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Benthic ostracod diversity and biogeography in an urbanized seascape

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ABSTRACT

Hong Kong is one of the most urbanized coastal cities in the world. Yet, despite extensive anthropogenic impacts, adjacent marine environments harbour tremendous biodiversity. We investigated how the diversity, taxonomic composition, and biogeography of meiobenthic ostracods in Hong Kong's coastal waters vary in response to natural and anthropogenic factors. Our regression models indicated that metal pollution and mud content were the main factors affecting meiofaunal diversity, with eutrophication also playing a role. The highest diversity was observed in the Victoria Harbour region at the center of Hong Kong's urbanized seascape, and the lowest diversities were observed in Mirs Bay, Port Shelter, and Tolo Harbour. Ostracod diversity and biogeography patterns are congruent with published studies of other soft-sediment fauna, which also identified a diversity peak in Hong Kong's urban center and a vast southern water biofacies characterized by muddy and turbid conditions. These results do not apply to organisms that prefer oligotrophic conditions, such as hard corals. For those taxa, eutrophic waters in western Hong Kong are generally not habitable and higher diversities are observed in less productive, eastern waters. Our findings indicate that meiofaunal ostracods can be used more broadly as a bioindicator for the diversity of soft sediment benthos.

1. Introduction

Benthic communities in highly urbanized settings are typically characterized by low diversity, low abundance, and by the dominance of a few taxa that can tolerate eutrophic conditions (see Frenzel and Boomer, 2005; Ruiz et al., 2005; Yasuhara et al., 2012a). Examples of this are seen around the world, including in the Chesapeake Bay near Washington DC (Karlson et al., 2000; Cronin and Vann, 2003; Willard and Cronin, 2007), Osaka Bay next to Osaka (Yasuhara et al., 2002; Yasuhara and Yamazaki, 2005; Tsujimoto et al., 2006; Yasuhara et al., 2007; Tsujimoto et al., 2008), and the Baltic Sea adjacent to many European large cities (Andrén, 1999; Ruiz et al., 2005). However, relatively few studies have examined benthic diversity in areas of South East

Asia that are adjacent to megacities like Hong Kong, Singapore, Manila, and Jakarta, among others (Fauzielly et al., 2013; Cybulski et al., 2020; Tan et al., 2021).

Hong Kong, with a land area of 1100 km² and a population of over seven million, is one of the most urbanized coastal cities in the world (Morton, 1989; Lai et al., 2016) and also one of the world's busiest container ports (Tanner et al., 2000). Victoria Harbour is located at the center of the City (Xu et al., 2011) (Fig. 1). The impacts of human activities on Victoria Harbour are significant: organic and heavy metal pollution and extensive land reclamation (Wong et al., 1995; Blackmore, 1998; Yin and Harrison, 2007; Tang et al., 2008; Liu et al., 2011). To respond to the challenges of rapid urbanization and resulting environmental degradation in many areas of Hong Kong, the marine water and

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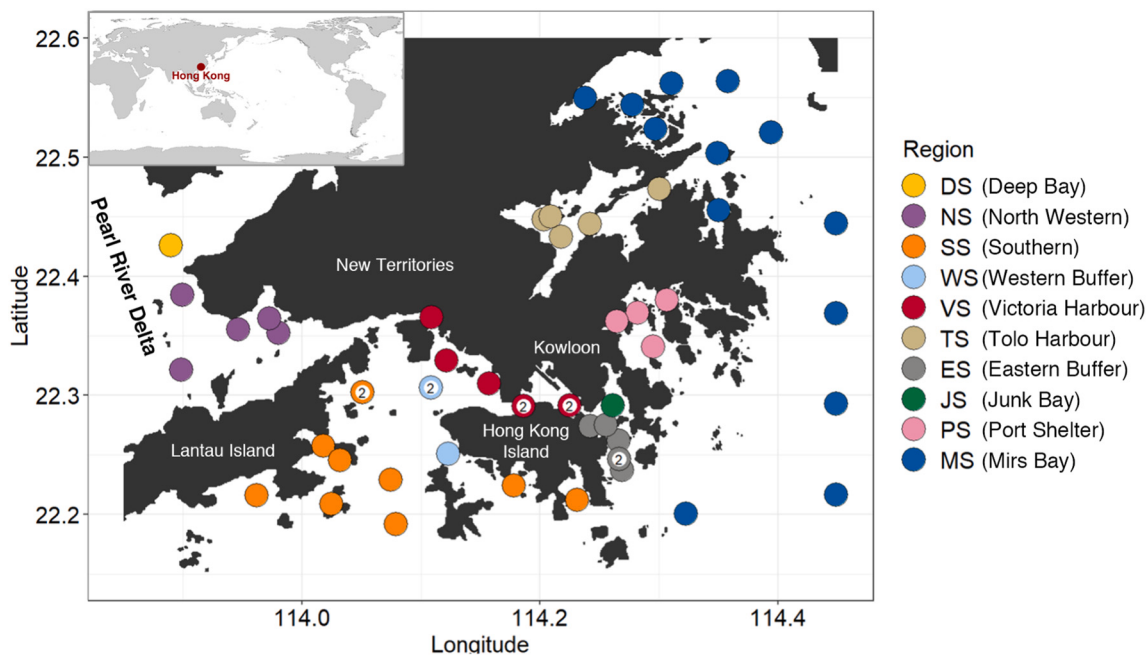


Fig. 1. Locality map of Hong Kong. Shown are the samples from the EPD marine water and sediment quality monitoring program sites (solid circles; ostracod data from Hong et al., 2019, 2021), and 10 newly studied samples for ostracods (open circles; There are two samples in a site within 50 m distance as indicated; ostracod data herein). Index map shows the location of Hong Kong.

sediment monitoring program of Hong Kong Environmental Protection Department (EPD) in 1986 began biweekly to monthly water quality and biannual sediment quality measurements (EPD, 2011).

