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Macrobenthic functional group analysis of ecological health of the intertidal artificial oyster reefs in the Yangtze Estuary, China

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The functional groups and the ecological health of a constructed oyster reef were investigated during April 2016 to October 2017 in the Yangtze estuary. The data of the long-term monitoring program (14 years) confirmed the feasibility of creating an artificial oyster reef in Yangtze estuary. One or two functional groups were absent from moderate salinity conditions (5.17% - 8.73%), while macrobenthos functional group species richness, abundance, and biomass were greatest under high salinity conditions. The feeding evenness index (j_{FD}) showed that 14 years after construction, the ecosystem health of the artificial oyster reefs were good or high, and was better under high salinity conditions. Ecological health was lower under moderate levels of salinity, possibly due to the absence of some functional groups. Redundancy analysis indicated ecosystem health was associated with changes in salinity and substrate factors. Our study confirmed artificial oyster reef construction is feasible and has a positive effect on estuarine ecosystem health.

KEYWORDS

functional group, macrobenthos, Yangtze estuary, oyster reef, ecological restoration

1 Introduction

Estuaries are highly productive, yet threatened ecological systems that deliver a range of ecosystem services and benefits to human society. However, the health of many estuaries is under threat from high levels of anthropogenic activities and pollution associated with the dense human populations that occur in these regions (Adger et al., 2005).

For example, sediment and organic material deposited by rivers in estuaries include nutrients, heavy metals, microplastics, and other contaminants (Chen et al., 2012; Han et al., 2017; Xu et al., 2018) that have led to biodiversity loss and reduced ecological resilience in estuarine environments (Lotze et al., 2006; Liu et al., 2018a).

The recognition of the essential role of estuaries for humans and other organisms has led to the initiation of ecological restoration programs (Lotze et al., 2006), where rehabilitation or construction oyster reefs represent an effective estuarine restoration technique. The biological structure formed by a large number of oysters is called an oyster reef, which is an important coastal habitat, because they provide many ecological services and bring multiple ecological benefits (Grabowski et al., 2022), such as water purification through water filtration and the removal of phytoplankton and heavy metals, and habitat provisioning through creation of three-dimensional habitats for macrobenthos and fish (Breitburg et al., 2000; Coen and Luckenbach, 2000; Grabowski and Peterson, 2007; Thomsen et al., 2007; La Peyre et al., 2014; Liu et al., 2018b). Thus, construction of oyster reefs has been increasingly used for shoreline protection and ecological restoration (La Peyre et al., 2014; Chakraborty, 2016; Walles et al., 2016).

The Yangtze Estuary is the largest in Asia and is one of the most globally important ecotones (Chen et al., 2012) that provides vital ecological services, including breeding grounds for aquatic organisms and stopover and overwintering sites for millions of migratory birds (Xu and Zhao, 2005). However, it is affected by rising sea-level and anthropogenic pressures that have culminated over centuries of overexploitation, reclamation, and pollution (He et al., 2011; Kirwan and Megonigal, 2013; Liu et al., 2016b; Yu et al., 2018); consequently, eutrophication and reduction in biodiversity have led to a deterioration in the ecological health of the Yangtze Estuary (Chai et al., 2006; Chen et al., 2012; Huang et al., 2021; Huang et al., 2022). In response, an oyster reef restoration project was initiated in the region (Quan et al., 2009), based on previous studies that showed the construction of artificial oyster reefs in the Yangtze River estuary was feasible and effective in the restoration of the ecosystem (Quan et al., 2009; Quan et al., 2012; Lv et al., 2016a; Liu et al., 2018b).

