**ECOLOGY**

**CO₂ leakage alters biogeochemical and ecological functions of submarine sands**

Massimiliano Molari,1*, Katja Guillini,2* Christian Lott,3 Miriam Weber,1,2 Dirk de Beer,4 Stefanie Meyer,1* Alban Ramette,16 Gunter Wegener,1,5 Frank Wenzhöfer,1,6 Daniel Martin,7 Tamara Cibic,8 Cinzia De Vittor,8 Ann Vanreusel,2 Antje Boetius1,5,6

Subseabed CO₂ storage is considered a future climate change mitigation technology. We investigated the ecological consequences of CO₂ leakage for a marine benthic ecosystem. For the first time with a multidisciplinary integrated study, we tested hypotheses derived from a meta-analysis of previous experimental and in situ high-CO₂ impact studies. For this, we compared ecological functions of naturally CO₂-vented seafloor off the Mediterranean island Panarea (Tyrrhenian Sea, Italy) to those of nonvented sands, with a focus on biogeochemical processes and microbial and faunal community composition. High CO₂ fluxes (up to 4 to 7 mol CO₂ m⁻² hour⁻¹) dissolved all sedimentary carbonate, and comigration of silicate and iron led to local increases of microphytobenthos productivity (+450%) and standing stocks (+300%). Despite the higher food availability, faunal biomass (−80%) and trophic diversity were substantially lower compared to those at the reference site. Bacterial communities were also structurally and functionally affected, most notably in the composition of heterotrophs and microbial sulfate reduction rates (−90%). The observed ecological effects of CO₂ leakage on submarine sands were reproduced with medium-term transplant experiments. This study assesses indicators of environmental impact by CO₂ leakage and finds that community compositions and important ecological functions are permanently altered under high CO₂.

**INTRODUCTION**

The atmosphere takes up large amounts of CO₂ from anthropogenic sources, resulting in global warming and increasing dissolution of CO₂ into seawater, with detrimental consequences for the ocean ecosystems (1). Ocean acidification is predicted to decrease seawater pH by 0.2 to 0.4 units by 2100 at unchanged rates of CO₂ emissions (2). To meet the international goal of limiting global warming to 1.5°C, the use of fossil fuels would have to end before 2040 (3) and may need to be complemented by mitigation technologies. One way to reduce industrial emissions is CO₂ capture and storage (CCS) in the subsurface, which includes subseabed reservoirs (4). This new maritime mitigation technology causes a need for assessments of ecological risk, especially from potential CO₂ leakage (5, 6). Besides reducing effectiveness of the technology, CO₂ leakage from subseaflow reservoirs could lead to extreme pore- and seawater acidification, with pH substantially lower than 7 (7), and thereby negatively affecting the local ecosystem. Current knowledge of high-CO₂ effects on marine ecosystems is mostly based on assessing the vulnerability of individual specimens and mesocosm communities to artificially enhanced CO₂ levels in seawater (8, 9). However, knowledge on long-term ecosystem-level responses and assessment of adaptation and resilience of communities are limited (10, 11). Thus, a crucial question remains whether CO₂ leaks can locally lead to profound and persistent changes of element cycling, as well as to negative effects on ecosystem functions and services, including biodiversity and productivity. This question calls for field studies of naturally complex, dynamic ecosystems under long-term high-CO₂ exposure (for example, caused by volcanic degassing) (12, 13). To our knowledge, this study is the first synchro-nous assessment (that is, occurring at the same time and place) of high-CO₂ effects covering all trophic levels from microbes to macrofauna in submarine sands. Sands make up a substantial proportion of shelf seas and play a critical role as biogeochemical filters at the land-sea boundary (14). We investigated for over 2 years the impact of CO₂ degassing on benthic biogeochemistry and community structure from microbes to macrofauna, focusing on carbon cycling (primary productivity and organic matter remineralization). In addition, we transplanted sediments between CO₂-vented and nonvented sites to assess the immediate effects of changing CO₂ levels within a year and to test whether we could reproduce the natural patterns. On the basis of a meta-analysis of previous high-CO₂ impact studies, we derived and tested the following hypothesis: CO₂ leakage enhances benthic primary production but negatively affects ecosystem functional diversity, with consequences for the benthic food web and carbon fluxes.

**RESULTS**

Identification of natural analog sites for the leakage scenario

The Aeolian archipelago in the southern Tyrrhenian Sea is a ring-shaped volcanic arc (fig. S1A), composed of 7 islands and 10 seamounts, associated with the Peloritan-Calabrian orogenic belt (15). Panarea, the smallest (3.3 km²) Aeolian island, represents the emergent part of a wide stratovolcano more than 2000 m high and 20 km across (16). In 2011, we surveyed a number of CO₂-vented sites around Panarea to identify those to be used as “natural laboratories” to assess pure CO₂ effects, finally selecting the eastern side of Basiluzzo Islet (a rhylolitic dome northeast to Panarea). Two sites (fig. S1B) best fulfilled the “natural laboratory”
criteria: (i) continuous, dispersed degassing of CO2 through sand causing low pH; (ii) similar oxygen availability and negligible co-emission of toxic substances or microbial energy sources such as sulfide and methane; and (iii) no significant temperature anomalies from hydrothermalism. The selected “CO2-R” and “CO2-G” sites showed comparable environmental conditions to the reference (Table 1), and rather evenly distributed gas leakage (fig. S1D; density of two to three gas bubble strings per m²). The reference site (REF) showed no gas emissions (fig. S1C).

| Table 1. Main environmental characteristics of sampling sites at Basiluzzo Islet (Panarea Island, Italy). | REF | CO2-G | CO2-R |
|---|---|---|---|
| Coordinates | N 38°39.827′ E 15°07.118′ | N 38°39.820′ E 15°07.137′ | N 38°39.749′ E 15°07.123′ |
| Water depth | m | 14–17 | 21 | 15–17 |
| Area | m² | 100 | 35 | 200 |
| Seagrass meadows | | Posidonia oceanica | Posidonia oceanica | Posidonia oceanica |
| Bottom water properties (10 cm asf) | | | | |
| Temperature | °C | 18.8–19.5 | 18.8–19.0 | 18.7–19.3 |
| Salinity | ‰ | 38 | 38 | 38 |
| ORP | mV | 245 (±75) | na | 133 (±64) |
| pH | | 7.8 | 7.3 |
| DIC | mmol liter⁻¹ | 2.1 (±0.1) | 2.3 (±0.1) | 25 (±0.2) |
| TA | meq kg⁻¹ | 2.3 (±0.1) | 2.3 (±0.1) | 2.4 (±0.2) |
| CaCO₃ | meq kg⁻¹ | 0.0 (±0.02) | 0.0 (±0.02) | 0.0 (±0.02) |
| Si(OH)₄ | mmol liter⁻¹ | 2.1 (±0.0) | 2.7 (±0.0) | 3.2 (±0.0) |
| NH₄ | µmol liter⁻¹ | 0.4 (±0.0) | 0.8 (±0.0) | 0.5 (±0.0) |
| NO₂⁻/NO₃⁻ | µmol liter⁻¹ | 0.1 (±0.0) | 0.2 (±0.0) | 0.7 (±0.0) |
| Fe | µmol liter⁻¹ | 0 | 0.5 | 0.4 (±0.0) |
| Mn | µmol liter⁻¹ | 0 | 0 | 0.5 |
| Sediment properties (0–10 cm layer) | | | | |
| Color | | Gray | Gray | Red (rusty) |
| Median grain size | | Coarse sand | Coarse sand | Coarse sand |
| Porosity | % | 30–44 | 38–43 | 38–43 |
| Carbonate content | mg g⁻¹ | 9.34 (±1.13) | 0.04 (±0.02) | 0.08 (±0.02) |
| Porewater fluxes | | | | |
| Gas bubbling | | Yes | Yes | Yes |
| CO₂ content | % | — | 90–97 | 97–99 |
| Gas flow | Liter m⁻² hour⁻¹ | — | 80 | 120 |
| Porewater flow | Liter m⁻² day⁻¹ | 11–69 | 12–45 | 11–85 |
| DIC flux | mol m⁻² day⁻¹ | 0.0–0.2 | 2.4–13.8 | 2.7–10.3 |
| Si(OH)₄ flux | mmol m⁻² day⁻¹ | 0.0–0.9 | 10.0–41.7 | 17.6–28.2 |

†Patch of bare sediment within seagrass bushes. ‡Average temperatures in 2011 to 2013 measured in situ with SEAGUARD at 30 cm asf. §Average (±SD; n = 4000) of 2012 data collected in situ with RBR sensors over 15 days at 2 cm asf. ¶Average of 2011 to 2013 measurements (n = 9). ‖Average of 2011 to 2013 measurements (n = 9). ‡‡Average of 2011 to 2013 measurements (n = 9). ‡‡‡Average of 2011 to 2013 measurements (n = 9). ‡‡‡‡Average of 2011 to 2013 measurements (n = 9). ‡‡‡‡‡Average of 2011 to 2013 measurements (n = 9). ‡‡‡‡‡‡Average of 2011 to 2013 measurements (n = 9). *P < 0.05. **P < 0.001; Welch's t test between REF and CO2-R.
Gas and bottom water chemistry

The gas bubbles emanating at CO2-R and CO2-G consisted mainly of CO2, with traces of CO and CH4 (0.32 and 0.01 parts per million, respectively). Tidal variations in venting, with enhanced CO2 leakage during low tide, caused peaks in pHT (total scale; Fig. 1A). The CO2 emission rates during low tide were 6.6 mol CO2 m$^{-2}$ hour$^{-1}$ at CO2-R and 4.2 mol CO2 m$^{-2}$ hour$^{-1}$ at CO2-G. Bottom water oxidation-reduction potential (ORP) at CO2-R was significantly lower than at REF (Table 1), being driven by effluxes of Fe$^{2+}$-enriched porewater at the vented sites (see section below). Bottom water O2 concentration varied slightly with the light
period due to photosynthesis, with the average being higher at CO2-R than at REF (Table 1).

