



Report of epibenthic macrofauna found from Haima cold seeps and adjacent deep-sea habitats, South China Sea

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Abstract

This work reports on a preliminary taxonomic study of epibenthic macroinvertebrates collected or observed by underwater video at the Haima cold seeps and in adjacent deep-sea habitats, including a mud volcano field and Ganquan Plateau, during an expedition in the South China Sea by the Chinese-manned submersible *Shenhai Yongshi* in May 2018. A total of 41 species belonging to 6 phyla were identified, among which 34 species were collected from the Haima cold seeps. Mollusks and crustaceans that are specialized in reducing habitats were predominant in biotopes of the Haima cold seeps, whereas sponges and cold-water corals and their commensals were prominent in communities of the mud volcano field and the slopes of Ganquan Plateau. The distribution and faunal composition of each taxonomic group are discussed.

Keywords Cold seep · Mud volcano · Ganquan plateau · Epibenthic macroinvertebrates · Faunal composition · South China Sea

Introduction

Cold seeps are areas of the seafloor, where hydrocarbon-rich fluid and gases leak from fissures and emerge through the sediments and into the water column, creating unique habitats. Such seepage was first discovered on the Florida Escarpment in the Gulf of Mexico (Paull et al. 1984). Since then, hundreds of cold-seep sites have been discovered and observed globally (e.g., Feng et al. 2018; German et al. 2011; Suess 2014), unveiling a specialist seepage macrofauna. Cold-seep macrofauna, being sustained by chemosynthetic

primary production, typically consists of a high abundance of symbiotrophic organisms (Barry et al. 1996; Carney 1994; Levin 2005; Sibuet and Olu 1998; Washburn et al. 2018). Hence, cold seeps exhibit a community structure that is distinct from that seen in the surrounding seafloor environment. During the past three decades, numerous studies have been performed, driven by efforts to explore these special habitats and their associated organisms. Some studies have focused on the taxonomy and phylogeny of organisms associated with cold seeps, aiming to discover new species and new distribution records, evaluate phylogenetic relationships, and reconstruct the origins and evolutionary histories of seepage faunas (e.g., Chen et al. 2018; Dong and Li 2015; Xu et al. 2019). Other studies have focused on the community structures and temporal dynamics of cold seeps, analyzing the correlation between pattern of biodiversity in the infaunal assemblages and environmental factors such as depth and seepage types (e.g., Bourque et al. 2017; Cunha et al. 2013; Levin et al. 2015; Washburn et al. 2018).

The South China Sea is a marginal sea in the western Pacific Ocean with passive continental margins in the west and north, where various cold-seep sites have been discovered, in particular on the continental slope (Fang et al. 2019; Feng and Chen 2015; Li 2015, 2017; Niu et al. 2017). The first active seepage site to be discovered in the

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South China Sea is known as Site F (also called Jiaolong Seep No. 1, Formosa Ridge or Taixinan cold seep), located in the northeastern region (Lin et al. 2007). In the summer (June–July) of 2013, the corresponding author (Xinzheng Li) participated in the first cruise with experimental applications of the Chinese-manned submersible “*Jiaolong*” in the South China Sea (China Ocean Voyage No. 31). The epibenthic community at Site F was described as being dominated by the alvinocaridid shrimp *Alvinocaris longirostris* Kikuchi & Ohta, 1995, the mytilid mollusk *Gigantidas platifrons* (Hashimoto & Okutani, 1994), the galatheid squat lobster *Shinkaia crosnieri* Baba & Williams, 1998, and other squat lobster and mollusk species (Li 2017). From 2013 to 2018, various Chinese research institutions carried out a series of surveys at this site and gathered numerous macrobenthic specimens using remote operated vehicles (ROV) and the manned submersible *Jiaolong*. To date, 33 epibenthic macroinvertebrate species have been reported from Site F and its rim area, six of which are new to science, revealing high biodiversity and most likely a high level of species endemism (Chan and Komai 2017; Dong and Li 2015; Gong et al. 2015; Li 2017; Sha 2019; Zhang et al. 2016; Zhang and Zhang 2017).

In 2013, a new active cold-seep site was discovered on Four-Way Closure Ridge, not far from Site F (Klaucke et al. 2015). In 2015, another active cold-seep field in the South China Sea was identified on the northwestern continental slope, characterized by patches of authigenic carbonate rocks protruding from the muddy seafloor, which contains at least two seepage sites (Fang et al. 2019; Liang et al. 2017). This area was named the Haima cold seeps and has attracted increasing attentions since its discovery. To date, four new bivalve species have been described based on specimens

collected through surveys made at Haima in the past four years (Chen et al. 2018; Jiang et al. 2019; Xu et al. 2019). However, information on other taxonomic groups and community structure found in this field is scarce.

The expedition (Fig. 1) to the Haima cold seeps conducted in May 2018 was organized by Tongji University and Chinese Academy of Sciences. Additional cold-seep sites were discovered, including one on the southwestern slope of Ganquan Plateau, close to the Haima field. The expedition also explored an adjacent mud volcano field, and the slopes of Ganquan Plateau (Fig. 2). The corresponding author (Xinzheng Li) took part in the expedition and dived with the submersible “*Shenhai Yongshi*” (*Deep-Sea Warrior*) at the Haima cold-seep sites. Abundant epibenthic macroinvertebrate specimens were collected or observed in situ using the manned submersible. Some stalked barnacles were subsequently described by Gan and Li (2019). The present study summarizes and reports the taxonomic findings on epibenthic macrofauna from Haima cold seeps and nearby ecosystems, based on specimens collected and photographed during the expedition, to provide an overview of the species composition and biodiversity characteristics of the macrofaunal assemblages in these deep-sea habitats.