It is likely that artificial and restored oyster reefs may be affected by human activities and pollution, such as over-fishing of the nursery beds and oil contamination, therefore, monitoring of such restoration schemes is crucial for their success. Macrobenthos represents a vital component of oyster reefs, and are ideal indicators for use in ecosystem health monitoring schemes due to their limited mobility, ease of collection, and high sensitivity (Borja et al., 2000; Borja et al., 2003; Gamito and Furtado, 2009; Liu et al., 2019); however, studies of oyster reefs have tended to focus on taxonomic macrobenthos community structure, abundance, biomass, diversity, and ecosystem health (Quan et al., 2009; Quan et al., 2012; Lv et al., 2016a; Liu et al., 2018b) that may be more appropriate in the evaluation of biodiversity rather than ecosystem health (Cummins et al., 2005; Peng et al., 2013). Macrobenthos functional groups represent taxa that have similar ecological characteristics, so assessment of their responses to environmental change is more meaningful than that of individuals or taxonomic families, because it provides a more accurate indication of ecosystem-level responses to ecological perturbations (Yang et al., 2017).

Ecological restoration may trigger the emergence of novel functional groups and turnover of existing functional groups (Lv et al., 2018). Previous studies of the Yangtze Estuary oyster reef have quantified biological resources, including macrobenthos and nektons, macrobenthic taxonomic diversity and ecological health (Quan et al., 2007; Quan et al., 2009; Quan et al., 2012; Lv et al., 2016a; Liu et al., 2018b); however, assessment of macrobenthos functional groups remains lacking. Here, we identified macrobenthos functional groups to assess the ecological status of the artificial oyster reef in the Yangtze estuary. Specifically, the objectives were to (1) assess the composition and abundance of macrobenthos functional groups; (2) measure long-term variation in macrobenthos functional group composition; (3) test the relationship between macrobenthos functional groups and environmental variables; and, (4) use macrobenthos functional groups to assess ecosystem health.

2 Materials and methods

2.1 Study site

The Yangtze estuary (29°30'N-32°00'N, 121°00'E-124°00' E) artificial intertidal oyster reef (c. 260 ha) was constructed in 2004 in the north passage of the southern Yangtze River channel (Figures 1A, B), and comprises two dikes (south and north) with 19 groins (S1-S9 in the south dike and N1-N10 in the north dike). The north dike (total length: 48 km) is located along the southern margin of Hengsha Island, and the south dike (total length: 49.2 km) is located along the northern edge of Jiuduansha National Nature Reserve. The artificial reef is vaulted in cross-section (Figure 1C).

2.2 Sampling

Macrobenthos were sampled in April and September 2016, and in August and October 2017 from three paired sites (N = 6) at the two dikes along an increasing salinity gradient, where salinity at sites S2 and N2 was 0.43 ± 0.03 and 0.43 ± 0.04 , respectively, S5 and N6 was 6.55 ± 1.38 and 7.86 ± 0.87 , respectively, and S8 and N9 was 16.43 ± 0.34 and 15.50 ± 1.09 , respectively. At low spring tide from three site (high, mid, and low tidal zones), four quantitative samples of macrobenthos



Study site locations. (A) Yangtze River estuary, China; (B) Sampling sites on the artificial intertidal oyster reef; (C) Schematic drawing of the sampling transect and cross-section of the artificial reef; (D) No organisms were recorded from the low-salinity sites (S2 and N2).

(25 × 25 cm) were collected from each site (Quan et al., 2009), where the high tidal zone was located on the crest of the reef, the mid zone was on the flank, and the low zone was at the reef base. In addition, as many qualitative samples as possible were collected from an area of 25.0 m² from the sampling center. Samples were immediately sieved (1.0-mm mesh) and the collected macrobenthos were preserved in 75% alcohol prior to subsequent identification at the laboratory to the lowest possible taxonomic level; individuals were counted and weighed to the nearest 0.01g wet weight. No organisms were recorded from both quantitative and qualitative investigations from the low salinity sites (S2 and N2) (Figure 1D).

Water temperature (WT), salinity (SAL), pH, and dissolved oxygen (DO) at the sampling sites were measured using a HQ 40D multi-parameter water quality meter (Hach, Loveland, Colorado, USA). The abundance of oyster (OA) and barnacles (BA) were defined as the substrate variables of the oyster reef.

2.3 Long-term variation in functional groups

Including the 2016 and 2017 data collected, changes over a 14year period in the macrobenthic functional groups were integrated in an analysis of dynamics from 2004 to 2017. Historical data points (2004–2014) were obtained from the literature, namely for September 2004, August 2005, August and November 2007, April and July 2008, May and September 2009, April and November 2011, July and October 2012, August and October 2013, August and October 2014 (Quan et al., 2009; Quan et al., 2012; Lv et al., 2016a).