The bottom water pH_T measured at 5 to 10 cm above seafloor (asf) was, on average, 7.9 at REF and 6.7 to 7.7 at CO2-R (Fig. 2B), compared to 7.5 to 7.9 at CO2-G (Table S1). Saturation states of calcite (Ω_{calc}) and aragonite (Ω_{ara}) were lower at the vented sites than at REF, but always >1 (Fig. 1C and Table S1). Only marble tiles exposed to the vented seafloor partially dissolved within 1 year (n = 6; dissolution rates, 0.02 to 9.94 mg day \(^{-1}\)), whereas those with a distance >85 m away from the venting area center (CO2-R) did not show dissolution (n = 42). Nutrient levels (phosphate, ammonium, nitrite, and nitrate) in the bottom water did not differ significantly between the CO2-impacted sites and REF (Table 1). The silicate content was also similar, with 1.2 to 1.8 µM in 2012 and 1.7 to 3.5 µM in 2013. However, iron and manganese bottom water concentrations increased at the CO2-vented sites and were highest at CO2-R (Table 1).

**Porewater chemistry**

At REF, the pH_T decreased slightly with increasing sediment depth at REF, from 7.9 (sediment surface) to 7.7 [2.0 cm below seafloor (bsf)] and then remained constant (Fig. 1B and table S1). In contrast, at CO2-G and CO2-R, a pH_T of ca. 5.5 was reached already at 2.5 and 0.5 cm bsf, respectively (Fig. 1B and table S1). CO2(aq) increased rapidly with sediment depth, from 0.02 mM (sediment surface) to 5.3 mM (2 cm bsf) at both vented sites. O2 penetrated to 2 cm bsf at REF and CO2-G sediments and to ca. 1 cm bsf at CO2-R (fig. S2A). ORP was constant at REF and decreased at CO2-G and CO2-R (to ~50 mV within 1 cm bsf; fig S2A). No sulfide was detected in the subsurface porewaters, but peaks of a few micromolar were measured directly at the sediment surface at REF and CO2-G and to a lesser extent at ca. 0.5 cm bsf at CO2-R (fig. S2A). Hydrogen concentrations were below 1 µM (detection limit, 0.3 µM) and constant down to 5 cm bsf at all sites (fig. S2A). Together, the chemical gradients indicated that CO2-R was more strongly vented than CO2-G.

In REF sediments, porewater total alkali (TA) was constant but increased substantially with depth at the CO2-vented sites (Fig. 1C). Ω_{calc} and Ω_{ara} decreased to <1 at the vented sites, whereas they remained around 2 and 1 at REF, respectively (Fig. 1C and table S1). Porewater at the vented sites was significantly enriched in silicate, iron, manganese, and, somewhat, phosphate (fig. S2B). Fe\(^{2+}\) was almost absent from REF porewaters, whereas it reached 0.5 to 1 mM at the vented sites, explaining the ORP dynamics in porewater and bottom waters and also the enhanced bottom water concentrations. In contrast, B, Ca, Na, Mg, Sr, Li, and K concentrations were similar at REF and CO2-vented sites.

**Sediment grain size, carbonate, and elemental composition**

All three sites were dominated by coarse sand with similar porosity and grain size distribution (Table 1). Concurrent with the observed undersaturation in calcite and aragonite in porewaters of the vented sites, the solid-phase carbonate content was about 100 to 200 times lower compared to those in REF sediments (Table 1 and Fig. 2). In accordance with the high porewater Fe concentration, also solid Fe was elevated at the vented sites (3.4 mg g \(^{-1}\) at CO2-R versus 0.4 mg g \(^{-1}\) at REF). Total organic carbon (TOC) was low (<0.1%) but approximately twofold higher in the surface sediments of CO2-R and CO2-G compared to REF (Fig. 2). Total organic nitrogen (TN) was also very low (<0.2 µg mg \(^{-1}\)) at all three sites, leading to a C/N ratio of ca. 4 to 7.5 in the surface sediments. Both TOC and TN were higher in 2013 than in 2012 (table S2A).

**Fluxes and remineralization rates**

Benthic chambers placed in between the bubble streams at the CO2-vented sites showed a decreasing pH in the enclosed water bodies with time, together with a substantial efflux of dissolved organic carbon (DIC) and silicate from the sediment (table S3). In comparison, at REF, no effluxes of silicate or DIC were detected, and the chamber water pH_T remained stable at 8.0 in incubations of up to 5 hours.

Benthic chamber measurements at the vented sites showed similar advective fluid flow rates as those at REF (Table 1). On the basis of ORP signals, porewater iron concentrations, and fluid flow rates, it is likely that a substantial iron efflux occurred at the vented sites. Thus, respiration rates were corrected for potential oxygen consumption by purely chemical Fe\(^{2+}\) oxidation, which amounted to 1 to 7% of the total oxygen consumption at the vented sites. At the time of chamber deployments, the seafloor at all sites showed net oxygen consumption, and respiration always exceeded photosynthetic O2 production even during daytime. However, both oxygen respiration and production were substantially higher at the vented sites compared to REF (Table 2). Diffusive oxygen fluxes calculated from microprofiler measurements (fig. S2) were <10% of the total fluxes but were also higher at CO2-R (2.6 mmol m \(^{-2}\) day \(^{-1}\)) than at REF (1.7 mmol m \(^{-2}\) day \(^{-1}\)).
Standard proxies for microbial activities were also influenced by high CO2. The β-glucosidase hydrolytic activity measured at substrate saturation ($V_{\text{max}}$) was significantly higher at CO2-G and CO2-R than at REF, whereas the aminopeptidase and esterase activities were significantly lower at the vented sites (Fig. 2). Its contribution to aerobic benthic respiration was 16% at REF and 0.04% at the vent sites. SRRs determined in vitro were substantially higher at REF than at both vent sites (Fig. 2). Its contribution to aerobic benthic respiration was 16% at REF and 0.04% at the vent sites.

**Microbial community patterns**

Benthic diatoms (Bacillariophyceae) dominated the microphytobenthos in the surface sediment layer (95 ± 4% of cells), and their abundances were up to three times higher at CO2-R than at CO2-G and REF. Abundances were ca. 50% lower in 2012 than in 2013 at all sites: 4709 ± 636 cells cm$^{-2}$ and 8932 ± 560 cells cm$^{-2}$ at CO2-R, 1552 ± 158 cells cm$^{-2}$ and 6079 ± 973 cells cm$^{-2}$ at CO2-G, and 1744 ± 150 cells cm$^{-2}$ and 4909 ± 218 cells cm$^{-2}$ at REF, respectively (table S2A). CO2-R also showed the highest content of chlorophyll a (Chl a) pigments (Fig. 2). Chl a made up 73 to 95% of the chloroplastic pigment equivalents (CPEs), indicating that the pigments originated mostly from living cells. At all sites, Chl a decreased with sediment depth, with 33 to 50% concentrated in the top 2 cm, and was positively correlated with TOC (Pearson’s $R = 0.4967$; $P < 0.001$; $n = 63$).

In contrast, total bacterial cell abundances were similar at all sites (Fig. 3A). The highest abundances occurred in the upper sediment layer (ca. 0.7 × 10$^9$ cells m$^{-2}$) and decreased with sediment depth. Bacteria dominated [54 to 71% of 4,6-diamidino-2-phenylindole (DAPI)-stained cells] over Archaea (3 to 1% of DAPI-stained cells).

With 454 massively parallel tag sequencing (MPTS), we recovered a total of 9674 bacterial operational taxonomic units (OTUs; 0.03) from all sediment layers combined. At the class level, bacterial communities of the vented sites were dominated by Flavobacteria, Gammaproteobacteria, Deltaproteobacteria, Caldilinea, and unclassified Cyanobacteria (Fig. 3A). Functional group analysis (table S4) showed that CO2 leakage stimulated primary producers (that is, Cyanobacteria and Chlorobia), aerobic and anaerobic organic matter–degrading bacteria (that is, Flavobacteria and Caldilinea), some metal-reducing bacteria (that is, Desulfuromonadales), and some ferrotrophic bacteria (that is, Rhodobacteraceae). Concurrent with the negative impact of high CO2 on SRR, the relative sequence abundances of sulfate reducers (that is, Desulfoacteriales) were reduced, as well as those of sulfur oxidizers (that is, Candidatus Thiomicos) and nitrifiers (that is, Nitrosospira, Nitrosoospira, and Nitrosococcus).

**pH and DIC were the main environmental parameters influencing the bacterial community structure, explaining more than 35% of the variance in OTU composition for all data sets (that is, amplified ribosomal intergenic spacer analysis (ARISA) and MPTS; table S5A). The three sites differed in bacterial community composition by 58 to 74% (that is, ARISA; fig. S4A). The principal source of variability was associated with the differences in CO2 flux and associated parameters (table S6).**

**Benthic invertebrate communities**

Total meiofauna density was much higher at REF than at the vented sites [1019 ± 354 individuals (Ind.) m$^{-2}$ versus 407 ± 237 Ind. m$^{-2}$], which was mirrored in all taxa: nematodes (Fig. 3B) and copepods, which dominated, but also nauplii, polychaetes, and tardigrades (table S7). Total meiofauna and nematode abundances decreased rapidly along the sediment profile (ANOVA; $P < 0.001$; $F_{3,71} = 19.0$ and $F_{3,71} = 11.6$, respectively), more steeply at the CO2-vented sites than at REF. At REF, this gradient was not reflected in nematode biomass, conversely at CO2-vented sites, the biomass was significantly higher in the top layer (0 to 2 cm) than in the other layers (ANOVA; $P < 0.001$; $F_{3,20} = 6.6$ for CO2-G and $F_{3,20} = 12.7$ for CO2-R). Community structure for meiofauna at the higher taxon level differed significantly between the three sites (tables S6 and S7) but was even more different at the nematode species level (table S7 and Fig. 4B). Year-to-year differences in the nematode.