Results and discussion

Faunal composition and species list

A total of 41 species (Table 1) were identified to genus or species level from specimens either collected or observed in situ during the expedition to the slopes of Ganquan Plateau, the Haima cold-seep sites, and a mud volcano field. These comprise 16 species of Crustacea; 14 Mollusca;

Fig. 1 Location map of the study area in the South China Sea. *H* Haima cold-seep field; *N* northeastern slope of Ganquan Plateau; *M* mud volcano field, *S* southwestern slope of Ganquan Plateau

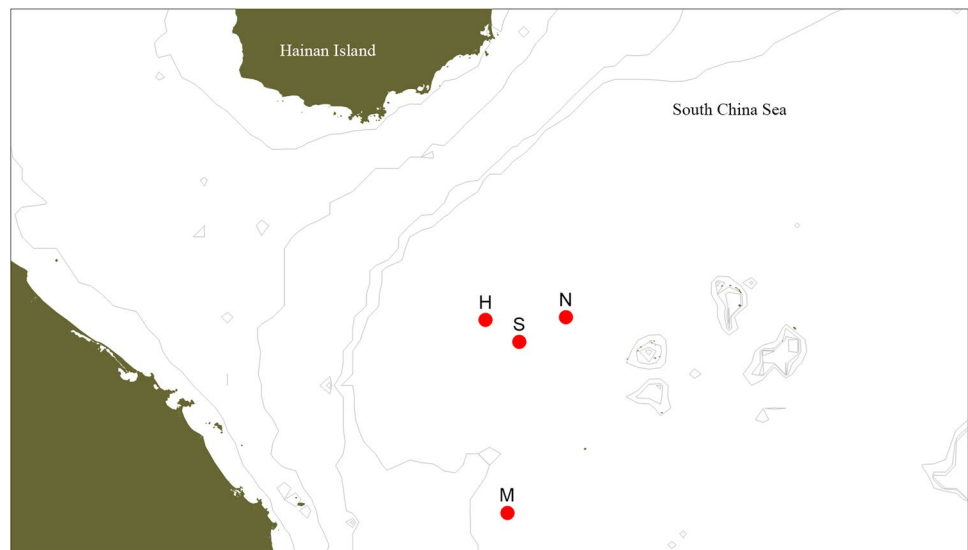
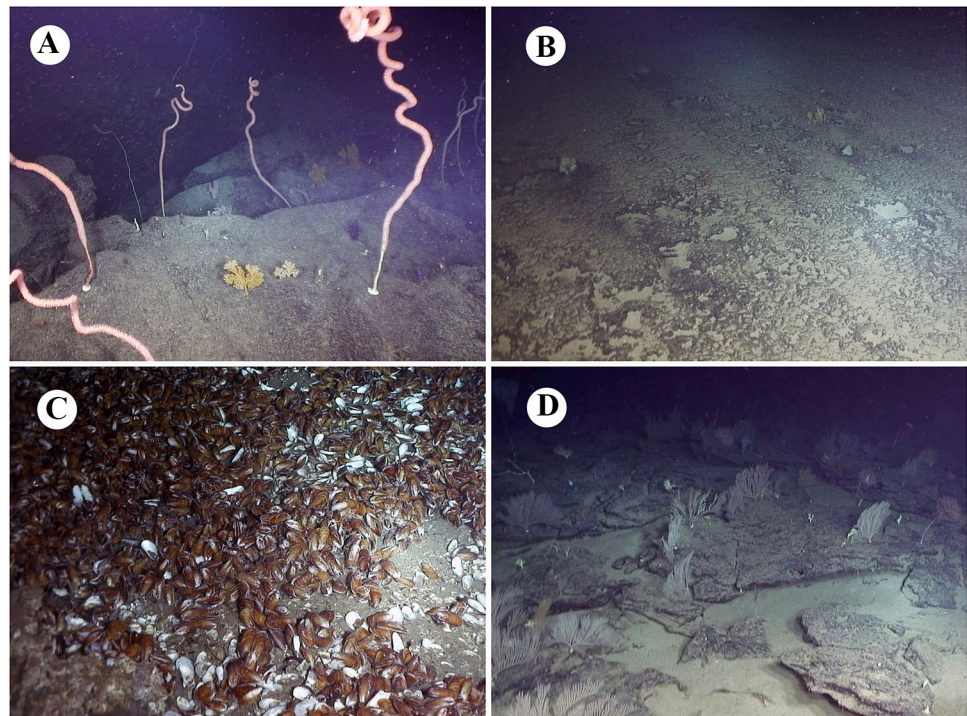


Fig. 2 Habitats investigated during the expedition. **a** Southwestern slope of Ganquan Plateau; **b** northeastern slope of Ganquan Plateau; **c** Haima cold-seep site; **d** mud volcano field



4 Cnidaria; 3 Porifera; 3 Annelida and 1 Echinodermata (Figs. 3, 4, 5 and 6). Some other specimens that are pending identification may include undescribed taxa and therefore are not listed or reported here.

The investigation area of the southwestern slope of Ganquan Plateau has a variety of microhabitats. The substrate basically comprised boulders on steep slopes and flat rock on gentle slopes. In addition, a small cold seep at the foot of the plateau was discovered for the first time. Six macrofaunal species were identified based on the specimens collected from this area, including two species of sponges (Fig. 4a, b), one species of cold-water coral (Fig. 4d), and one gastropod species (Fig. 5g) which was associated with the sponge. *Paraescarpia echinospica*, a symbiotrophic tubeworm, was the only organism observed in the small cold-seep site within this study area.

The investigation area of the northeastern slope of Ganquan Plateau is characterized by flat and rocky slopes extending across a large area. Six macrofaunal species were identified in this habitat, including two sessile organisms (a *Chonelasma* sponge (Fig. 4c) and a *Pseudochrysogorgia* cold-water coral), a squat lobster of the genus *Uroptychus* (Fig. 6g) that was associated with the cold-water coral, and a *Glycera* polychaete associated with an unidentified sponge. A *Cyrtomaia* crab (Fig. 3j) was collected from an unidentified sponge, but nature of their relationship is not clear. One species of *Palaemonella* shrimp was observed in high abundance on the sea bottom.