A growing number of studies have tried to identify organisms that are sensitive to anthropogenic impacts, which can thereby serve as bioindicators for benthic conditions in coastal and nearshore environments (Diaz and Rosenberg, 1995; Balsamo et al., 2012; Yasuhara et al., 2012a; Yasuhara et al., 2019). Ostracods are meiofaunal crustaceans that have been widely used to monitor the response of benthos to stressful conditions (Bodergat et al., 1997; Bodergat et al., 1998; Alvarez Zarikian, 2000; Yasuhara et al., 2003; Frenzel and Boomer, 2005; Ruiz et al., 2005; Yasuhara and Yamazaki, 2005; Yasuhara et al., 2007; Yasuhara et al., 2012a; Irizuki et al., 2015; Irizuki et al., 2018). They are well-suited to serve as bioindicators because of their high abundance and diversity, relatively short life span, sensitivity to environmental variation, and rich fossil records (Ruiz et al., 2005; Yasuhara et al., 2012a). For this study we investigated Recent ostracods in surface sediment samples in Hong Kong in order to understand biodiversity and biogeographic distributions and their relations to natural and anthropogenic environmental factors in a highly urbanized seascape.

2. Materials and methods

2.1. Study area and materials

Fifty-two surface sediment samples (Fig. 1) were collected from the EPD marine sediment quality monitoring sites in 2011 by Van Veen grab sampler. The ostracod data from these samples is from Hong et al. (2019, 2021). For the current study, we also gathered new data from ten Van Veen grab samples from five sites (i.e., two replicates within 50 m per site) (Fig. 1; Tang et al., 2008) in order to improve our sampling coverage in the polluted areas of central Hong Kong around Victoria Harbour, as these areas were under sampled in the previous study by Hong et al. (2021). The sampling area is divided into 10 different regions based on EPD's division of Water Control Zones in Hong Kong (EPD, 2011) (Fig. 1).

Ostracod specimens larger than 150 μm were picked from the ten

additional surface sediment samples. Smaller individuals were not analyzed since they are mostly early instar juveniles that are often not well preserved (because their shells tend to be thin and delicate) and/or are difficult to identify (see Yasuhara et al., 2009 and Yasuhara et al., 2017 for more details). All individuals were picked if the samples contained less than 200 specimens; ostracods were picked from a split if the sample contained more than 200 specimens. For counting and data analysis, an entire carapace or a single valve was considered to be one individual.

2.2. Environmental variables

For the additional 10 samples from Tang et al. (2008), water depth (*Dep*) was measured in situ using a YSI 6600 Sonde. The mud content (*Mud*) was determined using a laser diffraction particle sizer, a "Coulter® LS100". The concentration of metals (*Cu*, *Pb* and *Zn*) were analyzed using the nitric and perchloric acid digestion method (Ip et al., 2004) with inductively coupled plasma-atomic emission spectrometry (ICP-AES).

For the remaining 45 samples from Hong et al. (2021), the eutrophication- and pollution-related environmental parameters selected were surface-water chlorophyll-*a* (*Chl*), bottom-water dissolved oxygen (*O₂*), turbidity (*Tur*), and sediment metals (*Cu*, *Zn* and *Pb*). Other major environmental parameters were bottom-water mean winter temperature (*wT*), bottom-water salinity (*Sal*), and sediment mud content (*Mud*). We analyzed mean values from the EPD data over the entire monitoring period (1986–2011). The environmental data from both datasets were combined and summarized in Fig. 2.

2.3. Data analysis

2.3.1. Quantifying diversity

We used Hill numbers (Hill, 1973), $^qD = \left(\sum_{i=1}^S p_i^q \right)^{1/(1-q)}$ or the effective numbers of equally abundant species, to estimate ostracod diversity in each sample and EPD region. In the equation, the order $q \geq 0$ and $q \neq 1$, *S* is the number of species and p_i denotes the relative