### Table 2. Benthic oxygen fluxes.

Oxygen exchange (transparent chamber; net O$_2$ flux), oxygen consumption (masked chamber; O$_2$ respiration), and oxygen production [GPP = net O$_2$ flux + (O$_2$ respiration)] rates obtained from benthic chambers deployed in 2013; oxygen production–to–respiration (GPP/R) ratio, respiration per unit of total biomass (R/BTotal), and respiration per unit of heterotroph biomass (bacteria and animals; R/BHeterotrophs). Benthic masked chambers (n = 2) and average (mean with ±SD in parenthesis), maximum (Max), and minimum (Min) rates of transparent chambers (n = 3 to 4) and O$_2$ production (n = 6 to 8) are given. nt, not tested for significance level; ns, not significant (P > 0.05).

| Net O$_2$ flux (daylight)$^\text{mM}$ | O$_2$ respiration (masked)$^\text{mm}$ | O$_2$ production$^{**}$ | GPP/R | R/BTotal | R/BHeterotrophs |
|-------------------------------------|-------------------------------------|------------------------|-------|----------|-----------------|
| REF                                 | Mean                                | –7 (6)                 | 10 (8)$^\dagger$ | 0.6    | 0.06/0.14       | 0.07/0.18       |
|                                     | Max                                 | –11                    | –23   | 21       |                 |                 |
|                                     | Min                                 | –2                     | –10   | 0        |                 |                 |
| CO2-G                               | Mean                                | –58 (63)               | –188$^\ddagger$ | 130 (63)| 0.7    | 1.13            | 2.06            |
|                                     | Max                                 | –151                   | –173  |          |                 |                 |
|                                     | Min                                 | –15                    | na    | 37       |                 |                 |
| CO2-R                               | Mean                                | –18 (6)                | –55 (37)$^\ddagger$ | 0.8    | 0.16/0.49      | 0.41/1.14       |
|                                     | Max                                 | –24                    | –106  | 96       |                 |                 |
|                                     | Min                                 | –11                    | –38   | 15       |                 |                 |

$^\dagger$Average (±SD; n = 6) of O$_2$ production calculated from each Net O$_2$ flux using O$_2$ respiration from both masked chambers. $^\ddagger$At CO2-R, only one masked chamber was available. $^\ddagger$Average (±SD; n = 8) of O$_2$ production calculated from each O$_2$ flux using O$_2$ respiration from both masked chambers. **P < 0.01 (Welch’s t test between REF and CO2-R).
Fig. 3. Community composition of studied sampling site (top 5 cm of sediments). (A) Microbial cell numbers and bacterial community structure, as described by 454 MPTS, showing relative number of sequences for dominant bacterial classes (that is, OTUs > 0.1%) clustered according to similarity [based on the Bray-Curtis distance matrix, surface, and subsurface layer; analysis of similarities (ANOSIM); \( R = 0.948; P < 0.001 \)]. (B) Nematode density and biomass and relative abundance of nematode feeding groups. (C) Polychaete density, macrofauna biomass, and relative abundance of polychaete feeding groups. Error bars are ±SD; year and number of sampling are given in each plot; stars indicate significant differences between one or both CO₂-vented sites and the REF (ANOVA; \(^* P < 0.05, ^{**} P < 0.01, ^{***} P < 0.001\); for details, see table S2E). Ind., individuals; dwt, dry weight; uncl., unclassified.
plants (ANOSIM; R = 0.982; P < 0.001). The bacterial community of CO2-R transplanted into REF sediment (CO2-R/REF) was also significantly different to that of the original site (ANOSIM; R = 0.961; P < 0.001). Both communities that were transplanted within their own habitat remained similar significantly different to the undisturbed ones. (P < 0.001). The bacterial community of REF sediment (top, 10-cm layer) transplanted into CO2-R sediment (REF/CO2-R) was significantly different from the source community after 1 year (ANOSIM; R = 0.982; P < 0.001). The bacterial community of CO2-R transplanted into REF sediment (CO2-R/REF) was also significantly different to that of the original site (ANOSIM; R = 0.961; P < 0.001). Both communities that were transplanted within their own habitat remained similar to the undisturbed ones. (P < 0.001).

**Bacterial community of REF sediment (top, 10-cm layer) transplanted into CO2-R sediment (REF/CO2-R) was significantly different from the source community after 1 year (ANOSIM; R = 0.982; P < 0.001).**

**DISCUSSION**

Here, we focused on permeable sandy marine ecosystems that occupy large areas of the continental shelves, the target areas for submarine CCS (7). To assess potential ecological risks from CO2 leakage (5), we synchronously investigated the geochemical phenomena of CO2 leakage and its effects on community function and composition including different benthic size classes and trophic groups from microbes to macrofauna. Moreover, our multidisciplinary integrated approach allowed us to test the main hypothesis derived from a meta-analysis of previous experimental and in situ studies (Table 3): CO2 leakage locally enhances primary production in sandy sediments, but it negatively affects ecosystem functional diversity, with consequences for the benthic food web and carbon fluxes.

**Geochemical phenomena of CO2 venting**

Here, we compared the geochemical characteristics of a nonvented REF, representing the natural baseline, with two different CO2-vented sites, CO2-R and CO2-G, of similar hydrological and sedimentological characteristics (Table 1 and fig. S1). CO2 leakage was identified visually as decreased at the vent sites, which were instead dominated by predators and scavengers (2B; CO2-G) or epistrateum feeders (2A; CO2-R; Fig. 3B). This finding is in line with the highest diatom densities at CO2-R. pH was the most influencing environmental parameter for the nematode assemblage structure over the whole sediment profile, explaining more than 40% of the variance (table S5A).

Macrofauna was dominated by polychaetes at all sites, with relative abundances of 71 ± 8% (REF), 69 ± 45% (CO2-G), and 45 ± 36% (CO2-R). Polychaete abundances, as well as the whole macrobenthos biomass, were substantially lower at the vent sites compared to REF (Fig. 3C). The polychaete community structure also differed significantly between REF and the two vent sites (Tables S6 and S7 and fig. S4C). At the vent sites, all polychaetes were grazers or deposit feeders, whereas at REF, filter feeders, carnivores, and omnivores also occurred (Fig. 3C).

**Sediment transplantation**

Sediment was transplanted between REF and CO2-R, and after 1 year, sediment parameters, microbial activities, nematode density, and bacterial and nematode community structure were compared. In REF/CO2-R, porewater DIC, TA, silicate, and iron increased, and pH decreased, whereas the opposite trend occurred in CO2-R/REF (fig. S3 and table S8). Both pH and carbonate content of CO2-R/REF remained significantly lower than those of REF after 1 year (table S8 and Table 1). Still, the carbonate content increased 35-fold in the 0- to 2-cm sediment layer and doubled in the 4- to 6-cm sediment layer. In contrast, carbonates dissolved in REF/CO2-R both in the 0- to 2-cm sediment layer and in the 4- to 6-cm sediment layer (table S8).

As to microbial activities, the β-glucosidase activity increased in REF/CO2-R, whereas aminopeptidase and esterase activities decreased (fig. S3). SRR also decreased in REF/CO2-R and did not recover in CO2-R/REF (fig. S3). Chl a increased in REF/CO2-R and decreased somewhat in CO2-R/REF (fig. S3).

It took a year until the cross-transplanted bacterial and nematode communities resembled the respective background communities (Fig. 4). However, the nematode density decreased significantly in REF/CO2-R but did not increase in CO2-R/REF (fig. S3). pH and DIC were the main environmental factors responsible for the observed shifts in bacterial and nematode community structure in the transplanted sediments (fig. SSB).
escaping gas bubbles of >90% CO₂ content (fig. S1D). Similar to this natural analog, CO₂ upward migration through subseafloor sediment strata in the case of CCS leakage would result in the dissolution of the gas, leading to subsequent reactions with porewater and sediments, so that only a fraction of the gas would escape to the water column (17).

As a consequence of CO₂ dissolution, porewater pH and carbonate saturation would decrease, whereas DIC and TA will increase (18). We recorded all of these geochemical phenomena at the Basiluzzo vent sites: CO₂ venting through the sandy sediments resulted in a loss of solid-phase carbonate and a decrease in porewater pH (Table 1, Fig. 1B, and table S1), as well as in emission of acidified porewaters to the water column (Fig. 1A and table S3). Hence, the long-term geochemical consequence of CO₂ leakage through marine sediments would be the local decline of buffering capacity and a reduction of the mineral carbon sink (Fig. 5).

The CO₂-enriched porewater fluids at Basiluzzo did not contain elevated sulfide or methane concentrations typically associated with hydrothermalism. Boron, found at high concentrations in Panarea hydrothermal fluids (19), showed typical seawater concentrations in the Basiluzzo porewaters. The temperature anomaly in the surface sediments at the vent sites was negligible. However, we recorded some ORP dynamics in the bottom water above the vents, as well as substantially elevated iron, manganese, and silicate concentrations in the porewaters, derived from subseafloor hydrothermal and/or CO₂ reactions with the bedrock and overlying sands. Together with the high CO₂ fluxes, these high iron and silicate concentrations apparently enhanced productivity of the microphytobenthos. In an analog CCS leakage scenario, the CO₂ co-leakage will depend on the type of geological reservoir, and these could include mineral products of weathering from CO₂ exposure as well as hydrocarbons (18).