2.4 Macrobenthos functional groups and trophic levels

We defined five macrobenthos functional groups according to feeding strategy (Zhu and Lu, 2003; Peng et al., 2013; Yang et al., 2017) that comprised planktivore (Pl) herbivore (He); predator (Pr); omnivore (O); and, detritivore (D). From these, we defined four trophic levels (Figure 2), where the first level comprised primary producers (we did not survey this group), the second level comprised primary consumers (Pl, He, and D), the third level comprised omnivore secondary consumers of levels I and II, while the fourth level comprised predatory (Pr) secondary consumers of levels II and III.

2.5 Statistical analysis

We used CLUSTER (hierarchical clustering) and NMDS (two-dimensional nonmetric multidimensional scaling) in PRIMER v5.0 to test the degree of similarity of the macrobenthic functional groups (Clarke and Gorley, 2001). Macrobenthos functional group data were fourth-root transformed to calculate the community similarity according to the Bray-Curtis similarity index, and similarity analysis (ANOSIM 2-way) was used to test for tidal zone and site differences in the abundance of macrobenthos functional groups.

Detrended correspondence analysis determined linear redundancy analysis (RDA) was more appropriate than a unimodal analysis (maximum gradient length of the axes<3) to describe the effects of environmental variables on macrobenthos functional group abundance. With the exception of pH, environmental data were log10 (x+1) transformed prior to analysis.

A feeding evenness index (j_{FD}) was used to assess the ecological health status of the macrobenthos trophic levels, following the methodology reported by (Gamito and Furtado, 2009):

$$j_{FD} = H'_{FD} / Log_2 n \tag{1}$$

where *H*' is the Shannon–Wiener index and *n* is the number of trophic groups. Ecosystem health was determined as low ($j_{\rm FD} = <0.20$), poor ($j_{\rm FD} = 0.20-0.40$), moderate ($j_{\rm FD} = 0.40-0.60$), good ($j_{\rm FD} = 0.60-0.80$), and high ($j_{\rm FD} = >0.80$) (Gamito et al., 2012).

Two-way ANOVA, with *post-hoc* Turkey test, was used to test the effects of site and tidal zone on macrobenthos species richness and abundance, and biomass of macrobenthos functional group at P < 0.05. When assumptions of homogeneity of variances were not met, data were fourth-root transformed; if variances remained nonhomogenous, we applied the Scheirer-Ray-Hare extension of the Kruskal-Wallis test. Associations between environmental variables and macrobenthos were tested using correlation analysis at P < 0.05. Analyses were performed using SPSS statistics 19.0 and presented data are means \pm SEM.

3 Results

3.1 Temporal variation in functional group structure

The species composition of the macrobenthos functional groups varied between 2004 and 2017, where total number of species increased immediately after reef creation, before it declined in 2008, and then peaked in 2011 (36 species); species richness remained stable from 2012 to 2017 (Figure 3). The number of



functional groups increased with restoration time, from three in 2004 (O, He, and Pl) to five from 2009 onwards (Figure 3).

3.2 Study site and tidal zone variations in macrobenthos community

We identified three and four macrobenthos functional groups at the moderate salinity sites N6 and S5, respectively (Figure 4 and Table 1), and five groups at both high salinity sites (S8 and N9) during 2016-2017. O and He groups comprised higher numbers of species at the high salinity sites (S8 and N9) than at the moderatesalinity (S5 and N6) sites (P< 0.05, Table 2). There were more species in the O group in the low intertidal zone, but fewer in the mid zone; species richness varied among the three tidal zones (P < 0.05, Table 2).

3.3 Functional group abundance and biomass

Abundance of C, O, and He and biomass of C were higher in high-salinity (N9 and S8) sites than in the moderate salinity sites (N5 and S6) (P<0.05, Tables 3-6), and there was no effect of tidal zone on abundance and biomass of the five functional groups (Tables 5, 6).