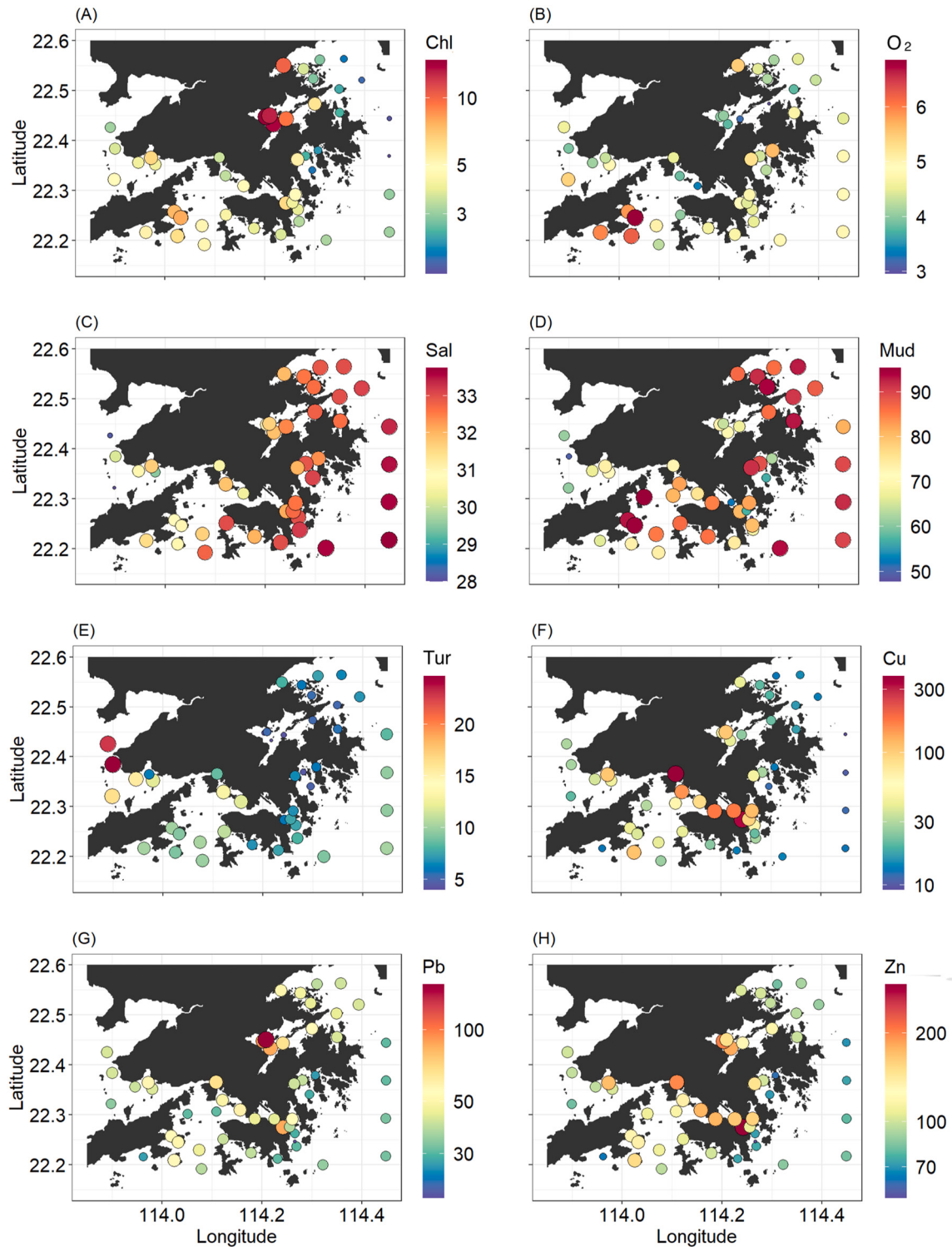


Fig. 2. Selected environmental factors, including (A) Chlorophylla ($\mu\text{g/L}$), (B) dissolved oxygen (mg/L), (C) salinity (psu), (D) mud content [$\%\text{w/w}$ ($<63\ \mu\text{m}$)], (E) turbidity (NTU), (F) copper concentration (mg/kg), (G) lead concentration (mg/kg) and (H) zinc concentration (mg/kg).

abundance of species i ($i = 1, 2, \dots, S$). Hill numbers offer several distinct advantages over other diversity indices (Chao et al. 2014b). The order q of the Hill numbers controls the sensitivity of the diversity metric to species relative abundance. When the order $q = 0$, Hill number (0D) reduces to species richness; when the order $q \approx 1$, Hill number (1D) measures the diversity of the abundant or common species; when the order $q = 2$, Hill number (2D) measures the diversity of highly abundant or dominant species (Chao et al., 2014a). The Hill numbers 1D and 2D

correspond to Shannon and Simpson diversity indices, respectively (Chao et al., 2014a). To make a fair comparison among samples and EPD regions, the Hill numbers (0D , 1D and 2D) were rarefied or extrapolated to the same sample completeness (93%), which was the largest sample coverage possible across all samples and EPD regions (Chao et al., 2020). The standard error and 95% confidence intervals of the Hill numbers were estimated by bootstrap resampling, which was repeated 1000 times.