**Effects of CO₂ venting on primary production and microphytobenthos**

The CO₂-vented sites had higher microphytobenthos standing stocks, higher Chl a content, and more TOC, mostly due to an enhanced productivity of benthic diatoms (Figs. 2 and 5), especially where CO₂ leakage was the highest (CO₂-R). This effect was reproduced by transplantation of reference sands to CO₂-impacted site (REF/CO₂-R), showing increased Chl a content after 1 year (fig. S3). Noncalcifying benthic primary producers are likely to profit from high CO₂ (Table 3) as a result of the reduction in the energy costs for carbon concentration mechanisms (20). In addition, the higher availability of nutrients (especially silicate and iron, but also phosphorus) in
the upward migrating fluids at the vent sites may stimulate microphytobenthos growth. Similar effects were recorded previously along a natural CO2 gradient at Volcano Island (Italy), where the microphytobenthos was found to be promoted by CO2 leakage, showing a twofold increase in Chl a concentrations and a two to four times higher diatom abundance under high CO2 (21). As to specific effects on benthic diatom genera, a previous study at Basiluzzo (22) suggested that the diatom genera *Fragilaria*, *Diploneis*, and *Amphora* are favored by CO2 leakage. *Fragilaria* was the most abundant diatom taxon at CO2-R and was observed to form colonies on the surface sands, likely as a response to the combination of CO2 and nutrient enrichment by porewater advection. This dominance of chain-forming diatoms at CO2 vent sites has been also reported in other coastal areas (21, 23). Furthermore, *Diploneis* seems to be an opportunistic genus that becomes more competitive in the presence of environmental stress (24). This genus, as well as members of *Amphora*, was represented by larger cells with heavily silicified frustules at CO2-R, compared to REF. This increase in microphytobenthos standing stock was also reflected in significantly higher oxygen production at CO2-R compared to REF (Table 2). The CO2 leakage caused a higher primary production to respiration ratio, keeping more carbon fixed in microphytobenthos biomass and in total organic carbon. Our results suggest that this effect is also due to reduced grazing pressure and altered microbial community function, as discussed below.

**Effects on faunal community biomass and composition**

Our study shows that CO2 leakage led to a significant decline in abundance and biomass and a change in community composition for meiofauna and macrofauna (Fig. 3, B and C). Being the most abundant taxa at all target sites, nematodes and polychaetes were particularly affected (fig. S4, B and C). Previous studies using benthic mesocosms and laboratory experiments found that acute CO2 leakage exposure changed macrofaunal or meiofaunal community abundance, biomass, and composition as a result of seawater acidification (duration of experiments, maximum of 20 weeks; pH levels, ≥ 5.6) (11, 25–28). Specific experiments focusing on nematode communities found no negative effects on abundance, composition, or diversity at pH ≥ 6 (28–30). For this highly abundant meiofaunal taxon, decreases in density (29, 31, 32) or an increasing mortality based on changed morphometrics (33) only occurred when seawater pH is <6. In our study of naturally CO2-vented sands, we detected substantial long-term effects on nematodes already below a porewater pH of 7. This effect was reproduced by the experimental transplantations, which lead to a significant decrease of nematode density and a shift in community structure as a direct consequence of CO2 leakage (fig. S3 and Fig. 3B). The nematode community did not fully recover to background density within 1 year. Furthermore, for several taxa of meiofauna and macrofauna, we show that these CO2 effects persist under long-time exposure and are not overcome by adaptation and community change. Our results confirm
previous findings on epibenthic macrofauna communities from other CO2 vents (Table 3) and indicate that few invertebrate taxa can cope with high CO2. Particularly, opportunistic species with short life spans and capable of rapid colonization in strongly disturbed habitats seemed to tolerate the extreme and chronic high partial pressure of CO2/low-pH conditions. These include polychaete species of the Capitella clade, some sponiosids, the interstitial hezonoid Microphthalminus tyrrhenicus, the paraonid Aricidea curtulii, and the nematode species Microlaimus comprius, Microlaimus honestus, Oncholaimus campyloriboides, and Daptonema micropiculum. The polychaete and nematode communities also showed a strong trophic shift, being more diversified under the baseline pH conditions (Fig. 3, B and C). Hence, our observations confirm that despite the additional energy availability due to the high microphytobenthic production at the CO2-vent ed sites, the associated benthic communities are negatively affected by high CO2, with declining densities and loss of functional diversity as main consequences.

Effects of elevated CO2 levels on microbial communities

In contrast to the faunal communities, we did not detect a significant change of bacterial or archaeal densities in CO2-vented sands compared to REF (Fig. 3A). Some bacterial taxa, like Oceanospirillacea, did not show differences in relative sequence abundance between vent and REFs and, hence, seem not to be affected by CO2 leakage (Table S4). However, we detected an overall substantial shift in community composition already at the phylum and class levels, which was increasingly pronounced at increasing taxonomic resolution (Fig. S4A). Previous studies using natural leakage analogs also found impacts on both microbial compositions already at pH < 7.7 (Table 3).

Here, the most striking change in community composition was the decline of relative abundances of Gammaproteobacteria by 50%. Members of this group typically dominate marine sediments but seem unable to cope well with high CO2, as previously reported from pelagic mesocosms (34–37), sedimentary CO2 vents (38, 39), and sponge and coral associates (40). In contrast, the Flavobacteriaceae, a bacterial family relevant in the degradation of marine algal organic matter (41), are not negatively affected by high-CO2 exposure and low pH according to our study, as well as to laboratory experiments (34, 42) and previous observations on natural CO2-vented sites (40, 43). As for the anaerobic bacteria inhabiting the deeper, more acidified sediments, the Caldilineales, a group involved in organic matter degradation under anaerobic and low-pH conditions (44, 45), were favored at the vented sites. Conversely, the relative sequence numbers of anaerobic bacteria, including sulfate reducers (order Desulfbacterales) and nitri fiers (genus Nitrospira, Nitrospina, and Nitrosococcus), decreased significantly at the CO2 vents. Despite the high Fe2+ availability in porewaters, no mats of iron-oxidizing bacteria were observed, and typical iron oxidizers were missing in the acidified sands. In the transplantations, the significant shift in bacterial community occurring after 1 year (Fig. 4A) was in line with the long-term changes but was not detectable in short-term incubations (that is, 2 weeks; data not shown). Transplantation from the vent site to the REF showed incomplete recovery within a year.

Consequences of CO2 leakage on local food webs and carbon fluxes

Acidification influences all cellular processes, including enzyme kinetics and membrane potentials, but different species are differently adapted to high CO2 levels. Our study shows significant, long-lasting effects of high CO2 on benthic biomass and composition, which alter biogeochemical functions at the ecosystem level (Fig. 5). Integrated with findings from previous experiments and field studies, we prove that these are consistent indicators of high-CO2 effects across different ecosystem types, organism size classes, and ecological functions.

Our study detected a substantial increase in microphytobenthos primary production and standing stock in relation to CO2 seepage and the comigration of nutrients such as silicate and iron. We expected this to compensate the metabolic costs of adaptation to high CO2 for the infaunal communities (46), thus favoring high faunal biomass of a few adapted types. Instead, both meiofauna and macrofauna communities significantly declined in biomass (that is, by up to 90%). Although replacement of typical species of shallow sandy sediments by opportunistic species with different trophic functionalities occurred (Fig. 3, B and C), this did not lead to similar levels of faunal density and biomass. Furthermore, we found a long-lasting shift in microbial community composition and function. The high-CO2 venting caused a decrease in the whole hydrolytic capacity of the benthic communities, as revealed by measurements of the potential activity of aminopeptidases and esterases (Fig. 2 and fig. S3). These results match those on coastal sediments in mesocosms (47). Only the β-glucosidase responded by increased hydrolytic activities, as reported in previous experiments with bacterioplankton (48–51) and here also with transplant experiments. This enzyme, responsible for polysaccharide degradation, may be enhanced as a consequence of the higher microphytobenthic productivity (22). Furthermore, anaerobic remineralization by sulfate reducers was almost fully repressed in the vented sites (Fig. 2 and fig. S3), matching the substantial decline in sulfate reducer sequences. A sensitivity of sulfate-reducing bacteria to high CO2 was also previously observed at CO2-vented sediments off Papua New Guinea (32). Other functional groups affected by CO2 based on relative sequence abundance were sulfate oxidizers and nitrifiers.

Together, the observed CO2-leakage effects had consequences on carbon remineralization per biomass (R/B). This ratio was higher in the vented sands, indicating that more organic carbon needs to be remineralized per unit of biomass, compared to the reference (Table 3 and Fig. 5). Similarly, an altered food web structure and an impaired carbon cycling have been recently reported for soils affected by natural CO2 leakages (53). The results of the transplantation were in full accordance with the observed field patterns, showing basic CO2 effects on community biomass, composition, and biogeochemical function that will not be overcome by long-term adaptation of the involved species. Rather, the selection of opportunistic or tolerant species caused long-term deviation from reference ecosystem-level functions in terms of productivity, standing stock, and remineralization rates. There was an overall increase of productivity but also a higher respiration rate per standing stock, thus decreasing the biological carbon sink function. Furthermore, the quantitatively more relevant geochemical carbon sink was weakened by the carbonate dissolution and by the long-term loss of buffering capacity due to CO2 leakage.

CONCLUSION

Our study shows that CO2 leakage substantively changed the carbonate chemistry in permeable sandy sediments, increasing mineral weathering and nutrient flux (for example, iron and silicate). This led to local shifts in bacterial communities and enhanced microphytobenthos growth but also to a decline in benthic meiofauna and macrofauna density and composition. Together, CO2 leakage altered the ecosystem functions in terms of remineralization and carbon transfer along the food web. Hence, there is a substantial risk that CO2 leakage from submarine CCS sites may locally lead to negative impacts on the ecosystem and the function
of the seafloor as carbon sink, when globally, it would help mitigate the detrimental consequences of climate change and acidification on ocean ecosystems. CO2 leakage from submarine CCS would be very difficult, if not impossible, to stop. Therefore, site selection and spatial planning for marine CCS should include the environmental assessment of habitats, community composition, and ecological functions, including the resilience of local marine species, to mitigate ecological risks.