3.4 Functional group community structure

Cluster analysis revealed similarities and differences in macrobenthos functional groups among sites and tidal zones, where there were three clusters comprising S8-L, S8-M, N9-L, S5-M, S8-H, N9-H, and N9-M; S5-H; and, N6-H, N6-M, and N6-L (Figure 5A). The nMDS analysis results were consistent with the cluster analysis, with a stress value = 0.08 indicating the results were reliable (Figure 5B). There were site differences in macrobenthos functional groups (ANOSIM 2 Global R = 0.719; P = 0.029; however, there were no tidal zone differences (Global R = 0.167; P = 0.34).

3.5 Ecological health

We found that j_{FD} values for the macrobenthos functional groups were<0.8 in the moderate salinity sites (S5 and N6), indicating ecological health was good, and >0.8 in the high salinity sites (S8 and N9), indicating high quality ecological health (Figure 6).

3.6 Relationship between macrobenthic functional group abundance and environmental variables

Monte Carlo tests for the first and all Redundancy analysis (RDA) canonical axes were significant (P = 0.004), indicating these parameters may be important in explaining the composition of the macrobenthos functional groups. The first axis explained 62.4% of total variation in the functional group data and all canonical axes accounted for 72.3% of the variance in the functional group abundance-environment relationship. Functional groups were correlated with environmental variables: groups O, Pl, D, and He were positively correlated with barnacle



Pl, planktivore



	Sites	Tidal zone	Functionalgroup	Reference
Coelenterata				
Haliplanella sp.	S5S8N6N9	HML	0	(Cai, 2015)
Sipuncula				
Phascolosoma esculenta	S8N9	HML	D	(Cai, 2015)
Annelida		·	·	
Perinereis aibuhitensis	S5S8N6N9	HML	0	(Zheng and Fan, 1986)
Neanthes japonica	S5S8N6N9	HML	0	(Ye, 2010)
Dentinephtys glabra	S8N6N9	HML	Pr	
Perinnereis nuntia	S8	L	0	
Hydroides ezoensis	S8N9	Н	PI	
Mollusca		!	1	
Nerita yoldi	S5S8N6N9	HML	0	
Littorina brevicula	S8N9	HML	Не	(Bao et al., 2008)
Littorinopsis intermedia	S8N6N9	HML	Ph	
Pyrene bella	S8N6N9	HML	Не	
Purpura clavigera	S8N9	HML	Pr	
Rapana bezoar	S8N9	HML	Pr	
Potamocorbula amurensis	\$5N9	HML	PI	(Lv et al., 2013)
Musculus senhousia	S8N9	HML	PI	(Liu et al., 2016a)
Modiolus flavidus	S8N9	HL	PI	(Li et al., 2014)
Vignadula atrata	S8N9	HML	Pl	
			•	(Continued)

	Sites	Tidal zone	Functionalgroup	Reference
Barbatia bistrigata	S8N9	HML	Pl	
Unkonwn species (GASTROPODA)	N6	L	Не	
Arthropoda		1	·	·
Corophium acherusicum	S5N6	L	Не	(Chen, 2001)
Metopograpsus quadridentatus	S5S8N6N9	HML	Не	(Zhang et al., 2017)
Macromedaeus distinguendus	S5S8N9	HML	Pr	
Eriocheir leptognathus	\$5\$8	HM	Pr	
Scylla serrata	N9	М	Pr	(Yuan et al., 2002)
Pinnotheres sinensis	N6	Н	0	
Alpheus japonicus	S8	Н	Pr	(Zhang et al., 2017)
Gnorimosphaeroma rayi	S5S8N6N9	HML	Pr	
Cleantioides annaandalei	S8	HML	Не	
Eohaustorius cheliferus	S5S8N6N9	HML	0	(Zhu and Lu, 2003)
Chordata		1	·	·
Liciogobius guttatus	S5S8N9	ML	Pl	(Li et al., 2014)
Echinodermata				
Amphiura vadicola	S8	L	D	(Li et al., 2013)
No organisms were recorded from the low-salinity site	es (S2 and N2) Pr, predator; I	D, detritivore; O, omnivore; He, I	herbivore; Pl, planktivore. H, high; M, mi	ddle; and L, Low tidal zone.