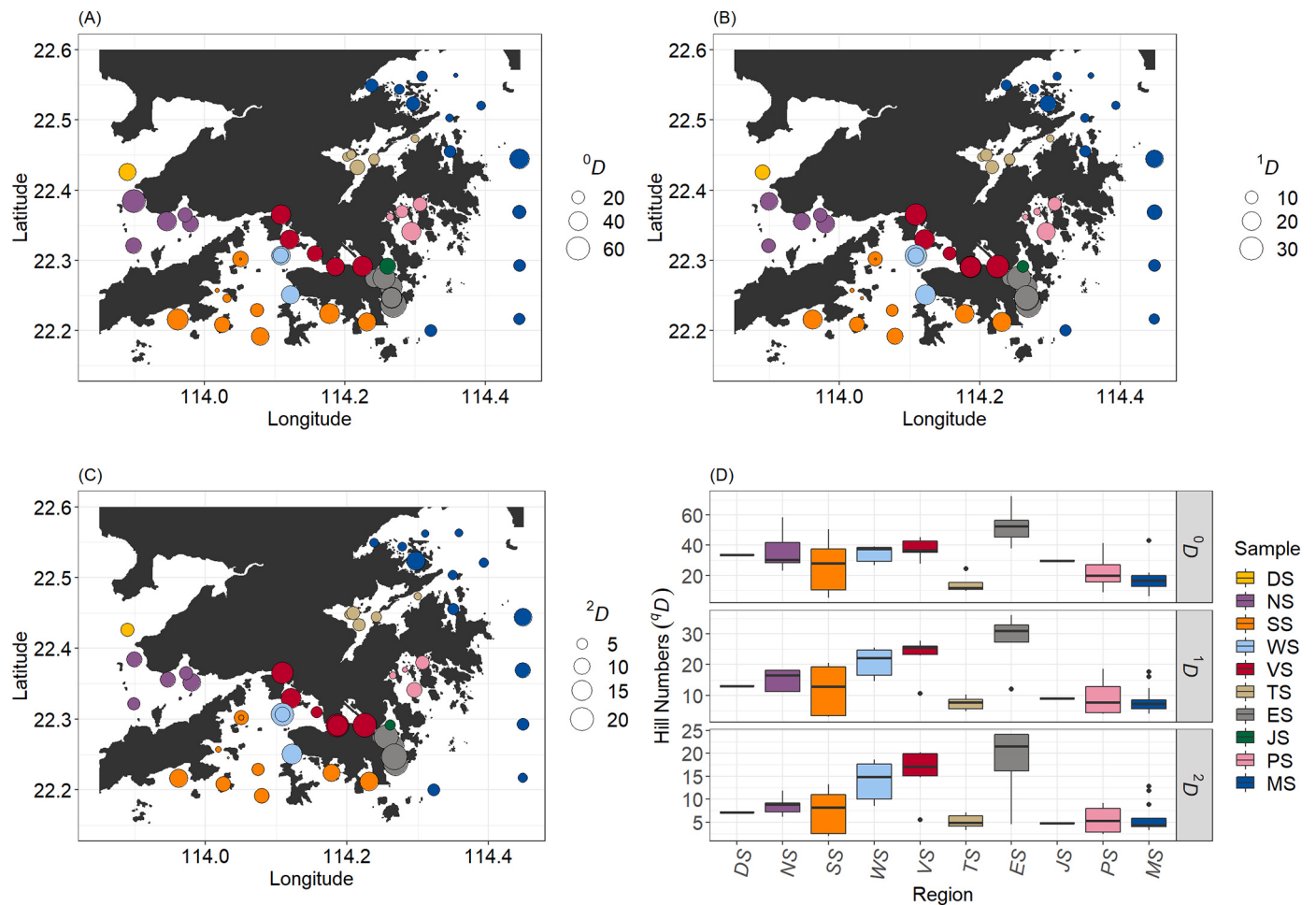


Fig. 3. Distribution (A-C) and boxplot (D) of Hill numbers 0D , 1D and 2D based on 93% sample coverage. Symbol colors in panels A-C correspond to the legend colors (i.e., EPD regions) in panel D.

2.3.2. Generalized least squares modelling

Generalized least squares modelling was used to determine the relationship between ostracod diversity (qD) and environmental parameters. All environmental parameters were log-transformed, zero centered, and normalized (divided by the standard deviation) before analysis. The normalized *Chl* were converted to absolute values to determine if there is a unimodal (or hump-shaped) relationship between diversity and productivity; i.e., whether diversity increases with productivity but declines at high productivity due to associated hypoxic conditions (cf. Yasuhara et al., 2012a). The best-fitting models were selected based on Akaike's Information Criterion (AICc) corrected for small sample size. The lower score indicates better model support considering both goodness-of-fit and model complexity (Anderson and Burnham, 2002). The Akaike weights were used to summarize the relative support for all candidate models (Anderson et al., 2000). We averaged parameter estimates over all models, with the estimates generated by each model weighted by their relative support (Anderson et al., 2000). This approach accounts for the uncertainty in model selection and generates appropriately broader confidence intervals than obtained by relying only on the single, best-supported model. The relative importance of various predictor variables was measured by the sum of Akaike weights of models that included the variables in question (Burnham and Anderson, 2002). The multicollinearity among predictor variable was examined by variance inflation factors (VIF) (Legendre and Legendre, 1998) and pairwise correlations (Yasuhara et al., 2012b). The correlation coefficients among predictors in the model were between -0.42 and 0.82 . The maximum VIFs among predictors from all 96

regression models were between 1 and 8.6 with an average of 2.8. The pairwise correlations and VIF suggest little to moderate multicollinearity ($VIF < 10$, Johnston et al., 2018). The spatial autocorrelation in model residuals was examined by Moran's I autocorrelation coefficient using the shortest distance among sampling sites as weight matrices. Out of the total 96 sets of model residuals, 53 were spatially autocorrelated (Moran's I test, $p < 0.05$). However, if we considered the top 10 best models (with the lowest AICc) for each of the 0D , 1D and 2D , only one was spatially autocorrelated ($p < 0.05$).