Five cold-seep sites were surveyed during the expedition. Large areas of mussel beds (dominated by *Gigantidas haimaensis*) were observed at these seeps, which is a typical feature of an active seepage ecosystem (Feng et al. 2018). In contrast, patches of relatively small authigenic carbonate hardgrounds were common around these sites, indicating an early stage of ecological succession in some places (Feng et al. 2018). In total, 24 species were identified from the specimens collected at Haima cold seeps, and most were mollusks (13 species) (Figs. 3e, 5a–f, h, i) and crustaceans (seven species) (Figs. 3f, g, i, 6d–f, h, i). The large vesicomid clam species *Calyplogena marissinica*, which was described first time by Chen et al. (2018) based on samples from the Haima cold seeps, was found and collected again by the corresponding author (Xinzheng Li) in the expedition of 2018. It is a very common species in the cold-seep areas. A red bloody fluid flows from freshly collected specimens of *C. marissinica*. This appearance is very similar to other species of the genus *Calyplogena* (Fujikura et al. 2012). The amphipod *Eurythenes maldoror* (Fig. 6d) and the isopod *Bathynomus jamesi* (Fig. 6e), both of which are opportunistic scavengers, were captured in cage traps. No coral species were observed in this habitat. The symbiotrophic tubeworm *Paraescarpia echinospica*, (Figs. 3d, 4g) occurred in small clusters surrounding authigenic carbonate rocks. The scaled polychaete *Branchipolynoe pettiboneae* (Fig. 4f) and the sea anemone *Actinernus* sp. (Fig. 4e) were very common within and around the cold-seep mussel beds.

Table 1 Checklist of the species currently identified to genus or species levels collected from deep-sea habitats of Haima cold seeps, Ganquan Plateau and a mud volcano field in May of 2018

Phylum	Class	Order	Family	Species	Location and habitat ^b	Figure	Remarks
PORIFERA	Hexactinellida	Lyssacinosida	Rosellidae	<i>Caulophacus</i> sp.	GP1	4a	Sessile living on rocks
			Euretidae	<i>Pleurochorium</i> sp.	GP1	4b	Sessile living on rocks
				<i>Chonelasma</i> sp.	GP2	4c	Sessile living on rocks
CNIDARIA	Anthozoa	Scleractinia	Dendrophylliidae	<i>Enallopsammia rostrata</i> (Pourtales, 1878)	GP1	4d	Sessile living on rocks
			Chrysogorgiidae	<i>Rhodaniridogorgia</i> sp.	MV	3a	Sessile living on rocks
				<i>Pseudochrysogorgia</i> sp.	GP2		Commensal with Euryalid ophiuroid
			Actiniaria	<i>Actinernus</i> sp.	HM	4e	Adherent to dead shells, especially abundant in the rim area of mussel beds
ANNELIDA	Polychaeta	Phyllodocida	Polynoidea	<i>Branchipolynoe pettiboneae</i> Miura & Hashimoto, 1991	HM	4f	Commensal, living within shells of mussels
			Glyceridae	<i>Glycera</i> sp.	GP2		Found in crevice of a Pheronematidae sponge
			Siboglinidae	<i>Paraescarpia echinospica</i> Southward et al., 2002	GP1, HM	3d, 4g	Usually clusters on edges of authigenic carbonate rocks and occasionally forms small assemblages on mussel bed
			Leptochitonidae	<i>Leptochiton tenuidontus</i> Saito & Okutani, 1990	HM	5a	Among mussels and ophiactids in mussel bed of Haima cold seeps
MOLLUSCA	Polyplacophora	Nuculanida	Malletiidae	<i>Malletia</i> sp.	HM	5b	Within mussel bed of cold seeps
			Solemyidae	<i>Solemya</i> sp.	HM	5c	Within mussel bed of cold seeps
			Mytilidae	<i>Gigantidas haimaensis</i> Xu, Feng, Tao & Qiu, 2019	HM	5d	This species is a chemosynthetic-special mussel, and is one of the most dominant species in the mussel beds of Haima cold seeps
			Propeamussiidae	<i>Propeamussium</i> sp.	MV, HM		Identified based on a small colorless specimen associated with dead sponge in mud volcano field. Plenty of <i>Propeamussium</i> scallops were also observed free-living upon mud in rim area of mussel beds
			Venerida	<i>Calyptogena marissinica</i> Chen, Okutani, Liang & Qiu, 2018	HM	5e	Very common in mussel-bed assemblages
Gastropoda	Patellogastropoda ^a	Trochida	Lucinidae	<i>Lucinoma</i> sp.	HM	5f	Within mussel bed of cold seeps
			Pectinodontidae	<i>Bathycyma lactea</i> Zhang et al., 2016	HM		Small in size and predominant in abundance in mussel-bed assemblages
			Calliostomatidae	<i>Tristichotrochus ikukoae</i> (Sakurai, 1994)	GP1	5g	Identified based on a small specimen associated with a Hexasterophora sponge
			Margaritidae	<i>Margarites</i> sp.	HM		Within mussel bed of cold seeps
			Provannidae	<i>Provanna glabra</i> Okutani, Tsuchida & Fujikura, 1992	HM	5h	Within mussel bed of cold seeps
Neogastropoda	Raphitomidae		<i>Phymorhynchus buccinoides</i> Okutani, Fujikura & Sasaki, 1993	HM		Within mussel bed of cold seeps	

Table 1 (continued)