TABLE 1 Continued

abundance, oyster abundance, salinity, and temperature, and negatively correlated with pH, whereas group C was negatively correlated with dissolved oxygen (Figure 7).

Discussion

Restored and artificial oyster reefs play a vital role in improving the ecological health status of estuarine environments, because they improve water quality and provide a three-dimensional structure that may be colonized by macrobenthos and fish (Breitburg et al., 2000; Thomsen et al., 2007). In the first year (2004) since its creation, the artificial oyster reef in the Yangtze estuary was found to have become host to three functional groups (comprising five species) of macrobenthos; the species richness of the functional groups rapidly increased with time, with the exception of 2008, before it peaked in 2011 and remained stable from 2012. This increase in functional group species richness occurred after oyster abundance had stabilized (2011) (Liu et al., 2018b) and prior to the increase in macrobenthos taxonomic diversity that occurred from 2012 to 2016-2017 (Liu et al., 2018b). These results support previous studies that have indicated the oyster reefs in the Yangtze Estuary be develop into self-sustaining systems (Walles et al., 2016; Liu et al., 2018b).

Water salinity is a key factor that affects the distribution of macrobenthos in estuarine habitats (Holland et al., 1987; Liu

TABLE 2 Study site and tidal zone effects on macrobenthic functional group species richness during 2016-2017.

Functional group		Sites	Tidal zone				
	df	F or H	Р	df	F or H	Р	
С	3	5.870	0.12	2	0.168	0.92	
D	3	0.609	0.89	2	0.429	0.81	
0	3	10.329	< 0.001	2	4.212	0.023	
Ph	3	16.546	< 0.001	2	0.873	0.427	
Pl	3	6.045	0.11	2	0.472	0.79	

										N9		
Pr	346.0 ± 140.2	249.0 ± 116.2	_	501.0 ± 165.2	263.0 ± 103.2	348.0 ± 89.9	40.0 ± 26.8	164.5 ± 126.8	168.0 ± 73.8	652.0 ± 146.3	491.0 ± 126.9	247.0 ± 70
D	0.0 ± 0.0	0.0 ± 0.0	_	0.0 ± 0.0	1.0 ± 1.0	3.0 ± 1.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	1.0 ± 1.0	0.0 ± 0.0	3.0 ± 3.0
0	83.0 ± 36.5	180.0 ± 66.4	_	342.0 ± 15.8	171.0 ± 57.3	180.0 ± 21.3	7.0 ± 5.7	209.0 ± 70.8	368.0 ± 73.2	355.0 ± 56.4	258.0 ± 88.9	329.0 ± 50.8
He	3.0 ± 3.0	12.0 ± 7.1	_	166.0 ± 29.5	53.0 ± 24.6	282.0 ± 59.8	11.0 ± 7.5	13.0 ± 5.5	19.0 ± 10.4	75.0 ± 32.3	57.0 ± 26.0	74.0 ± 33.1
Pl	1.0 ± 1.0	15.0 ± 5.0	_	122.0 ± 6.2	89.0 ± 36.1	132.0 ± 14.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	61.0 ± 36.1	42.0 ± 24.7	76.0 ± 44.2
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TABLE 3 Abundance of macrobenthos functional groups in the constructed intertidal oyster reef in Yangtze River Estuary during 2016-2017.

H, high tidal zones; M, mid tidal zones; L, low tidal zones.

TABLE 4 Biomass of macrobenthic functional groups in the constructed intertidal oyster reef in Yangtze River Estuary during 2016-2017.