2.3.3. Multivariate analysis

We applied hierarchical cluster analysis (based on Ward's minimum variance) to measure the dissimilarity among ostracod faunas from different sites. Diversity decomposition was used to obtain dissimilarity measures. We used Sørensen-type dissimilarity measures, denoted as $1-C_{qN}$ for diversity order q and N assemblages, following Chao et al. (2014a). The measure $1-C_{qN}$ quantifies the effective average proportion of non-shared species in an assemblage. For $q = 0$, $1-C_{0N}$ reduces to the classical richness-based Sørensen dissimilarity (Sørensen, 1948). For $q = 1$, $1-C_{1N}$ reduces to Shannon-entropy-based Horn dissimilarity (Horn, 1966); for $q = 2$, $1-C_{2N}$ reduces to Morisita-Horn dissimilarity (Morisita, 1959). These later two measures intuitively quantify the compositional dissimilarity between two sets of species relative abundance vectors, abundant species and dominant (or highly abundant) species, respectively.

A non-Metric Multidimensional Scaling (nMDS; Gower, 1966) was performed, based on Sørensen, Horn and Morisita-Horn dissimilarities to

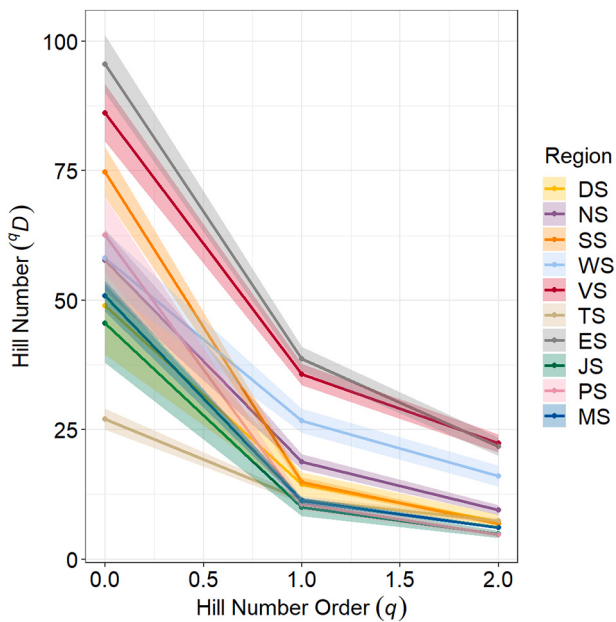


Fig. 4. Hill number profile of each EPD region based on 93% sample coverage. The shaded areas show 95% confidence interval of the profile. The overall elevation of the profile indicates the diversity based on Hill number across different order q . The levelness of the line suggests species evenness of the assemblage because a complete leveled diversity profile would indicate that 0D , 1D and 2D are all the same.

examine the relationships between assemblages. The correlations between sample ordination scores and the environmental variables were visualised by fitting the correlation coefficient as vectors on the nMDS graphs. Characteristic species (i.e., those with high occurrence or relative abundance) were mapped onto the nMDS plot using weighted average scores (WAS) (i.e., mean ordination scores weighted by species relative abundance). Stress values were calculated to quantitatively weigh the level of ‘goodness of fit’ between the original input data matrix and the ultrametric matrix of the resultant nMDS scatter plots (Kruskal, 1964; Clarke, 1993). Distance-based Redundancy Analysis (dbRDA) was used to fit environmental parameters to compositional dissimilarity (Legendre and Anderson, 1999). dbRDA converted dissimilarity into principal coordinates before conducting a traditional RDA and thus is suitable to a wide range of dissimilarity indices commonly used by ecologists (including the Sørensen, Horn and

Morisita-Horn dissimilarity in this study). The RDA is analogous to multiple regression but can be applied to composition data. The best subset of environmental variables explaining the ostracod composition was selected by ‘forward selection’ to maximize the adjusted R^2 of the dbRDA at every step and stopping when the adjusted R^2 starts to decrease (Blanchet et al., 2008).

We analyzed the full biological data (55 samples) with reduced environmental information (i.e., *Dep*, *Mud*, *Cu*, *Pb*, and *Zn*), and then the reduced biological data (45 samples) with full environmental information (i.e., *Dep*, *Mud*, *Cu*, *Pb*, *Zn*, *Chl*, *O₂*, *Tur*, *Sal*, and *wT*). All analyses were implemented in RStudio (Team, 2016). We used ‘iNEXT’ to estimate ostracod diversity (Chao et al., 2014b; Hsieh et al., 2016) and ‘vegan’ for our multivariate analyses (Oksanen et al., 2020). Figures and maps were constructed using ‘ggplot2’ (Wickham, 2012).