Phylum	Class	Order	Family	Species	Location and habitat ^b	Figure	Remarks
ARTHROPODA	Hexanauplia	Lepadiformes	Buccinidae	<i>Plicifusus</i> sp.	HM	3e	Outside of the mussel bed. The species, based on a large specimen (> 9 cm in length), was observed creeping on muddy seafloor in rim of a mussel bed
			Muricidae	<i>Scabrotrophon scitulus</i> (Dall, 1891)	HM	5i	Within mussel bed of cold seeps
			Poecilasmatidae	<i>Glyptelasma gigas</i> (Annandale, 1916)	MV	6a	Adherent to Isidridae coral
				<i>Poecilasma litum</i> Pilsbry, 1907	MV	6b	Adherent to carapace of <i>Metanephrops neptunus</i> (Bruce, 1965)
	Malacostraca	Amphipoda		<i>Poecilasma obliqua</i> Hoek, 1907	MV	6c	Adherent to the 3rd maxilliped of <i>Metanephrops neptunus</i>
			Eurytheneidae	<i>Eurythenes maldoror</i> d'Udekem d'Acoz & Havermans, 2015	HM	6d	Captured by trapping cage deployed on the seafloor between two seep sites. The cage was put during dive of the day ahead, and collected during the dive of the next day by the corresponding author (Xinzheng Li)
	Isopoda		Cirrolaniidae	<i>Bathynomus jamesi</i> Kou, Chen & Li, 2017	HM	6e	Captured together with <i>Eurythenes maldoror</i> using trapping cage by corresponding author (Xinzheng Li)
			Alvinocarididae	<i>Alvinocaris longirostris</i> Kikuchi & Ohta, 1995	HM	3f, 6f	One of the dominant crustaceans in West Pacific chemosynthetic ecosystems. This species was observed in swarms within seepage mussel beds
	Decapoda		Nematocarcinidae	<i>Nematocarcinus</i> sp.	HM	3g	Identified based on an uncollected individual; inhabiting on muddy seafloor in rim of the mussel bed
			Palaemonidae	<i>Palaemonella</i> sp.	GP2		Observed in high abundance suspending upon the seafloor
			Nephropidae	<i>Metanephrops neptunus</i> (Bruce, 1965)	MV	3h	Common in mud volcano area, inhabiting under stones or in gas holes. Stalked barnacles were recovered from a specimen of <i>Metanephrops neptunus</i>
			Lithodidae	<i>Paralomis</i> sp.	HM	3i	Identified based on a specimen captured using the submersible's mechanic arm; in situ observed crawling in the rim area of mussel bed
			Chirostyliidae	<i>Uropychus setosidigitalis</i> Baba, 1977	GP2	6g	Associated with <i>Paedochrysgorgia</i> corals. The specimen, with the chelipeds lost, was temporarily assigned to <i>U. setosidigitalis</i> , although its rostrum was relatively longer than that of the holotype
			Munidopsidae	<i>Munidopsis laevis</i> Baba & Saint Laurent, 1992	HM	6h	Typical species in chemosynthetic habitats, widely distributed in West-Pacific chemosynthetic environment
			<i>Munidopsis pilosa</i> Henderson, 1885	HM	6i	Widely distributed in Indo-West Pacific. In the deep waters of Taiwan Island, it was found associated with sunken wood (Baba et al. 2009). This is the first discovery of the species from cold-seep environment	
			<i>Munidopsis trifida</i> Henderson, 1885	GP1		Widely distributed across Pacific and Indian Oceans. The specimen collected from Ganquan Plateau has been parasitized by an isopod species in gill chamber	

Table 1 (continued)

Phylum	Class	Order	Family	Species	Location and habitat ^b	Figure	Remarks
ECHINO- DER- MATA		Ophiuroidea	Inachidae	<i>Cyrtomaia</i> sp.	GP2, MV	3j	The specimen in northeastern slope of Ganquan Plateau was captured upon a <i>Chonelasma</i> sponge. Another small specimen was observed inhabiting on a reefs in mud volcano area
			Geryoniidae	<i>Chaceon</i> sp.	MV	3k	Common in chemosynthetic environment, which is seen as an opportunist in this kind of habitat. The species was observed crawling on sea bottom in the mud volcano field
			Ophiothamnidae	<i>Histampica</i> sp.	HM		Inhabiting in mussel bed with other ophiuroids species

^aSubclass^bLocation and habitat: GP1: slope of southwestern Ganquan Plateau, depth 1300–1412 m, with the substrates basically boulders and flat rocks, and the only exception of a cold-seep site where the tubeworm *P. echinospica* were collected, GP2: rocky slope of northeastern Ganquan Plateau, depth 586–910 m, with the substrate flat rocks, HM: within mussels bed of Haima cold-seep sites, depth 1380–1390 m, with the substrate muddy seafloor, MV: mud volcano area, depth 500–810 m, with the substrates flat rocks on the otherwise muddy seafloor

A mud volcano field was newly found during the expedition, and its adjacent seabed was investigated. Here, a large area of flat rocks was present on the otherwise muddy seafloor, providing a hard substrate that harbored abundant corals. Eight species were identified based on faunal collections at this mud volcano field. One cold-water coral, *Rhodaniridogorgia* sp. (Fig. 3a), was identified from video images. Six crustacean species, including three species of stalked barnacle (Fig. 6a–c), were either collected or observed by video. A scallop bivalve of the genus *Propeamussium* was the only mollusk observed in this habitat.

Diversity and distributions of the major macrofaunal groups

Nearly all the sponges observed and collected during the expedition were distributed on rocky slopes of the Ganquan Plateau, whereas no sponges were found at the Haima cold seeps. Similarly, all cold-water corals were collected or observed on the slopes of Ganquan Plateau or at the mud volcano field, where the seafloor was largely composed of boulders or continuous flat reefs, respectively. This is in line with the sessile lifestyle of these two animal groups. Two anemone species were observed in the cold-seep habitat: *Actinernus* sp., which was present in high abundance attached to empty shells, and a species of Hormathiidae, which was found on the seafloor adjacent to one of the seepage site.