**MATERIALS AND METHODS**

**Site selection**

Locations with natural subseafloor CO2 venting are studied as analogs to assess the impacts of CO2 leakage as a risk of CCS technology for marine ecosystems (12). Natural CO2-vented sediments occur in active volcanic/tectonic regions and are driven by hydrothermal circulations (12). The selection of purely CO2-vented sediments for studies of high-CO2 effects is crucial to reduce confounding factors, such as presence of gases representing microbial energy sources (that is, H2, H2S, CH4, and C2H6), toxic elements (that is, metals), and temperature anomalies (54, 55). For example, some of the Mediterranean CO2 vents, such as the one in Levantine Bay of Volcano Island and some at Panarea, show high temperatures around 95°C and a relatively high H2S gas content of ca. 2%, complicating the isolation of pure CO2 effects at venting sites by favoring growth of extremophilic and thiotrophic microbial communities (56, 57). Before our study, we evaluated different CO2-vented habitats around Panarea Island to choose appropriate study sites for investigating the pure effects of CO2 leakage and seawater acidification on benthic life. During the first expedition (29 May to 10 June 2011), nine gas emission sites and two potential REFs were explored by diving in water depths of around 20 m depth. Larger areas were surveyed by manta towing (58). The divers sampled sediments and water asf for pH and sulfide measurements, measured temperature, and took documentary pictures and videos. At the study sites off Basiluzzo Islet, shallow sand flats were interspersed with seagrass meadows at 14- to 21-m depth. The CO2-impacted sites studied here were identified visually by the ebullition of CO2 from the seafloor. We found the east of Basiluzzo Islet to be suitable as a submarine natural analog to assess biogeochemical and ecological consequences of CO2 leakage because of the purity of the gas escaping from the seafloor, combined with the absence of cofounding hydrothermal effects altering microbial communities (Table 1). The experimental fieldwork was carried out in 2012 (02 to 21 June) and 2013 (01 to 14 June).

**Data analysis**

pH as a key variable for high CO2 is presented in pH2. We calibrated all sensors with commercial National Bureau of Standards buffers and corrected these to pH2 by subtracting 0.13 units, as described in Zeebe and Wolf-Gladrow (59). To assess significant differences between two or more independent data sets across the different sites, we used Welch’s t test (for example, sensors network and metadata analysis) or different ANOVAs (for example, biogeochemical and microbiological data and functional groups) (for details, see Statistical Analyses in the Supplementary Materials and Methods).

**Chemical analyses of vent gases and fluids, seawater, porewater, and sediments**

**Gases, seawater, and vent fluids**

Gas bubbles were sampled by scuba divers using exetainers and gas-collecting tubes made from glass, and analyzed via gas chromatography for CO2, O2, Ar, N2, CH4, C2H6, C3H8, COS, SO2, and H2. Hydrogen sulfide concentrations of these bubbles were determined by electrochemical sensor measurements directly after sampling (for details, see the Supplementary Materials). Seawater was sampled with a 5-liter Niskin bottle at ca. 30 cm asf and with 50-ml glass syringes at 5 to 10 cm asf. Benthic chamber water samples (that is, mixture of overlying bottom water and porewater) were collected with 50-ml glass syringes directly connected to benthic chambers at different times and analyzed for pH, nutrient [NH4+, PO43−, NO3−, NO2− + NO3−, and Si(OH)42−], sulfide, and Fe/Mn concentrations in the sediments were assessed by extracting porewater at 2-cm depth intervals with a TUBO device and with Rhizons (SMS type MOM, 19.21.21F; mean pore size, 0.15 μm; Rhizosphere Research Products) attached to 10-ml syringes. From three replicate push cores per site and per year, samples (2-cm intervals, maximum length up to 10 cm) were preserved for analyses of granulometry, porosity, TOC, TN, CPE (including Chl a), and calcium carbonate (CaCO3) content. The samples were preserved and analyzed as described in Bör et al. (61) (for details, see the Supplementary Materials). Chl a concentrations were converted to microphytobenthos biomass using a 1:40 Chl a–to–C biomass ratio (62).

**In situ measurements**

**Time-lapse recordings**

To monitor gas flow visually in situ, time-lapse photography was conducted (videos available at https://doi.pangaea.de/10.1594/PANGAEA.825241) (for details, see the Supplementary Materials).

**Oceanographic measurements**

A SEAGUARD recording current meter (Aanderaa Data Instruments) was used in 2011, 2012, and 2013 to monitor for 24 to 72 hours current velocity, temperature, salinity/conductivity, pressure, turbidity, and oxygen concentrations within the water column at about 30 cm asf.

**Bottom water chemistry loggers**

At five selected sites, the pH, oxygen (O2), ORP, and pressure (tides) were measured using five RBR loggers (RBR-Datalogger XR-420 D; RBR; www.rbr-global.com) with the sensors at 2 cm asf.

**Porewater chemistry**

A sediment microsensor profiler was equipped with sensors for pH (Microelectrodes Inc.) (63), O2 (64), CO2(aq) (Microelectrodes Inc.), ORP (a Pt wire, exposed tip is 50 μm thick and 0.5 mm long), temperature (Pt100; UST Umweltsensortechnik GmbH), H2 (Unisense), and H2S (65). The profiler was deployed at CO2-R, CO2-G, and REF in 2012. The CO2 sensors were damaged during the measurement at ca. 2 cm bsf at CO2-R; all other sensors worked successfully.

**Diffusive oxygen fluxes**

Diffusive oxygen fluxes were calculated from the in situ oxygen microprofiles using Fick’s law of diffusion as described previously (66). **Total oxygen uptake rates**

Benthic chambers were inserted into the seafloor to measure total fluxes of oxygen, nutrients, pH, and DIC within a defined sediment and seawater volume. At CO2-vented sites, the chambers were deployed between bubble streams to avoid the formation of internal headspace. We repeatedly subsampled the overlying water with syringes during daylight from transparent and masked chambers to assess
oxygen exchange (net \( \text{O}_2 \) flux) and oxygen respiration, respectively. Oxygen production, that is, gross primary production (GPP), was estimated as net \( \text{O}_2 \) flux + \( (\text{O}_2 \) respiration). To correct for anoxic fluid efflux from the seafloor, bags were attached to each chamber to measure the cumulative incubation volume (further information in the Supplementary Materials). Because of relatively high respiration rates, only incubations <6 hours were used to calculate oxygen fluxes.

**Sulfate reduction rates**

SRRs, as a proxy of anaerobic microbial respiration, were measured ex situ on sediments collected with push corers and sliced in 2-cm intervals (67).

**Extracellular enzymatic activities**

EEAs were analyzed by incubating the top 2 cm of the sediments with substrates for \( \beta \)-glucosidase, chitobiase, leucine aminopeptidase, and esterase (see further details in the Supplementary Materials) (61).

**Calcite dissolution rates (marble tiles)**

The corrosion of \( \text{CaCO}_3 \) structures by \( \text{CO}_2 \) was determined from the weight loss of marble tiles exposed at 0.5 m asf at REF \( \text{CO}_2\text{-G} \) and \( \text{CO}_2\text{-R} \) from June 2012 to June 2013. Once retrieved, the marble tiles were dried and weighed, and the dissolution rates were estimated as the difference in weight between pre- and postdeployment tiles, divided by exposure time (362 to 364 days; for further details, see the Supplementary Materials).

**Microbiological analyses**

**Sample collection and DNA extraction**

Samples from 0 to 2 cm bsf were obtained by 20 Sarstedt tubes (50 ml), and three push cores were collected for additional sections between 0 and 10 cm bsf per site and per year. Samples were kept frozen at \(-20^\circ\text{C}\) until subsequent analyses. DNA was extracted from 1 g of sediment per sample using the FastDNA SPIN Kit for Soil (Qbiogene), including a subsequent heating step to increase yield and final elution of the DNA in tris-EDTA buffer.

**Bacterial community structure**

The high-throughput fingerprinting technique ARISA [according to Ramette (68)] was applied to all sediment samples (0 to 2 cm bsf layer, \( n = 23 \); 2 to 4 cm, 4 to 6 cm, 6 to 8 cm, and 8 to 10 cm bsf layers, \( n = 3 \), per site and per year). In addition, 454 MPTS [according to Sogin et al. (69)] was used for the 0 to 2 and 4 to 6 cm bsf layers collected in 2012 (\( n = 3 \)) (for details, see the Supplementary Materials).

**Total microbial and microphytobenthos cell counts**

Sediment samples were fixed in 2% (for microbes) and 4% (for microphytobenthos) buffered formaldehyde/seawater and stored at 4°C until subsequent analysis. Microbial abundance was estimated by epifluorescence microscopy after staining with acridine orange (61). Catalyzed reporter deposition fluorescence in situ hybridization was applied for bacterial and archael cell enumeration. For microphytobenthos, only viable cells were counted under an inverted light microscope (Leica Microsystems AG) using 32× to 40× objective (final magnification, \( \times 320 \) to \( \times 400 \)) (see the Supplementary Materials).

**Fauna sampling and analysis**

Meiofauna samples (size class, 0.032 to 1 mm) were collected with pre-cut (2-cm horizontal) and taped push cores with an inner diameter of 4.7 cm in 2011 (\( n = 3 \) per site; upper 8 cm) or 5 cm in 2012 and 2013 (\( n = 3 \) per site; upper 8 cm). Samples were fixed and preserved in 4% buffered formaldehyde/seawater. Macrofauna samples (size class, >1 mm) were collected with push corers (inner diameter of 6.4 cm; \( n = 5 \) per site; upper 5 cm). Meiofauna and macrofauna organisms were identified to phylum or class level under a stereoscopic microscope. All meiofaunal nematodes and macrofaunal polychaetes were further identified to species level and allocated to functional feeding groups (see the Supplementary Materials). Depending on reproduction rate and tolerance to disturbance, nematodes were also allocated to a colonizer-persister (cp) category based on Bongers et al. (70): cp-2, short generation time, high reproduction rate, and very tolerant to disturbances; cp-3, characteristics intermediate to cp-2 and cp-4 and relatively sensitive to disturbances; cp-4, long generation time and sensitive to pollutants; and cp-5, long life span, low reproduction rate, and very sensitive to pollutants and other disturbances. The nematodes’ maturity index was determined as an ecological measure of environmental disturbance (see the Supplementary Materials) (70).