									N9			
Pr	2.1 ± 1.1	3.8 ± 1.2	-	28.7 ± 15.6	22.6 ± 15.3	89.3 ± 45.2	25.3 ± 20.5	17.4 ± 13.4	1.5 ± 1.5	8.2 ± 4.0	6.1 ± 3.4	8 ± 5.7
D	0.0 ± 0.0	0.0 ± 0.0	-	0.0 ± 0.0	0.0 ± 0.0	0.1 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0	0.0 ± 0.0
0	9.1 ± 5.2	23.5 ± 13.1	-	38.3 ± 15.3	42.5 ± 18.9	19.3 ± 5.4	11.9 ± 5.6	53.3 ± 29.6	54.2 ± 33.6	115.9 ± 16.2	75.5 ± 6.4	72.5 ± 5.0
He	0.2 ± 0.2	1.3 ± 1.0	-	8.6 ± 4.5	8.2 ± 3.9	16.1 ± 9.5	13.6 ± 7.9	7.4 ± 4.0	0.3 ± 0.3	2.1 ± 1.0	0.9 ± 0.3	2.1 ± 1.0
Pl	0.1 ± 0.1	1.9 ± 0.7	-	6.1 ± 3.2	8.9 ± 3.3	21.8 ± 6.2	11.3 ± 8.0	8.9 ± 5.3	0.0 ± 0.0	1.1 ± 0.8	0.8 ± 0.5	1.1 ± 0.7
H bio	h tidal zonas	M mid tidal zo	noc I I	ow tidal zonas								

H, high tidal zones; M, mid tidal zones; L, low tidal zones.

et al., 2018a), where species richness, abundance, and biomass increase with increase in salinity (Yuan et al., 2002). In this study, the increase in salinity (from S5 and N6 to S8 and N9) resulted in an increase in species richness, abundance, and biomass of macrobenthic functional group (P< 0.05; Figure 7), indicating salinity affects the distribution of macrobenthos functional groups in the artificial Yangtze Estuary oyster reefs. Adaptation to salinity varies among species (Yuan et al., 2002); for example, *Phascolosoma esculenta*, which was only collected

from the high salinity sites (S8 and N9), belongs to the D functional group, and has a survival salinity of 5-30 ‰, with a salinity range for embryo development of 17.10-30.15 ‰ (Jin et al., 2011). We found that functional group D, whose main species comprise polychaetes and Sipuncula that are sensitive to fluctuations in environmental conditions (Dauvin and Ruellet, 2007; Cai, 2015), was absent from the moderate salinity study sites (S5 and N6). There are marked seasonal changes in natural environmental variables, such as salinity and dissolved oxygen,

TABLE 5 Effect of study site and tidal zone on macrobenthos functional group abundance in the constructed intertidal oyster reef in Yangtze River Estuary during 2016-2017.

		Sites		Tidal zone		
	df	F or H	Р	df	F or H	Р
Pr	3	4.814	0.007	2	1.388	0.264
D	3	0.582	0.9	2	0.449	0.8
0	3	3.817	0.019	2	1.554	0.227
Не	3	12.491	<0.001	2	1.278	0.292
Pl	3	7.221	0.07	2	0.0989	0.95

		Sites		Tidal zone		
	df	F or H	Р	df	F or H	Р
Pr	3	2.998	0.045	2	0.016	0.984
D	3	0.42	0.94	2	0.044	0.98
0	3	4.304	0.23	2	0.124	0.94
He	3	2.565	0.46	2	0.068	0.97
Pl	3	2.567	0.46	2	0.157	0.92

TABLE 6 Effect of study site and tidal zone on macrobenthos functional group biomass in the constructed intertidal oyster reef in Yangtze River Estuary during 2016-2017.

and tidal conditions in the area where study sites S5 and N6 were located: it is possible that mud trapped at the bottom of the oyster reef is removed by tidal action, resulting in a lack of food source required by functional group D.

Salinity not only influences estuarine macrobenthos distribution patterns, but also affects growth and survival of oysters (Hulathduwa and Brown, 2006; Liu et al., 2018b). Effects of salinity on oyster size and abundance vary with region, therefore oyster substrate of the artificial reefs is likely to similarly vary with region, and may lead to diverse distribution patterns of macrobenthos functional groups. Our RDA confirmed that salinity, barnacle abundance, and oyster abundance (substrate parameters) were the main factors that influenced the distribution of macrobenthos functional groups in the Yangtze River estuary artificial oyster reefs (Figure 7). Although there were differences in growth and abundance of oysters between regions and at different tidal zones within a region, there was no difference in macrobenthos functional group species richness, abundance or biomass, with the exception of species richness of group D among