Most of the mollusk and polychaete species were found living in the seepage area, where the seabed was mainly mud sediment, providing an ideal habitat for mud-preferring mollusks and polychaetes. High chemosynthetic primary production in cold-seep sites sustains a flourishing seepage community in which chemosynthetic-specialist mussels predominate with high biodiversity and abundance, typically forming large areas of mussel beds. At least eight chemosynthetic-specialist mollusks or polychaetes were collected during the expedition. Only two gastropod species, namely *Tristichotrochus ikukoe* and a species of Cancellariidae, were present on the slope of Ganquan Plateau, associated with a sponge and a cold-water coral, respectively.

Crustaceans generally have a strong ability to move, which allows them to adapt to various deep-sea habitats. A high biodiversity of crustaceans was observed in this expedition. However, few species were shared among the four types of habitat, each of which had a distinct crustacean fauna. The rocky slopes of Ganquan Plateau supported plentiful gorgonians; coral-associated squat lobsters, such as species of *Uroptychus* and *Sternostylus*, were also very common. Additionally, the more complicated topographical environment attracted crabs which could shelter and breed under rocks and in gaps. The rocky substrate in the mud volcano area also provides a habitat for crabs and coral-associated

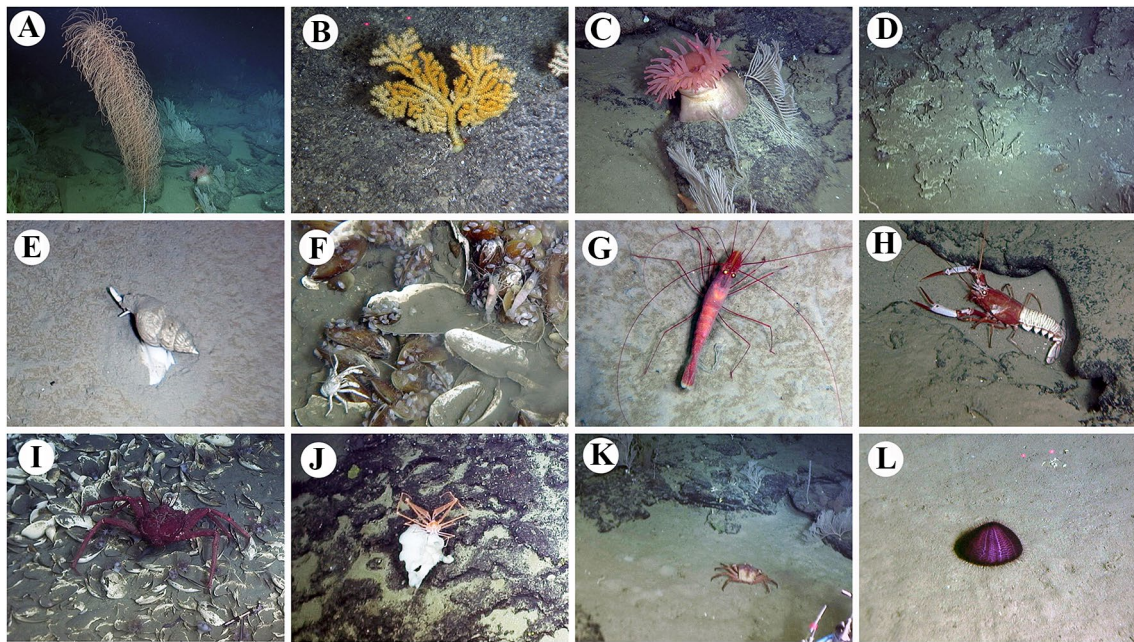


Fig. 3 In situ photographs of epibenthic macrofauna taken during the expedition. **a** *Rhodaniridogorgia* sp., mud volcano field; **b** *Acanthogorgia* gen. et sp. indet., southwestern slope of Ganquan Plateau; **c** *Actinostolidae* gen. et sp. indet., mud volcano field; **d** *Paraescarpia echinospica*, Haima cold-seep site; **e** *Plicifusus* sp., on muddy seafloor around Haima cold-seep site; **f** *Alvinocaris longirostris*) and squat lobster (probably *Munidopsis lauensis*), on mussel bed

in Haima cold-seep site; **g** *Nematocarcinus* sp., on muddy seafloor around Haima cold-seep site; **h** *Metanephrops neptunus*, mud volcano field; **i** *Paralomis* sp., in rim of mussel bed in Haima cold-seep site; **j**, *Cyrtomaia* sp., on sponge *Chonelasma* sp., northeastern slope of Ganquan Plateau; **k** *Chaceon* sp., mud volcano field; **l** *Echinothuriidae* gen. et sp. indet., on muddy seafloor around Haima cold-seep site

