037</i> |
| Mk | 3 | 0.75 | | | 115 | 40.7 | 3.78 | | | |
| SMV | 9 | 2.25 | | 1610 ± 218.7 | 280 | 56.6 | 4.58 | | | |
| CA | 180 | 0.25 | 1567 | | 64 | 14.9 | 1.41 | 0.340 | 72.5 | <i>Siboglinum spld</i> |
| CA | 218 | 0.25 | 1091 | | 21 | 10.5 | 1.65 | 0.541 | 51.4 | <i>Siboglinum cf. poseidoni</i> |
| CA | 2 | 0.50 | | | 76 | 17.3 | 2.05 | | | |
| CR | 157 | 0.25 | 70 | | 21 | – | 2.59 | 0.852 | 18.6 | Desmosomatidae und. |
| B | 133 | 0.25 | 198 | | 25 | 18.0 | 1.98 | 0.615 | 46.0 | <i>Siboglinum splc</i> |
| P | 145 | 0.25 | 157 | | 37 | 30.5 | 3.06 | 0.847 | 15.9 | Desmosomatidae und. |
| DMV | 5 | 1.25 | | 2466 ± 1204.5 | 124 | 24.9 | 2.58 | | | |
| All | 16 | 4.00 | 7036 | | 366 | 51.6 | 4.25 | | | |

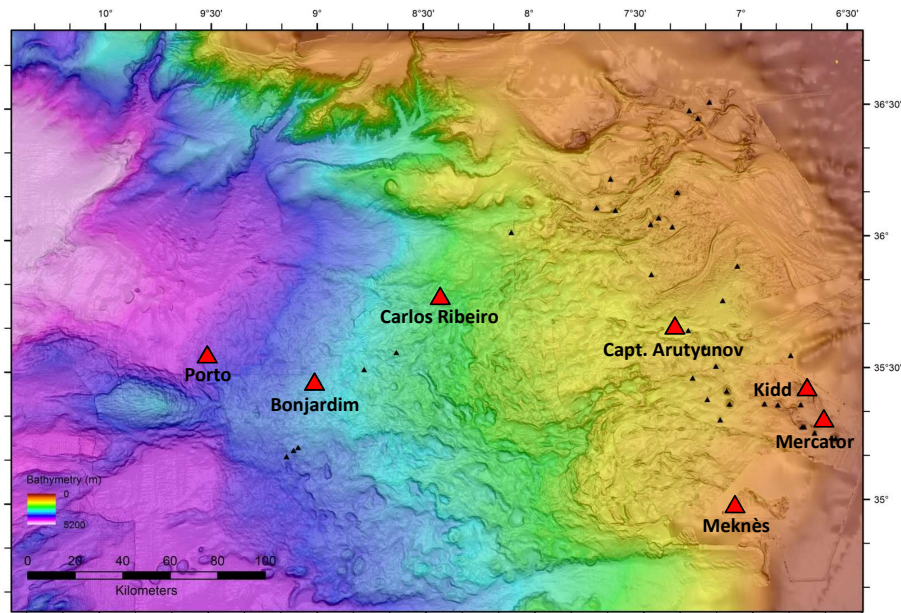


Fig. 1. Map of the study area in the Gulf of Cadiz. Large red triangles: location of the seven studied mud volcanoes; small black triangles: other mud volcanoes in the area; the two reference stations (not shown) are located in the vicinity of Mercator and Kidd mud volcanoes. Bathymetry from Zitellini et al. (2009).