3. Results

3.1. Diversity

3.1.1. α -diversity of each sample

The ES region had the highest species richness (or diversity of rare species, 0D), followed by the other EPD regions to the west (i.e., VS, WS, SS, NS, and DS), and then the rest of the EPD regions to the northeast (i.e., TS, PS, and MS, Fig. 3A). When only the α -diversity of abundant (1D) and highly abundant species (2D) were considered, the ES region still had the highest α -diversity, followed by the VS and then the WS regions (Fig. 3B and C). The α -diversity of abundant (1D) and highly abundant species (2D) in the western (i.e., SS, NS, and DS) and eastern EPD regions (i.e., TS, PS, and MS) was considerably lower than in the three high diversity regions (i.e., ES, VS and WS).

3.1.2. γ -diversity of each EPD region

The Hill numbers ($q = 0, 1, 2$) pooled by each EPD region were significantly higher in the ES and VS regions than in all other regions (i.e., non-overlapping 95% confidence interval, Fig. 4); ES and VS were not significantly different from each other (i.e., overlapping 95% confidence intervals). The WS region had relatively greater diversity of abundant and dominant species (i.e., Hill number of order $q = 1$ and 2 were only lower than the ES and VS regions) than some other regions, but relatively low diversity of rare species (0D - ${}^1D = 31$ species; approximately 54% of all the species); 0D was not significantly different from NS, PS, MS, DS and JS regions. In general, Tolo Harbour & Channel (TS), Mirs Bay (MS), and Port Shelter (PS) had the lowest γ -diversity (Figs. 3 and 4).

Table 1

Environmental controls of ostracod Hill numbers 0D , 1D and 2D in Hong Kong. Model averaging on multiple linear regressions of Hill numbers and environmental factors. RI = relative importance. Asterisks show significant results ($p < 0.05^*$, $p < 0.01^{**}$, $p < 0.001^{***}$).

Hill No.		Estimate	Std. Error	Adjusted SE	z value	Pr(> z)		RI
0D	(Intercept)	28.8	1.32	1.35	21.4	0.000	***	
	Mud	-6.7	1.57	1.61	4.1	0.000	***	1.00
	Cu	11.1	3.07	3.13	3.5	0.000	***	1.00
	Pb	-5.0	2.78	2.83	1.8	0.075		0.94
	Zn	-6.8	4.39	4.47	1.5	0.130		0.93
	Dep	2.4	1.68	1.71	1.4	0.165		0.85
	(Intercept)	14.6	0.64	0.66	22.2	0.000	***	
1D	Cu	8.9	1.54	1.57	5.7	0.000	***	1.00
	Dep	2.5	0.75	0.76	3.3	0.001	**	0.99
	Zn	-5.6	2.22	2.25	2.5	0.013	*	0.98
	Mud	-2.1	0.86	0.87	2.4	0.017	*	0.96
	Pb	-2.5	1.42	1.44	1.7	0.082		0.91
	(Intercept)	9.5	0.49	0.51	18.8	0.000	***	
	Cu	7.0	1.18	1.20	5.9	0.000	***	1.00
2D	Dep	1.9	0.58	0.60	3.2	0.001	**	0.99
	Zn	-4.9	1.70	1.73	2.8	0.005	**	0.99
	Pb	-1.3	1.10	1.11	1.1	0.253		0.76
	Mud	-0.5	0.64	0.65	0.8	0.403		0.58