the tidal zones. These results indicate that while there are differences in substrate habitat with salinity and among tidal zones within a salinity range, the distribution pattern of macrobenthos functional groups varies with salinity, but not with different tidal zone, such that tidal zone and associated effects on substrate elicit little effect on the artificial oyster reef macrobenthic functional groups in the Yangtze Estuary. This finding is inconsistent with those of other local natural habitats, such as Jiuduansha and Quanzhou Bay (Zhu and Lu, 2003; Zhuo et al., 2014), possibly because the habitat of artificially constructed oyster reef lacks diversity. We suggest enrichment and diversification of artificial oyster reef habitats may improve the recruitment of functional group types to increase biodiversity and ecosystem stability.

In general, healthy environments contain most types of feeding guild, whereas degraded environments tend to be dominated by a few trophic groups, with others absent (Gamito and Furtado, 2009; Peng et al., 2013). In this study, we found one or two groups were absent from the moderate





salinity study sites (S5 and N6), where they included functional groups that play an important role in estuarine food web nutrient cycling and energy flow. For example, Pl is the main source of nutrition for predators and omnivores, and plays an important role in controlling the growth of plankton, such as cyanobacteria (Figure 2). The absence of this functional group from the moderate salinity study sites indicates compromised ecological health, as supported by our results for $j_{\rm FD}$, and previous results for AMBI, M-AMBI, and ABC curve (Liu

et al., 2018b). Our ANOSIM 2 analysis showed site differences in ecological health, where it was better in the high salinity study sites. These results indicate that 14 years after construction, the habitat status of the artificial oyster reefs in the Yangtze Estuary was good or high, and was better under high salinity conditions (S8 and N9) than under moderate salinity (S5 and N6). Longterm artificial oyster reef monitoring confirms the feasibility of artificial oyster reef construction in the Yangtze estuary, and can be applied to estuarine areas or tidal flats within the same salinity



range, such as east shoal of Nanhui, which are mainly affected by reclamation (Liu et al., 2018a). Coupled with the complex habitat of natural intertidal zone, it is inevitably more conducive to the ecological restoration of artificial oyster reef in these areas.

We found that macrobenthos functional group structure was more heterogeneous under high salinity conditions (S8 and N9) than under moderate levels of salinity (S5 and N6). Substrate parameters and environmental variables, such as dissolved oxygen, pH, temperature, and salinity, were similar between S5 and N6, and S8 and N9. However, functional group species richness, abundance, and biomass, and $j_{\rm FD}$ of the southern dike (S5 and S8) was higher than in the north dike (N6 and N9), possibly due to site conditions. The southern region of the oyster reef in the Yangtze Estuary is proximate to the ecologically protected Jiuduansha Nature Reserve (Zhang et al., 2018) that may affect the structure and ecological environment of macrobenthos (Ly et al., 2016b). In contrast, the lack of such protection measures in the north dike may render the oyster and macrobenthos in this area susceptible to human activities, such as overfishing. Therefore, we suggest the establishment of a nature reserve to protect this important oyster reef habitat in the Yangtze estuary.

In summary, artificial oyster reef was feasibility and has positive effect on the ecological health in the Yangtze estuary. Further, the ecological health was worse in medium salinity than high salinity, may due to the absence of functional groups. The release of species of absent functional group can be taken in the medium salinity region. The artificial oyster reef may be disturbed by human activities, thus a nature reserve should be established to protect this important habitat in the Yangtze estuary.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Author contributions

YNZ: Conceptualization, Methodology, Resources, Formal analysis, Investigation, Data curation, Visualization, Writing original draft, Writing - review and editing. HW: Conceptualization, Methodology, Resources, Visualization, Data curation, Writing - review and editing. YLZ: Conceptualization, Methodology, Resources, Visualization, Funding acquisition, Writing - review and editing. JD: Writing - original draft, Writing - review and editing. ZZ: Investigation, Data curation, Writing - review and editing. HZ: Conceptualization, Methodology, Writing - review and editing. ZL: Conceptualization, Methodology, Resources, Visualization, Supervision, Writing - review and editing, Project administration, Funding acquisition. All authors contributed to the article and approved the submitted version.

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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.

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