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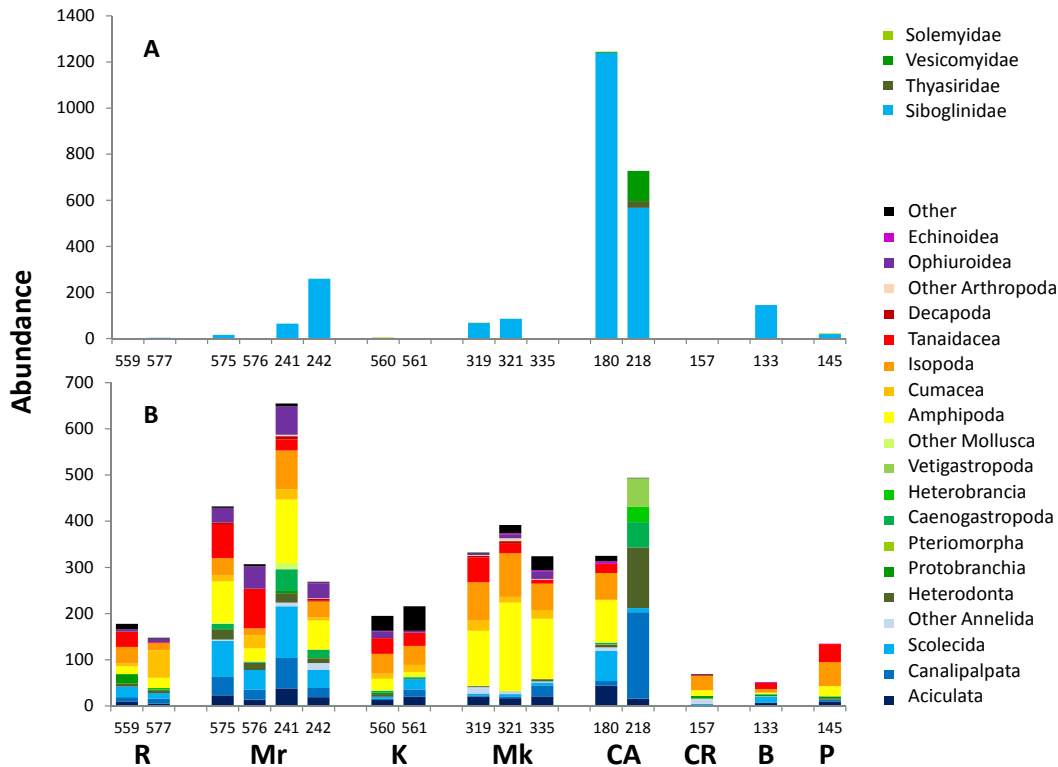


Fig. 2. Abundance per boxcore sample (0.25 m²). **(A)** Chemosymbiotic species (mixotrophic excluded); **(B)** Heterotrophic species. R: Reference sites; Mr: Mercator MV; K: Kidd MV; Mk: Menès MV; CA: Captain Arutyunov MV; CR: Carlos Ribeiro MV; B: Bonjardim MV; P: Porto MV. Numbers are the three last digits of the station code (Table 1).

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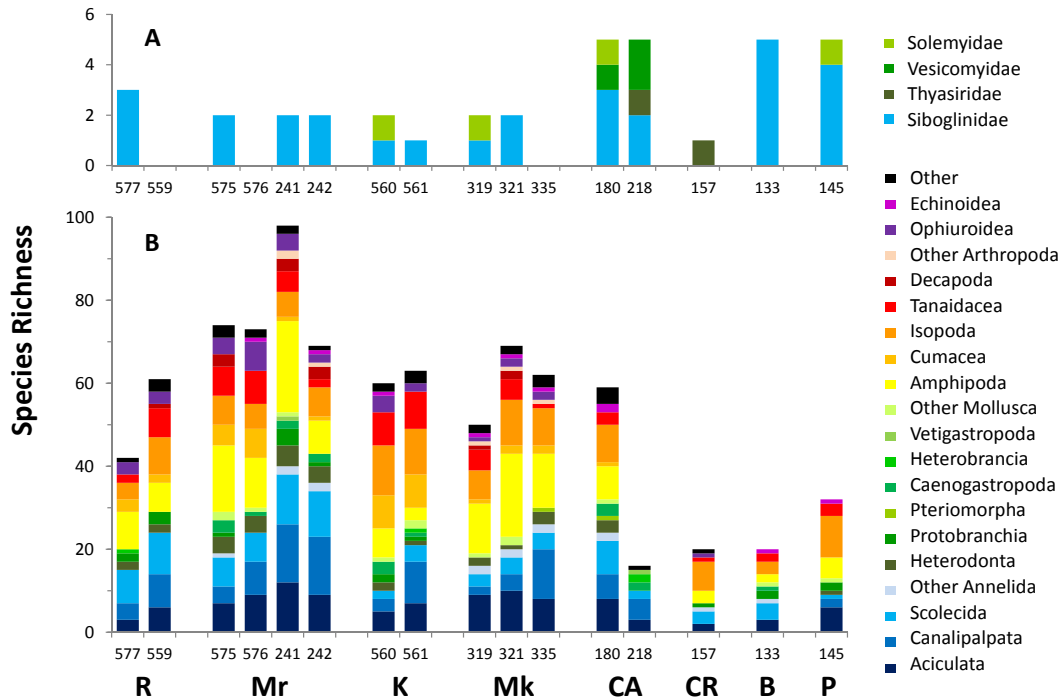


Fig. 3. Species richness per boxcore sample (0.25 m²). **(A)** Chemosymbiotic species (mixotrophic excluded); **(B)** Heterotrophic species. R: Reference sites; Mr: Mercator MV; K: Kidd MV; Mk: Menès MV; CA: Captain Arutyunov MV; CR: Carlos Ribeiro MV; B: Bonjardim MV; P: Porto MV. Numbers are the three last digits of the station code (Table 1).