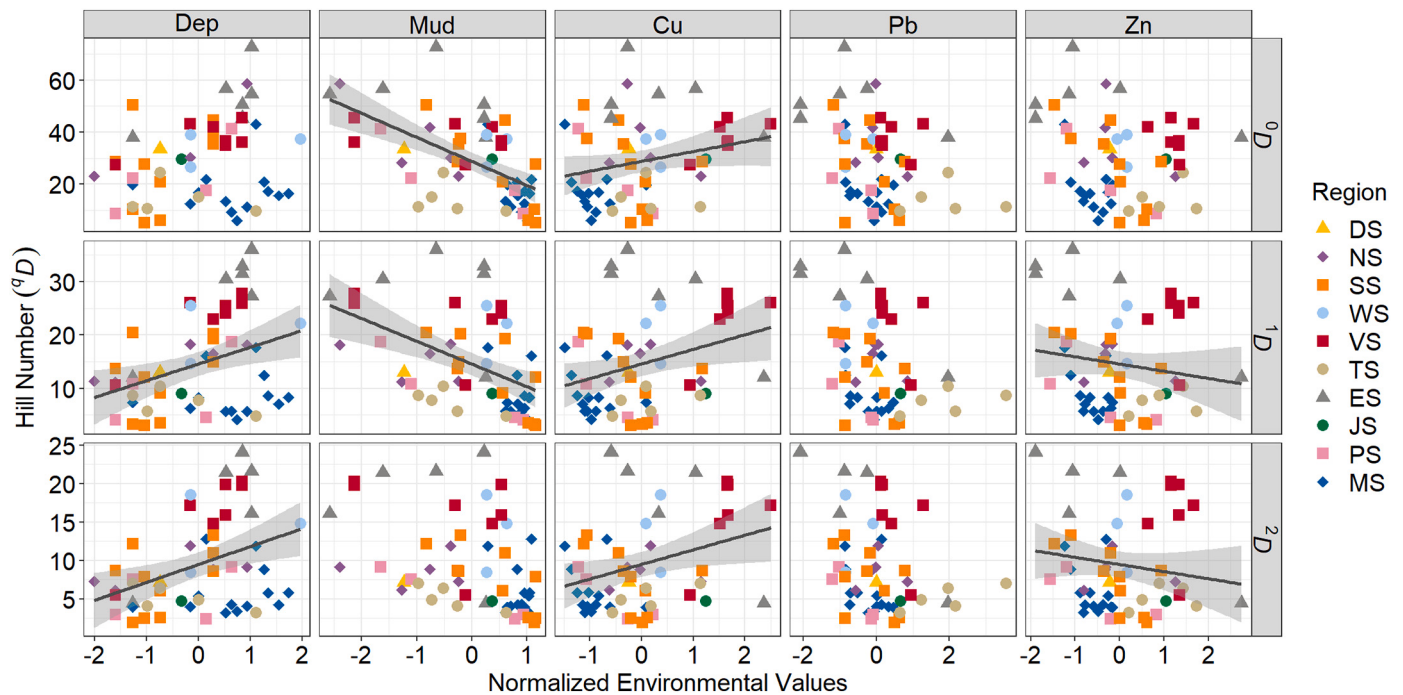


Fig. 5. Hill numbers 0D , 1D and 2D as functions of water depth (*Dep*), mud content (*Mud*), as well *Cu*, *Pb* and *Zn* concentrations in the surface sediments.

3.2. Generalized least squares modelling on α -diversity

GLS modelling and model averaging indicate that *Cu* concentration is positively associated with α -diversity (i.e., 0D , 1D , 2D) (Table 1, Fig. 5). In contrast, *Zn* concentrations are negatively associated with the diversity of abundant (1D) and dominant (2D) species (Table 1, $p < 0.05$, Fig. 5). Moreover, *Cu* had the largest positive effect size (i.e., regression coefficient), and *Zn* had the largest negative effect size (Table 1). Besides these metals, *Dep* also had significant positive effect on 1D and 2D , and *Mud* had a significant negative effect on 0D and 1D (Table 1, $p < 0.05$, Fig. 5). Each of these factors had high relative importance (RI > 0.96) based on model averaging, and thus contributed substantially to the averaged model (Table 1). When the ten additional samples were removed in order to consider more environmental predictors in the GLS model and model averaging, similar results were observed (Table S1). Metals, such as *Cu* for 0D and 1D and *Cu* and *Zn* for 2D , and *Mud* for 0D and 1D were statistically significant and had high relative importance, even for the reduced ostracod dataset (Fig. S1). The absolute value of the normalized *Chl* concentration, *Abs(Chl)*, did not have significant effects on diversity in the averaged models (i.e., for 0D , 1D and 2D) (Table S2). However, when we removed metals from the GLS and model averaging, *Abs(Chl)* was significant for 0D and 1D , and *Mud* was significant for 0D , 1D and 2D (Table S2); as values of *Chl* increased or decreased relative to the mean, the diversity of rare and abundant species declined.

3.3. Multivariate analysis

3.3.1. Composition based on presence/absence of species

The resulting nMDS plot (Fig. 6) showed positions of ostracod assemblages, colored after their clustering. nMDS analysis based on Sørensen dissimilarity ($1 - C_{0N}$) showed distinct separations of biofacies 2 (blue symbols) from 3 (green symbols), and 2 from 4 (purple symbols, Fig. 6A, B). Biofacies 2 was associated with increasing *Cu*, whereas biofacies 3 was associated with increasing *Pb* and *Zn* and biofacies 4 was associated with increasing *Mud* (Fig. 6B). Biofacies 2 mainly included stations located within the ES and VS regions. Several species occurred in 100% of the sites in biofacies 2 (Table 2). *Loxiconcha malayensis* (#75) and *Xestoleberis* spp. (#145) had weighted averages scores (WAS)

that overlapped with biofacies 2 (Fig. 6B). Biofacies 3 extended across the TS region and northern part of MS region. Biofacies 3 and 4 were similar (i.e., from the same root on the dendrogram). Together, biofacies 3 and 4 occupied the south-western and north-eastern sites off the Kowloon Peninsula and Hong Kong Island (Fig. 6A). *Sinocytheridea impressa* (#132) had 100% occurrence and WAS that overlapped with both biofacies 3 and 4 (Table 2, Fig. 6B). *Spinileberis quadriaculeata* (#134) also had 100% occurrence (Table 2) and WAS that overlapped with biofacies 3 (Fig. 6B). Biofacies 1 was close to, or overlapped with, other biofacies, and was distributed across the southern portion of the sampling area (Fig. 6A). Increasing *Dep* was the main environmental character that separated biofacies 1 from other biofacies; note that, when the ten additional samples were removed to include more environmental factors, increasing *Tur* was the main environmental factor (Fig. 7B). Only *Neomonoceratina delicata* (#88) had 100% occurrence in biofacies 1 (Table 2). It should be noted that the relatively high nMDS stress value (0.27) indicated the derived ordination configuration may not fully represent the multivariate data (Kruskal, 1964).