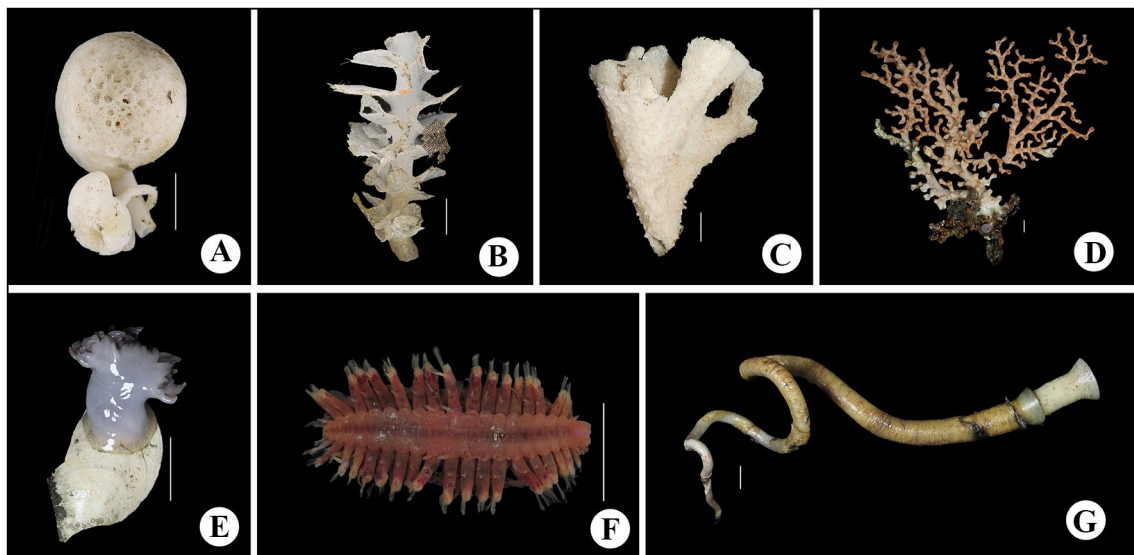


Fig. 4 Colour images of freshly collected specimens. **a** *Caulophacus* sp.; **b** *Pleurochorium* sp.; **c** *Chonelasma* sp.; **d** *Enallopsammia rostrata*; **e** *Actinernus* sp.; **f** *Branchipolynoe pettiboneae*; **g** *Paraescarpia echinospica*. Scale bar = 5 cm (a, c, e); 1 cm (b, d, f, g)

crustaceans; solid rocks on an otherwise muddy seabed provided habitat for hole-dwelling species, such as the crayfish *Metanephrops neptunus*. By contrast, the cold-seep sites were colonized by species of squat lobster and alvinocaridid

shrimp that are specialists in reducing habitats. Examples include *Munidopsis lauensis* and *Alvinocaris longirostris*, which are generally dominant in cold-seep communities.

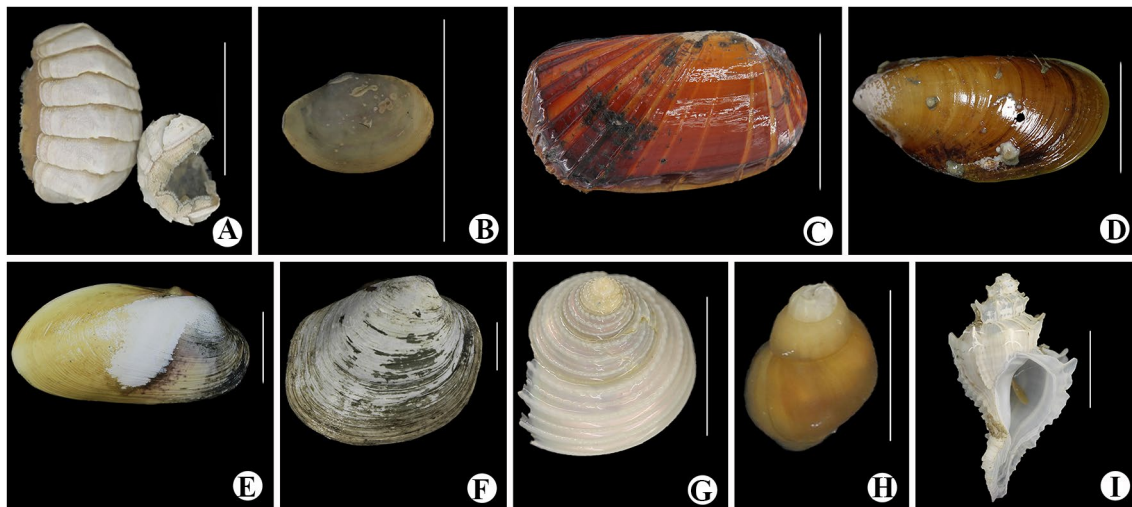


Fig. 5 Colour images of freshly collected specimens. **a** *Leptochiton tenuidontus*; **b** *Malletia* sp.; **c** *Solemya* sp.; **d** *Gigantidas haimaensis*; **e** *Calyptogena marissinica*; **f** *Lucinoma* sp.; **g** *Tristichotrochus iku-koae*; **h** *Provanna glabra*; **i** *Scabrotrophon scitulus*. Scale bar = 1 cm (a, b, f, g, h, i); 5 cm (c, d, e)