**Sediment transplantation experiments**

Sediments were transplanted in situ within and between REF and \( \text{CO}_2\text{-R} \) sites: (i) reimplanted at the same site (within habitat: REF/REF and \( \text{CO}_2\text{-R}/\text{CO}_2\text{-R} \) to control for transplantation effects, and (ii) reimplanted to the other habitat type (across habitat: REF/\( \text{CO}_2\text{-R} \) and \( \text{CO}_2\text{-R}/\text{REF} \)) to assess the effect of the different environmental setting. Samples for microbial community analyses (that is, ARISA) were collected immediately after transplantation (\( n = 3 \) per treatment), after 2 weeks (\( n = 1 \) per treatment), and after 1 year (\( n = 3 \) per treatment). Samples for porewater composition, sediment geochemistry, microbial activity, and nematode abundance and community structure were collected 1 year after transplantation (\( n = 3 \) per treatment). Samples were taken and analyzed as described in the above sections, except for nematodes that were identified at genus level.

**SUPPLEMENTARY MATERIALS**

Supplementary material for this article is available at http://advances.sciencemag.org/cgi/content/full/4/2/eaao2040/DC1

Supplementary Materials and Methods

fig. S1. Maps and seafloor images of the sampling area.

fig. S2. Bottom water and porewater chemistry at the study sites off Basiluzzo Islet.

table S1. Bottom water and porewater chemistry at the study sites off Basiluzzo Islet.

table S2. Outcome of one-way ANOVA.

table S3. \( \text{pH} \), change and in situ fluxes obtained from benthic chambers deployed in 2013.

table S4. Change in relative sequence number of highly abundant, putative functional groups of bacteria based on 454 MPTS OTUs annotation (OTU > 0.1\%).

table S5. Outcome of distance-based multivariate regression analysis.

table S6. Outcome of permutational ANOVA for environmental variables (Env. Setting).

table S7. Relative abundances of macrofauna taxa, polychaete species, meiofauna taxa, and nematode species at the three investigated sites.

table S8. Differences in porewater and sedimentary environmental settings in medium-term transplant experiments (\( n = 3 \)).

table S9. Overview of \( \text{CO}_2/\text{pH} \) impacts on benthic organisms at natural \( \text{CO}_2 \) vents, as observed in previous studies.

References (72–160)

**REFERENCES AND NOTES**

1. H.-O. Pörtner, D. M. Karl, P. W. Boyd, W. W. L. Cheung, S. E. Lluch-Cota, Y. Nojiri, D. N. Schmidt, P. O. Zavialov, Ocean systems, in Climate Change 2014: Impacts, Adaptation, and Vulnerability. Part A: Global and Sectoral Aspects. Contribution of Working Group II to the Fifth Assessment Report of the Intergovernmental Panel on Climate Change, C. B. Field, V. R. Barros, D. J. Dokken, K. J. Mach, M. D. Mastrandrea, T. E. Bilir, M. Chatterjee, K. L. Ebi, Y. O. Estrada, R. C. Genova, B. Girma, E. S. Kissel, A. N. Levy, S. MacCracken, P. R. Mastrandrea, L. L. White, Eds. (Cambridge Univ. Press, 2014), pp. 411–484.
2. J.-P. Gattuso, K. J. Mach, G. Morgan, Ocean acidification and its impacts: An expert survey. Clim. Change 117, 725–738 (2013).
3. IEA, Energy and Climate Change. World Energy Outlook Special Report (WEO2015), 15 June 2015.
4. S. Holloway, Storage of fossil fuel-derived carbon dioxide beneath the surface of the Earth. Annu. Rev. Energy Environ. 26, 145–166 (2001).
5. J. Blackford, J. M. Bull, M. Cevatoglu, D. Connelly, C. Houton, R. H. James, A. Lichtschlag, H. Stahl, S. Widdicombe, I. C. Wright, Marine baseline and monitoring strategies for carbon dioxide capture and storage (CCS). Int. J. Greenh. Gas Control 38, 221–229 (2015).
6. H. Al Botnen, A. M. Oman, I. Thorsset, T. Johannessen, G. Alendal, The effect of submarine CO2 vents on seawater: Implications for detection of subsea carbon sequestration leakage. Limnol. Oceanogr. 60, 402–410 (2015).
7. D. G. Jones, S. E. Beaubien, J. C. Blackford, E. M. Foekema, J. Lions, C. De Vittor, J. M. West, S. Widdicombe, C. Houton, A. M. Queirós, Developments since 2005 in understanding potential environmental impacts of CO2 leakage from geological storage. Int. J. Greenh. Gas Control 40, 350–377 (2015).
8. S. Widdicombe, S. Dupont, M. Thorndyke, Experimental design of perturbation experiments Laboratory experiments and benthic mesocosm studies, in Guide to best Practices for Ocean Acidification Research and Data Reporting, U. Riebesell, V. J. Fabry, L. Hansson, J.-P. Gattuso, Eds. (Publications Office of the European Union, 2010), pp. 113–122.
9. J. Blackford, H. Stahl, J. M. Bull, B. J. Bergès, M. Cevatoglu, A. Lichtschlag, D. Connelly, R. H. James, J. Kita, D. Long, M. Naylor, K. Shitashima, D. Smith, P. Taylor, I. Wright, M. Akhurst, B. Chen, T. M. M. Gurney, C. Houton, M. Hayashi, H. Kaieda, T. G. Leighton, T. Sato, M. D. J. Sayer, M. Suzumura, K. Tait, M. E. Vardy, P. R. White, S. Widdicombe, Detection and impacts of leakage from sub-seafloor deep geological carbon dioxide storage. Nat. Clim. Change 4, 1011–1016 (2014).
10. J.-P. Gattuso, A. Magnan, R. Bille, W. W. L. Cheung, E. L. Howes, F. Joos, D. Allemand, L. Bopp, S. R. Cooley, C. M. Eakin, O. Goehl-Guldford, R. P. Kelly, H.-O. Pörtner, A. D. Rogers, J. M. Baxter, D. Laffoley, D. Osborn, A. Rannkovic, J. Rochette, U. S. Runnala, S. Treyer, C. Turley, Contrasting futures for ocean and society from different anthropogenic CO2 emissions scenarios. Science 349, aa4722 (2015).
11. S. Widdicombe, C. L. McNeill, H. Stahl, P. Taylor, A. Queirós, J. Nunes, K. Tait, Impact of sub-seabed CO2 leakage on macrobenthic community structure and diversity. Int. J. Greenh. Gas Control 38, 182–192 (2015).
12. F. Inagaki, M. M. Kuyper, U. Tsunogai, K.-i. Ishibashi, K.-i. Nakamura, T. Treude, S. Ohkubo, M. Nakaseama, K. Gena, H. Chiba, H. Hirayama, T. Nunoura, K. Takai, B. B. Jørgensen, K. Horikoshi, A. Boetius, Microbial community in a sediment-hosted CO2 lake of the southern Okinawa Trough hydrothermal system. Proc. Natl. Acad. Sci. U. S. A. 103, 14164–14169 (2006).
13. J. M. Sunday, K. E. Fabricius, K. J. Kroeze, K. M. Anderson, N. E. Brown, J. P. Barry, S. D. Connell, S. Dupont, B. Gaylord, J. M. Hall-Spencer, T. Klinger, D. Long, M. Milazzo, A. M. Queirós, Response of two marine bacterial isolates to high CO2 concentration. PLoS ONE 7, e47035 (2012).
14. K. Takeuchi, Y. Fujioka, Y. Kawasaki, S. Shiayaama, Impacts of high concentration of CO2 on marine organisms; a modification of CO2 ocean sequestration. Energy Convers. Manage. 38, S337–S341 (1997).
15. S. L. Dallmann, P. J. Somerfield, S. Widdicombe, M. C. Austen, M. Nimmo, Impacts of ocean acidification and burrowing urchins on within-sediment pH profiles and subtidal nematode communities. J. Exp. Mar. Biol. Ecol. 365, 46–52 (2008).
16. J. P. Barry, T. Tyrrell, L. Hansson, G. K.-P. Plattner, J.-P. Gattuso, Experimental design of perturbation experiments, in Guide to best Practices for Ocean Acidification Research and Data Reporting, U. Riebesell, V. J. Fabry, L. Hansson, J.-P. Gattuso, Eds. (Publications Office of the European Union, Luxembourg, 2010), pp. 53–136.
17. H. Ishida, Y. Watanabe, Y. Kuroda, K. Nakano, K. Furusawa, Y. Shiayaama, In situ enclosure experiment using a benthic chamber system to assess the effect of high concentration of CO2 on deep-sea benthic communities. J. Oceanogr. 61, 835–843 (2005).
18. J. W. Flegel, D. S. Johnson, K. R. Carman, P. B. Weisenhorn, A. G. Thistle, J. P. Barry, The response of nematodes to deep-sea CO2 sequestration: A quantile regression approach. Deep Sea Res. Part I Oceanogr. Res. Pap. 57, 696–707 (2010).
19. E. Kousse, A. Wichels, L. Gimenez, M. Lunar, M. B. Schilhab, G. Gerdes, Small changes in pH have direct effects on marine bacterial community composition: A microcosm approach. PLOS ONE 7, e47035 (2012).
20. A.-S. Roy, S. M. Gibbons, H. Schunck, S. Owens, J. G. Caporaso, M. Spering, J. I. Nissimov, S. Romac, L. Bittner, U. Riebesell, J. LaRoche, J. A. Gilbert, Ocean acidification shows negligible impacts on high-latitude bacterial community structure in coastal pelagic mesoscoops. Biogeosci. Discuss. 9, 1313–1334 (2012).
21. A. Monier, H. S. Findlay, S. Charvet, C. Lovejoy, Late winter under ice pelagic microbial communities in the high Arctic Ocean and the impact of short-term exposure to elevated CO2 levels. Front. Microb. 5, 490 (2014).
22. A. Chauhan, A. Pathak, R. Rodolfo-Metalpa, M. Milazzo, S. J. Green, J. M. Hall-Spencer, Metagenomics reveals planktonic bacterial community shifts across a natural CO2 gradient in the Mediterranean Sea. Genome Announc. 3, e01543 (2015).
23. K. Yanagawa, Y. Morono, D. de Beer, M. Haeckel, M. Sunamura, T. Futagami, T. Hoshino, T. Terada, K.-i. Nakamura, T. Urabe, G. Rehder, A. Boetius, F. Inagaki, Metabolically active microbial communities in marine sediment under high-CO2 and low-pH extremes. ISME J. 7, 555–567 (2013).
24. D. Kerfahi, J. M. Hall-Spencer, B. M. Tripathi, M. Milazzo, J. Lee, M. Adams, Shallow water marine sediment bacterial community shifts along a natural CO2 gradient in the Mediterranean Sea off Vulcano, Italy. Microb. Ecol. 67, 819–828 (2014).
25. K. M. Morow, D. G. Bourne, C. Humphrey, E. S. Bottle, P. Laffy, J. Zanenfeld, S. Uthicke, K. E. Fabricius, N. S. Webster, Natural volcanic CO2 seeps reveal future trajectories for host-microbial associations in corals and sponges. ISME J. 9, 894–908 (2015).
26. J. P. Bowman, The marine clade of the family Flavobacteriaceae: The genera Aequorivita, Aemobacter, Cellulophaga, Croceibacter, Formosa, Gelidibacter, Gellisia, Maribacter, Menosina, Muricauda, Polaribacter, Psychroflexus, Psychrosphenus, Robiginitalea, Salgengibacter, Tenacibaculum, Ulvibacter, Villettibacter and Zobelia. Prokaryotes Vol. 7, 677–694 (2006).
27. E. Teira, A. Fernández, X. A. Álvarez-Salgado, E. E. García-Martin, P. Serret, C. Sobrino, Response of two marine bacterial isolates to high CO2 concentration. Mar. Ecol. Prog. Ser. 453, 27–36 (2012).
28. F. F. Raulf, K. Fabricius, S. Uthicke, D. de Beer, R. M. M. Abed, A. Ramette, Changes in microbial communities in coastal sediments along natural CO2 gradients at a volcanic vent in Papua New Guinea. Environ. Microbiol. 17, 3678–3691 (2015).
29. T. Yamada, Y. Sekiguchi, S. Hanada, H. Imachi, A. Ohashi, H. Harada, Y. Kamagata, Anaerolimna thermosilosa sp. nov., Levilinex salinocatlicola gen. nov., sp. nov. and Leptolina tardativallis gen. nov., sp. nov., novel filamentous anaerobes, and description
of the new classes Anoequalinaceae nov. and Caldimellinaceae nov. in the bacterial class Euryarchaeota. Int. J. Syst. Evol. Microbiol. 56, 1331–1340 (2006).