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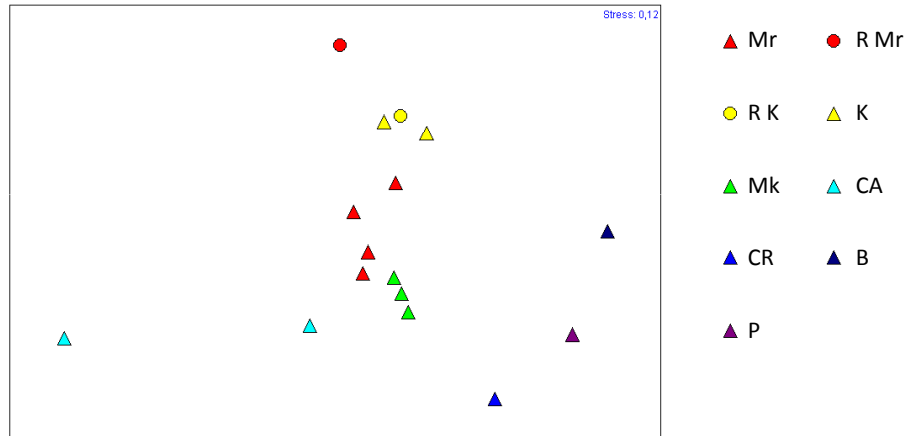


Fig. 4. MDS plots for the comparison between samples from Reference sites (Mercator R and Kidd R), Shallow mud volcanoes (Mercator, Kidd and Meknès) and Deep mud volcanoes (Capt. Arutyunov; Carlos Ribeiro, Bonjardim and Porto).

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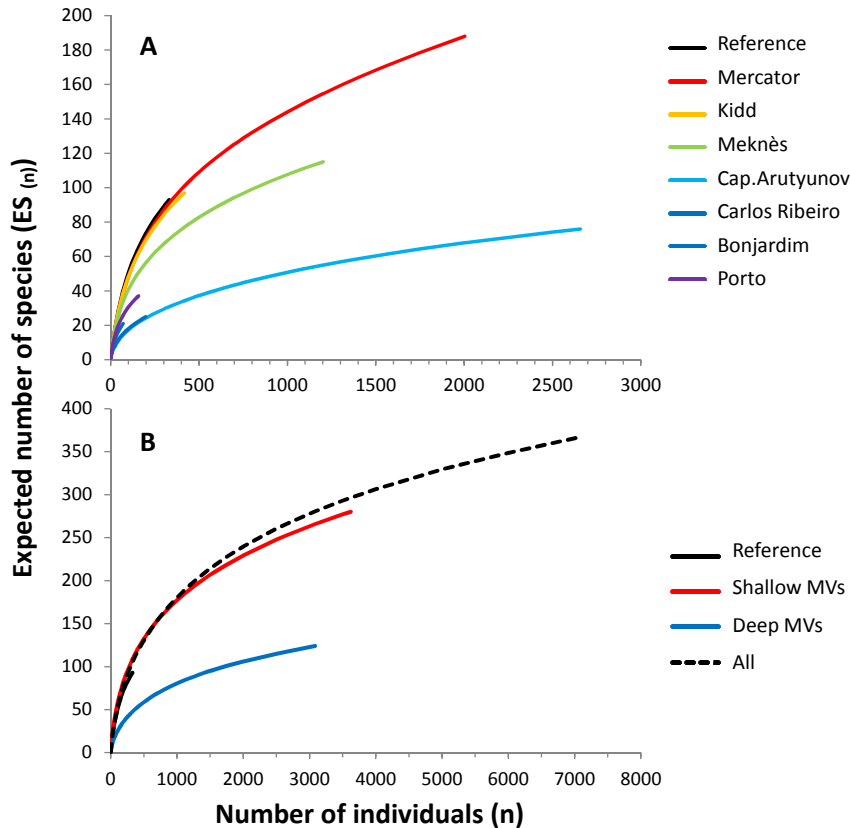


Fig. 5. Comparison of rarefaction curves (Hurlbert's expected number of species) for the assemblages pooled by mud volcano (top) and pooled by sub-region (bottom).