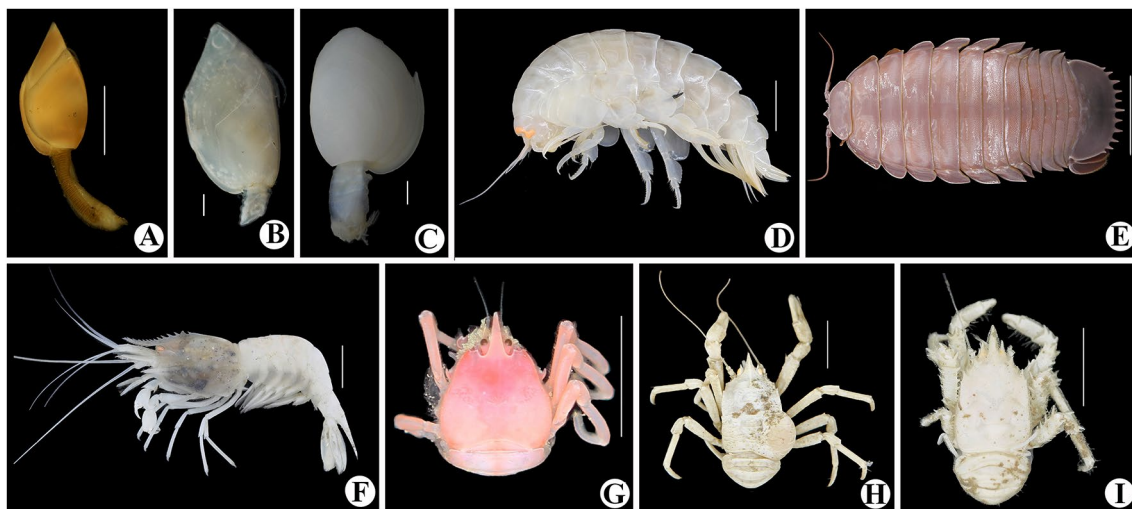


Fig. 6 Colour images of freshly collected specimens. **a** *Glyptelasma gigas*; **b** *Poecilasma litum*; **c** *Poecilasma obliqua*; **d** *Eurythenes maldoror*; **e** *Bathynomus jamesi*; **f** *Alvinocaris longirostris*; **g** *Uroptychus setosidigitalis*; **h** *Munidopsis lauensis*; **i** *Munidopsis pilosa*. Scale bar = 5 mm (a); 1 mm (b, c); 1 cm (d, f, g, h, i); 5 cm (e)

King crabs, acting as vagrant predators, were also very common in the cold-seep communities.

Members of the Echinodermata constituted an important part of the overall faunal biodiversity of the Haima cold-seep areas. However, the taxonomic identity of these animals was difficult to determine and only one species could be identified to genus level. Ophiuroids (brittle stars) were the most diverse subgroup and were observed in high abundance in the cold-seep assemblages. They included species of *Histampica*, Ophiacanthidae, and Amphiuroidae. Conversely, species of Euryalidae were associated with gorgonians

at Ganquan Plateau, and some species of Ophioneuridae, Ophiuridae and Ophiacanthidae were associated with sponges. A species of sea cucumber, probably belonging to the Chiridotidae, was common in and around cold-seep sites. Sea urchins were scarcely observed, but one large-sized individual representing a species of Echinothuriidae (Fig. 31) was captured on the muddy seabed beside a mussel bed.

Preliminary description of community structures and faunal comparison

The community structure of a cold-seep habitat can be influenced by many factors, such as depth, seepage type, and gas composition. Many studies have focused on endobenthic fauna assemblages (micro- and meiobenthos) to quantitatively evaluate their communities in different cold-seep habitats (e.g., Bourque et al. 2017; Cunha et al. 2013; Levin et al. 2015; Washburn et al. 2018).

Although nearly half of the taxa reported here have not been identified to genus or species level, this investigation lends preliminary insights into the community structures of the different types of habitat explored during the expedition. Accordingly, the Ganquan Plateau slopes, characterized by hard substrates, were colonized by high numbers of cold-water corals and sponges along with their associated crustaceans and ophiuroids, revealing faunal community structures similar to those at seamounts. To a large extent, it was the abundance and biodiversity of sessile invertebrates, rather than motile animals, determined the overall epibenthic community structure in these rocky habitats. However, the southwestern slope of Ganquan Plateau, which is deeper and nearer to the Haima cold seeps than the northeastern slope, had a cold-seep microhabitat site; therefore seep-associated faunal species, such as the tubeworm *Paraescarpia echinospica*, were present. The Haima cold-seep faunal assemblages on the muddy seabed revealed typical seepage communities, composed of species that are specialized for life in reducing habitats and opportunistic predators but devoid of cold-water corals and sponges. The mud volcano field, characterized by flat reefs on an otherwise muddy seafloor, shares some community characteristics with the Ganquan Plateau slopes. However, no seep-associated organism was observed in the mud volcano field, probably because the gas release there is less intensive than that in cold-seep sites and thus does not support a chemosynthesis-based assemblage.

The epibenthic community structure of the Haima cold-seep field is distinct from that of Site F in the northeastern South China Sea (southwest to Taiwan Island). Site F is notable for its *Gigantidas platifrons*–*Shinkaia crosnieri* community, in which the mussel *G. platifrons* and the squat lobster *S. crosnieri* thrives in high abundance (Li 2015, 2017). The Haima cold-seep field, which is approximately 1000 km from Site F, was nearly devoid of *S. crosnieri* and has low occurrence of *G. platifrons*, with *G. haimaensis* being the dominant mussel species. Likewise, siboglinid tubeworms were common in the Haima field but scarce at Site F. The squat lobster *Munidopsis lauensis* and the alvinocaridid shrimp *Alvinocaris longirostris* were common at both cold-seep areas. The community structures at the Haima cold-seep field and the Site F are depicted in Fig. 7.