45. A. Schippers, D. Kock, C. Höpf, G. Kóweker, M. Siegert, Quantification of microbial communities in subsurface marine sediments of the Black Sea and off Namibia. Front. Microbiol. 3, 16 (2012).

46. S. L. Garrard, M. C. Gambi, M. B. Scipione, F. P. Patti, M. Lorenti, V. Zupor, D. M. Paterson, M. C. Buia, Indirect effects may buffer negative responses of seagrass invertebrate communities to ocean acidification. J. Exp. Mar. Bio. Ecol. 461, 31–38 (2014).

47. E. Rastelli, C. Corinaldesi, A. Dell’anno, A. Tamaro, S. Greco, M. Lo Martire, L. Carugati, A. M. Queris, S. Widdicombe, R. Danovaro, CO2 leakage from carbonate capture and storage (CCS) systems affects organic matter cycling in surface marine sediments. Mar. Environ. Res. 122, 158–168 (2016).

48. H-P. Grossart, M. Allgaier, U. Passow, U. Riebesell, Testing the effect of CO2 concentration on the dynamics of marine heterotrophic bacterioplankton. Limnol. Oceanogr. 51, 1–11 (2006).

49. J. Piontek, M. Lunau, N. Händel, C. Borchard, M. Wurst, A. Engel, Acidification increases extremely high CO2 exposure. Limnol. Oceanogr. 59, 103–119 (2008).

50. M. M. Sala, F. L. Aparicio, V. Balagué, J. A. Boras, E. Borrull, C. Cardelús, L. Cros, A. Gomes, A. López-Sanz, A. Malts, R. A. Martinez, M. Mestre, J. Movilla, H. Sarmento, E. Vázquez-Domínguez, D. Vaqué, J. Pinhasi, A. Calbet, E. Calvo, J. M. Gasol, C. Pelejerio, C. Marrasé, Contrasting effects of ocean acidification on the microbial food web under different trophic conditions. ICES J. Mar. Sci. 73, 670–679 (2016).

51. C. Hassennack, A. Fink, A. Lichtschlag, H. E. Tegtemeyer, D. De Beer, A. Ramette, Quantification of the effects of ocean acidification on sediment microbial communities in the environment: The importance of ecosystem approaches. FEMS Microbiol. Ecol. 92, fiw027 (2016).

52. F. Beulig, T. Urich, M. Nowak, S. E. Trumbore, G. Sleekher, G. D. Gillflan, K. E. Fjelland, K. Küsel, Altered carbon turnover processes and microbiomes in soils under long-term extremely high CO2 exposure. Nat. Microbiol. 1, 15025 (2016).

53. F. Wenzhöfer, O. Holby, R. N. Guld, H. K. Nielsen, J. K. Gundersen, In situ microsensor studies of a shallow water hydrothermal vent at Milos, Greece. Mar. Chem. 69, 43–54 (2000).

54. S. Vizziènz, B. Di Leonardo, V. Costa, C. D. Tromati, F. Luzzo, A. Mazzola, Trace element bias in the use of CO2 vents as analogues for low pH environments: Implications for contamination levels in acidified oceans. Estuar. Coast. shelf Sci. 134, 19–30 (2013).

55. J. P. Amend, K. L. Rogers, E. L. Shock, S. Gurrieri, S. Ingagiuggio, Energetics of chemolithoautotrophy in the hydrothermal system of Vulcano Island, southern Italy. Geology 1, 37–58 (2003).

56. R. E. Price, D. E. Larrone, F. Italiano, I. Savoy, T. Pichler, J. P. Amend, Subsurface hydrothermal processes and the bioenergetics of chemolithoautotrophy at the shallow-sea vents off Panarea Island (Italy). Chem. Geol. 407–408, 21–45 (2015).

57. S. English, C. Wilkinson, V. Baker, Survey Manual for Tropical Marine Resources (Australian Institute of Marine Science, ed. 2, 1997).

58. R. Zeebe, D. Wolf-Gladrow, Calculation of CO2 and NH4+ in marine and freshwaters. Nitrogen 45, 169 (1995).

59. M. L. Sogin, H. G. Morrison, J. A. Huber, D. M. Welch, S. M. Huse, P. R. Neal, J. M. Arrieta, M. A. Rohwer, E. Roesch, S. Han, A. K希望大家好！
93. H. Utermöhl, Zur Vervollkommnung der quantitativen Phytolankton-Methode. Int. Ver. The. 9, 1–38 (1958).

94. F. F. Round, R. M. Crawford, D. G. Mann, The Diatoms: Biology and Morphology of the Genera (Cambridge Univ. Press, 1990).

95. T. Cibic, O. Blasutto, S. Fonda Umani, Biodiversity of settled material in a sediment trap in the gulf of Trieste (northern Adriatic Sea). Hydrobiologia 580, 57–75 (2007).

96. F. Widdel, F. Bak, Gram-Negative Mesophilic Sulfate-Reducing Bacteria, in The Prokaryotes, A. Balows, H. G. Trüper, M. Dworkin, W. Harder, K. H. Schleifer, Eds. (Springer-Verlag, 1992), pp. 3352–3378.

97. J. W. Seinhorst, A rapid method for the transfer of nematodes from fixative to anhydrous glycerin. Nematologica 4, 67–69 (1959).

98. K. Gallini, T. N. Bezerra, U. Eisenack-Flückner, T. Deprez, G. Fonseca, O. Holovachov, D. Leduc, D. Mijlant, T. Moens, J. Sharma, N. Smol, A. Tchesunov, V. Mokleivsky, J. Vanaverbeke, A. Vanreusel, V. Venekens, M. Vinck, NemyS: World Database of Free-Living Marine Nematodes. (2017); http://nemys.ugent.be.

99. G. Read, K. Fauchald, World Polychaeta database (2016); http://www.marinespecies.org/polychaeta.

100. J. C. F. Gil, The European fauna of Annellida Polychaeta, (Fac. Ciências. Univ. Lisboa, 2011); http://hdl.handle.net/10451/4600.

101. L. Andrassy, The determination of volume and weight of nematodes. Acta Zool. Hung. 2, 1–15 (1956).

102. C. Heip, M. Vincx, G. Vranken, The ecology of marine nematodes. Nematologica 40, 23–39 (1993).

103. H. Utermöhl, Zur Vervollkommnung der quantitativen Phytoplankton-Methodik. FEMS Microbiol. Ecol. 15, 846–858 (1996).

104. B. T. Hargrave, V. E. Kostylev, C. M. Hawkins, Benthic epifauna assemblages, biomass and respiratory in the Gully region on the Scotian Shelf, NW Atlantic Ocean. Mar. Ecol. Prog. Ser. 270, 55–70 (2004).