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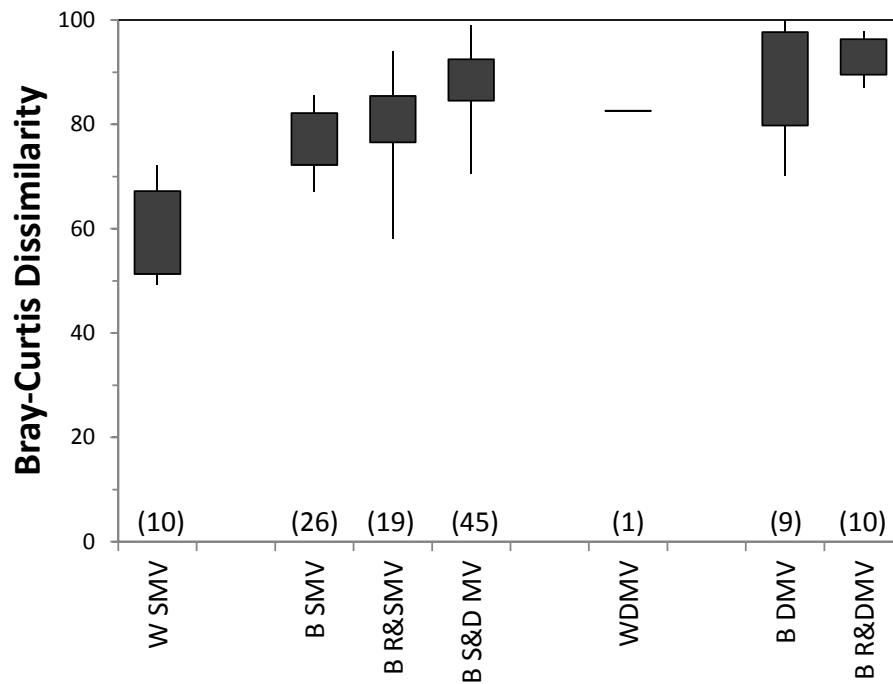


Fig. 6. β -diversity analysis based on Bray-Curtis dissimilarity between all possible pairs of individual samples. Bars show the range (line) and 25–75 percentile (box) of pairwise comparisons in each category (number of pairwise comparisons in brackets). W SMV: within Shallow MVs; B SMV: between Shallow MVs; B R&SMV: between Reference sites and Shallow MVs; B S&DMV: between Shallow and Deep MVs; W DMV: within Deep MVs; B DMV: between Deep MVs; B R&DMV: between Reference sites and Deep MVs.

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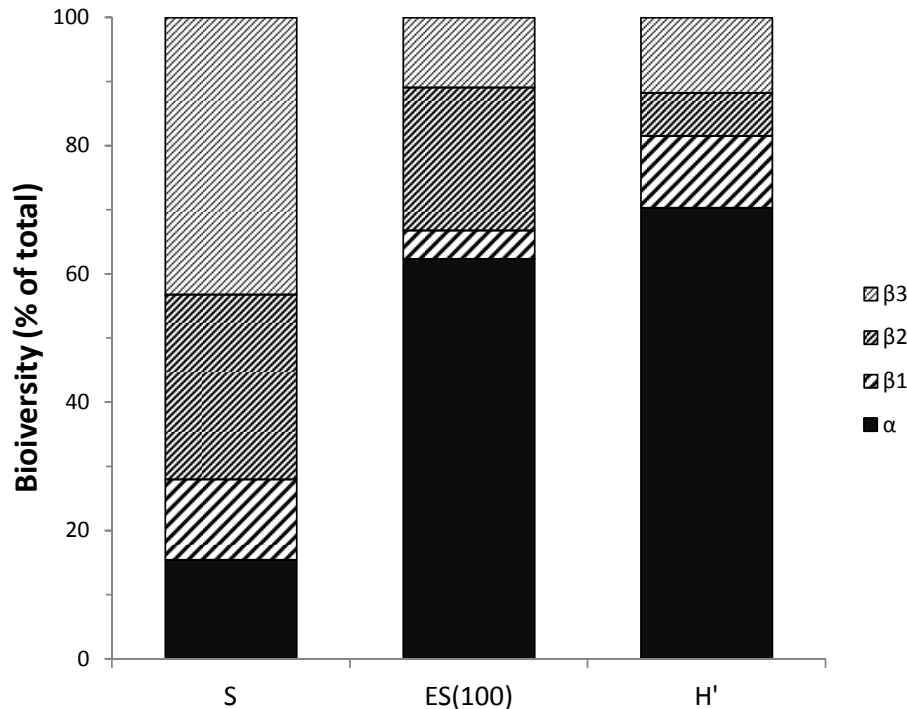


Fig. 7. Partition of diversity for different indices. S: number of species; ES(100): Hurlbert expected number of species per 100 individuals; H' : Shannon-Wiener diversity (ln-based). β_1 : β -diversity within mud volcanoes; β_2 : β -diversity between mud volcanoes β_3 : β -diversity between sub-regions.

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