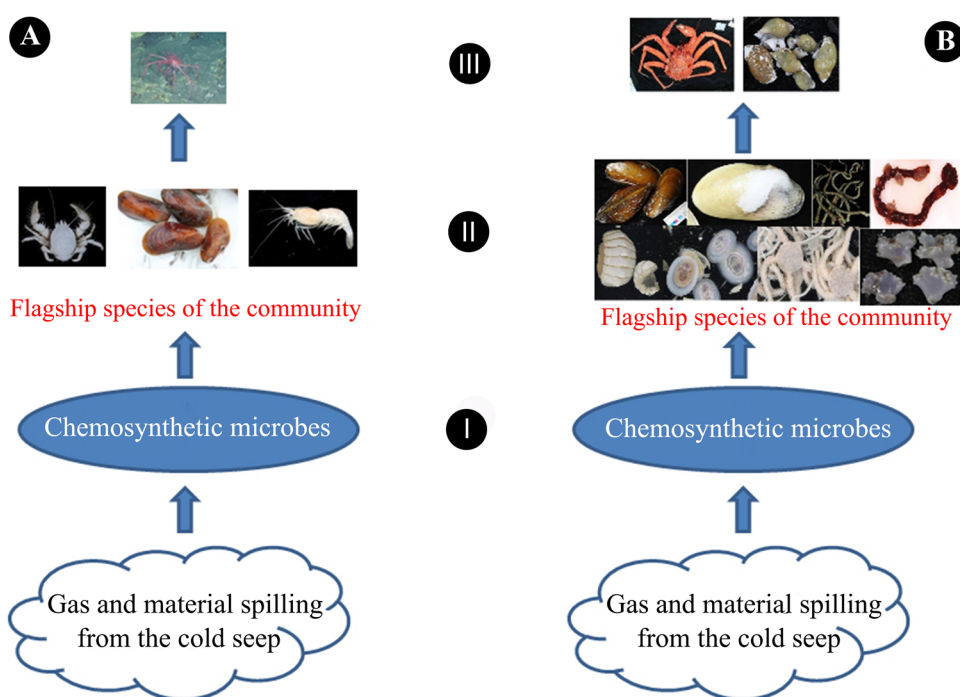
At first approximation, the community structure at Haima cold seeps are roughly similar to that of chemosynthesis-based communities in Sagami Bay (Fujikura et al. 2012), although the presence of the tubeworm *Paraescarpia echinospica* links the Haima cold seeps to chemosynthetic habitats at the Nansei-shoto (Ryukyu) Trench. In contrast, the community structure at Site F is roughly similar to that of the hydrothermal vents in the Okinawa Trough owing to the *Gigantidas platifrons*–*Shinkaia crosnieri* assemblage (Feng et al. 2018; Li 2015, 2017). The cause of such differences between cold-seep communities is still unknown. In accordance with the fieldwork experiences of the corresponding author (Xinzheng Li) at Site F in 2013 and at the Haima cold seeps in 2018, the sulfide smell of the substrate samples from the Haima cold-seep areas was much stronger than that from Site F. This phenomenon may imply that there are differences in the substrate materials, concentrations of reducing compounds, or levels of redox potentials, caused by differences in primary productivities of chemosynthetic microbes. Quantitative environmental data would be needed to verify this.

Our future research will build on this preliminary taxonomic work, especially to (1) better determine community connectivity of epibenthic macroinvertebrates among different seepage sites in the South China Sea, and (2) reveal the factors that govern the community structures and ultimately determine the biodiversity at cold seeps. In conclusion, the cold seeps at both Haima and Site F merit further research, with comparisons based on more sampling and field observations, as well as comprehensive analyses that consider biological, chemical, geographic, and physical oceanographic evidence.

Materials and methods

The investigated areas (Fig. 1) in the South China Sea are located on the northwestern continental slope and included three geomorphological environments: (1) the Haima cold-seep field, with five sites investigated, covering a depth range of 1380–1390 m; (2) a mud volcano field, at depths of 500–810 m; and (3) the slopes of Ganquan Plateau, at depths of 586–910 m on the northeastern slope, and at 1300–1412 m on the southwestern slope. At Haima cold-seep field, two sites were investigated on each dive in the manned submersible, and samples from each site were loaded together in a sampling box. Therefore, the macrofaunal specimens collected from all the five sites were examined and analysed as a single bulk sample to represent the Haima cold-seep habitat. On the southwestern slope of Ganquan Plateau, several microhabitats were present, ranging from rocky cliff to a small muddy cold-seep site in a limited area, where only siboglinid tubeworms were observed, but were

Fig. 7 Schematic diagram of the community structures of the Site F cold-seep site (a) and Haima cold-seep field (b). **a** I, chemosynthetic microbes; II, flagship species, making up the landscape of the community, left, *Shinkaia crosnieri*, middle, *Gigantidas platifrons*, right, *Alvinocaris longirostris*; III, top predator, *Lithodes longispina* Sakai, 1971. **b** I, chemosynthetic microbes; II, flagship species, making up the landscape of the community, above, from left to right, *Gigantidas haimaensis*, *Calyptogena marissinica*, *Paraescarpia echinospica*, indeterminate species of Chiridotidae; below, from left to right, *Leptochiton tenuidontus*, *Bathymacrea lactea*, *Histampica* sp., *Actinernus* sp.; III, top predators, *Paralomis* sp., even ? *Plicifusus* sp.



integrated as part of the overall fauna of the southwestern slope.

Sampling and in situ observations were performed with the manned submersible “*Shenhai Yongshi*”. Videos and photos (Fig. 2) were taken using high-definition underwater cameras deployed on the submersible. During the cruise, only epibenthic macroinvertebrates were qualitatively collected and analyzed. The specimens were visually detected and collected, and were therefore mostly larger than 5 mm in size. Cold-water corals, sponges, mollusks, sea urchins, and large crustaceans were directly grabbed using the mechanical arms of the submersible; small crustaceans like squat lobsters and shrimps were sampled using nets manipulated by the mechanical arms. Some adherent organisms, such as small gastropods and ophiurids, were collected together with rocks and corals. Specimens were photographed immediately after being transported to the deck of the mother vessel, and then directly frozen for conservation. Most of the specimens were retained by the research for taxonomic examination; however, specimens deposited in Tongji University and uncollected organisms observed in situ were identified only from videos and photographs.

Taxonomic studies of the specimens were mainly based on morphological method. DNA barcoding was employed to confirm the identification of some species. The samples are deposited in the Marine Biological Museum of Chinese Academy of Sciences in Qingdao, China.

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Author contributions DD wrote most part of the manuscript and prepared Figs. 1, 2, 3, 4, 5 and 6; XL, corresponding author, provided the outline for this manuscript, wrote part of the manuscript, prepared Fig. 7 and Table 1, improved the text; other co-authors, MY, LG, YL, JS, ZG, QK, NX, JZ, as well as DD and XL, identified parts of samples, respectively, and provided some literature.

Compliance with ethical standards

Conflict of interest The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Animal and human rights statement No animal and human rights are involved in this article.

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