105. W. Wieser, Die beziehung zwischen mundhöhlengestalt, ernährungsweise und polychaeta. Nematologica 58, 249–272 (2011).

106. A. Balows, H. G. Trüper, M. Dworkin, W. Harder, K. H. Schleifer, Eds. (Springer-Verlag, 1996).

107. I. Andrassy, The determination of volume and weight of nematodes. Acta Zool. Hung. 2, 439–483 (1953).

108. P. A. Jumars, K. M. Dorgan, S. M. Lindsay, Diet of worms emended: An update of polychaete feeding guilds. Annu. Rev. Mar. Sci. 7, 497–520 (2015).

109. R.Development-Core-Team, R: A Language and Environment for Statistical Computing. Found. Stat. Comput. Vienna, Austria 2014; http://www.r-project.org/.

110. K. Clarke, R. N. Gorley, R. K. Clarke, R. K. Clarke, R. N. Gorley, Primer v6: User Manual/Tutorial (Plymouth Marine Laboratory, 2006).

111. M. J. Anderson, R. N. Gorley, R. K. Clarke, M. C. Gambi, Spatial-temporal variability of polychaete colonization at volcanic CO2 vents indicates high tolerance to ocean acidification. Mar. Biol. 161, 2909–2919 (2014).

112. R. Rodolfo-Metalpa, F. Houlbique, É. Tambutté, F. Boisson, C. Baggini, F. P. Patti, R. Jeffere, M. Fine, A. Foggo, J.-P.Gattuso, J. M. Hall-Spencer, Coral and mollusc resistance to ocean acidification adversely affected by warming. Nat. Clim. Change 1, 308–312 (2011).

113. T. Arnold, C. Mealey, H. Leahey, A. W. Miller, J. M. Hall-Spencer, M. Milazzo, K. Maers, Intertidal epilithic bacteria response of foraminifera to high-CO2 conditions in the Mediterranean Sea. Proc. Natl. Acad. Sci. U.S.A. 107, 14520–14525 (2010).

114. J. D. Taylor, R. Ellis, M. Milazzo, J. M. Hall-Spencer, M. Cunliffe, Intertidal epilithic bacteria activity in sea anemones under high pCO2 conditions. FEMS Microbiol. Ecol. 69, 620–627 (2010).

115. T. Cibic, O. Blasutto, S. Fonda Umani, Biodiversity of settled material in a sediment trap in the gulf of Trieste (northern Adriatic Sea). Hydrobiologia 580, 57–75 (2007).

116. R. Danovaro, Prokaryote diversity and virus abundance in shallow hydrothermal systems in sea water version 2 (2011); http://nemys.ugent.be.

117. I. lidbury, J. v. d. Bern, J. M. Hall-Spencer, C. B. Munn, M. Cunliffe, Community-level response of coastal microbial biofilms to ocean acidification in a natural carbon dioxide vent system. Ocean. Biol. 23, 440–446 (2008).

118. D. Meron, E. Atias, L. Iasur Kruh, H. Elifantz, D. Minz, M. Fine, E. Banin, The impact of reduced pH on the microbial community of the coral Acropora eurystoma. ISME J. 5, 51–60 (2011).

119. A. Cunliffe, M. C. Gambi, Settlement pattern of Posidonia oceanica epibionts along a gradient of ocean acidification: An approach with mimics. Mediterr. Mar. Sci. 15, 498–509 (2014).

120. E. Ricevuto, K. J. Kroeker, F. Ferrigno, F. Micheli, M. C. Gambi, Spatio-temporal variability of polychaete colonization at volcanic CO2 vents indicates high tolerance to ocean acidification. Mar. Biol. 161, 2909–2919 (2014).

121. L. Basso, I. E. Hendriks, A. B. Rodríguez-Navarro, M. C. Gambi, C. M. Duarte, Extreme pH conditions at a natural CO2 vent system (Italy) affect growth, and survival of juvenile pen shells (Pinna nobilis). Estuaries Coasts. 15, 1986–1999 (2015).

122. R. Rodolfo-Metalpa, F. Houlbique, É. Tambutté, F. Boisson, C. Baggini, F. P. Patti, R. Jeffere, M. Fine, A. Foggo, J.-P.Gattuso, J. M. Hall-Spencer, Coral and mollusc resistance to ocean acidification adversely affected by warming. Nat. Clim. Change 1, 308–312 (2011).

123. L. Donnarumma, C. Lombardi, S. Cocito, M. C. Gambi, Settlement pattern of Posidonia oceanica epibionts along a gradient of ocean acidification: An approach with mimics. Mediterr. Mar. Sci. 15, 498–509 (2014).

124. D. Meron, E. Banin, Changes in microbial communities associated with the sea anemone Anemonia viridis in a natural pH gradient. Microb. Ecol. 65, 269–276 (2013).

125. E. Manini, G. M. Luna, C. Corinaldesi, D. Zeppilli, G. Bortoluzzi, G. Caramanna, F. Raffa, R. Danovaro, Prokaryote diversity and virus abundance in shallow hydrothermal vents of the Mediterranean Sea (Panarea Island) and the Pacific Ocean (North Sulawesi-Indonesia). Microb. Ecol. 55, 626–639 (2008).
Microbial and biogeochemical data. K.G. and D.M. analyzed porewater element composition at doi.pangaea.de/10.1594/PANGAEA.847916, and archived on the PANGAEA database: doi.pangaea.de/10.1594/PANGAEA.871453. Porewater element composition can be found at doi.pangaea.de/10.1594/PANGAEA.847916, and archived using the brokerage service of the German Planck Society and by the Flemish Fund for Scientific Research (grant number 124214N).

Acknowledgments: The authors would like to thank B. Merkel (TU Freiberg) for the sediment element analysis in 2011, F. Italiano (Istituto Nazionale di Geofisica e Vulcanologia (INGV) Palermo) for the vent fluid analysis in 2011, S. Beaubien and S. Lombardi (UniRoma1) for the on-site analysis of H2S gas in 2012, N. Bigalke (GOEMAR) for the estimation of the gas emission rates, M. Haeckel and R. Surberg (GEOMAR) for the porewater element analysis, J. Gil [Centre d’Estudis Avançats de Blanes, Consejo Superior de Investigaciones Científicas (CEAB-CSIC)] for his help in identifying some problematic polychaetes, and D. Wolf-Gladrow for the helpful discussions of the carbonate system. The authors are also grateful for the technical support by M. Meiners, E. Weiz-Bersch, M. Alisch, W. Stiens, and R. Stiens (HGF-MPG Joint Research Group on Deep Sea Ecology and Technology); help in field work and laboratory activities by N. Vaene, B. Beuselinck, A. Van Kenhove, G. De Smet, F. Sedano Vera, N. De Jesr, and L. Lins (UGent Marine Biology); scientific diving assistance by B. Unger, M. Schneider, H. Kuhluss, and A. Eich (HYDRA Institute for Marine Sciences); and logistic support by A. Fogliuzzi (Amphibia). Funding: This work was funded by the European Union Seventh Framework Programme (FP7/2007-2013) under grant agreement number 265847 [Sub-seabed CO2 storage: Impact on Marine Ecosystems (ECO2)] and supported by the Max Planck Society and by the Flemish Fund for Scientific Research (grant number 124214N).

This study is also a contribution of D.M. to the research project MarSymBiotics (reference number CTM2013-43287-P), funded by the Spanish “Agencia Estatal de Investigación” (AEI), and PopCoMics (CTM2017-88080), funded by the AEI and the European Funds for Regional Development (FEDER) and to the Consolidated Research Group on Marine Benthic Ecology (2014SR120) of the Generalitat de Catalunya. Author contributions: This study was designed by A.B. and A.R. Sampling activities and in situ experiments were performed by M.W., C.L., K.G., S.M., M.M., D.d.B., F.W., and A.R. M.M., D.d.B., and F.W. analyzed the in situ measurements. M.M. and G.W. analyzed the microbial and biogeochemical data. K.G. and D.M. analyzed the meiofauna and macrofauna. T.C. and C.D.V. analyzed the microphytobenthos. C.L. and M.W. analyzed the videos and images of the study sites. A.B., A.V., D.d.B., and M.W. contributed reagents/materials/analysis tools. The paper was written by M.M., K.G., and A.B. with contributions and final approval of all co-authors. Competing interests: The authors declare that they have no competing interests. Data and materials availability: All data were archived on the PANGAEA database: doi.pangaea.de/10.1594/PANGAEA.871453. Porewater element composition can be found at doi.pangaea.de/10.1594/PANGAEA.847916, and sediment element composition at doi.pangaea.de/10.1594/PANGAEA.847825. MPTS sequences were deposited on the European Nucleotide Archive under accession number PRJEB21026. The sequences were archived using the brokerage service of the German Federation for Biological Data (71). All data needed to evaluate the conclusions in the paper are present in the paper and/or the Supplementary Materials. Additional data related to this paper may be requested from the authors.

Submitted 29 June 2017
Accepted 5 January 2018
Published 7 February 2018
10.1126/sciadv.aao2040

Citation: M. Molari, K. Guilini, C. Lott, M. Weber, D. de Beer, S. Meyer, A. Ramette, G. Wegener, F. Wenzhöfer, D. Martin, T. Cibic, C. De Vittor, A. Vanreusel, A. Boetius, CO2 leakage alters biogeochemical and ecological functions of submarine sands. Sci. Adv. 4, eaa02040 (2018).
CO₂ leakage alters biogeochemical and ecological functions of submarine sands

Massimiliano Molari, Katja Guillini, Christian Lott, Miriam Weber, Dirk de Beer, Stefanie Meyer, Alban Ramette, Gunter Wegener, Frank Wenzhöfer, Daniel Martin, Tamara Cibic, Cinzia De Vittor, Ann Vanreusel and Antje Boetius

Sci Adv 4 (2), eaao2040.
DOI: 10.1126/sciadv.aao2040