3.3.2. Composition of abundant species

The nMDS analysis based on abundant species (i.e., Horn dissimilarity or $1 - C_{1N}$) shows a clearer separation of clusters with lower nMDS stress ($= 0.16$) (Fig. 6C and D). Similar to the analysis based on Sørensen dissimilarity, biofacies 1 extended across the southern portion of our sampling area and was associated with increasing *Dep*; note that, when the ten additional samples were removed to include more environmental factors, both *dep* and *Tur* were the main environmental factors (Fig. 7D). The WAS for *Neomonoceratina delicata* (#88), *Pistocythereis bradyi* (#108), and *Keijella kloempritsensis* (#65) overlapped with biofacies 1 and accounted for 19.4%, 9.0%, 4.7% of the abundance, respectively (Table 2). Biofacies 2 mainly occupied the ES and VS regions and was associated with increasing *Cu*. *Neonesidea* spp. (#89) and *Loxiconcha malayensis* (#75) were the characteristic species (i.e., WAS overlapped with biofacies 2) and contributed 6.3% and 5.5% of the abundance, respectively. Biofacies 3 covered PS and TS regions and was associated with increasing *Pb* concentration. Together, biofacies 2 and 3 were separated from biofacies 1 and 4 by increasing metal concentrations (*Pb*, *Zn*, *Cu*) and decreasing *Mud* and *Dep* (Fig. 6D). *Sinocytheridea impressa*

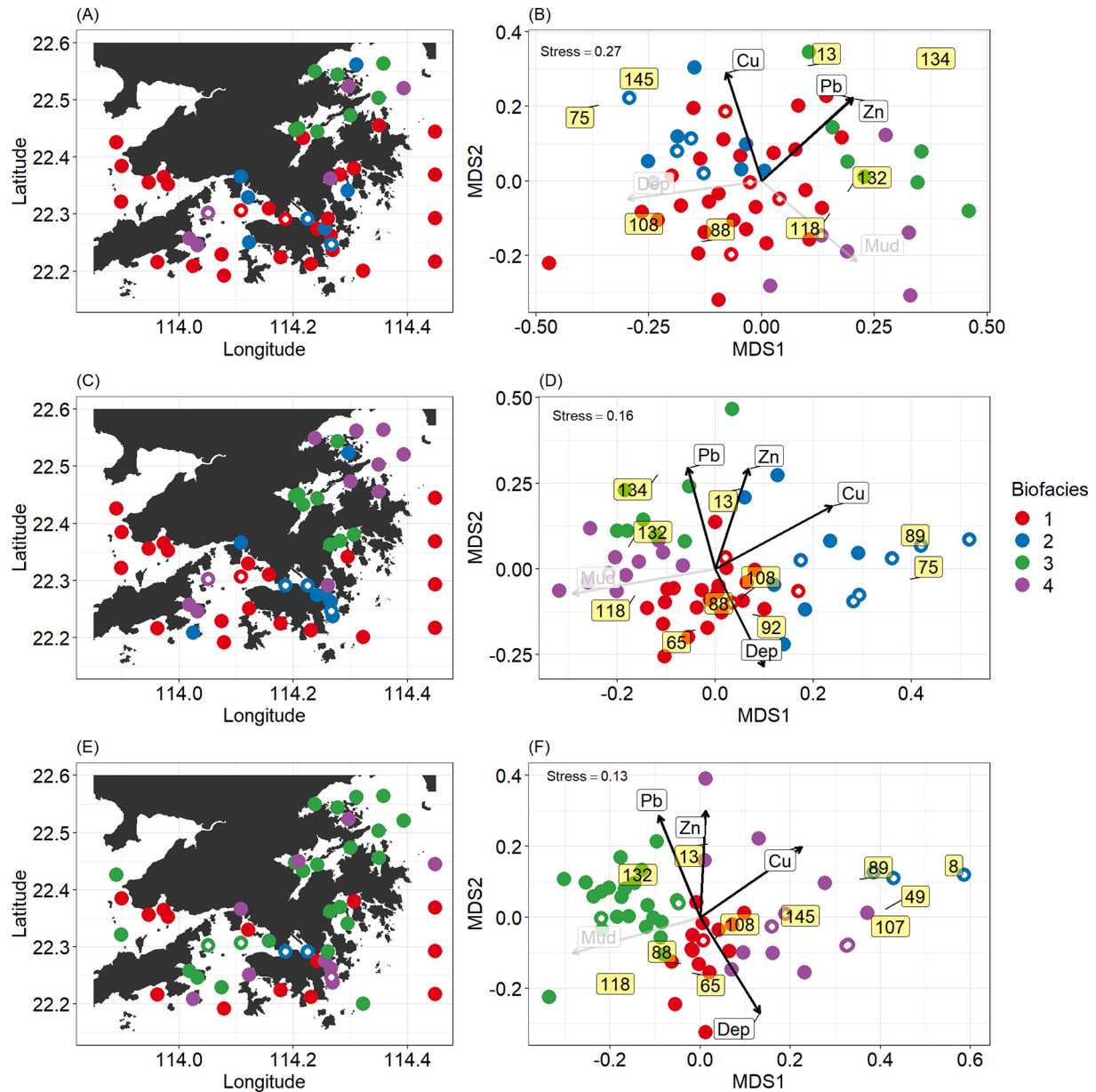


Fig. 6. Map (left panels) and nMDS ordination (right panels) showing biofacies based on (A, B) Sørensen, (C, D) Horn, and (E, F) Morisita-Horn dissimilarity and Ward minimum variance cluster analysis. The white labels and vectors on the nMDS ordination (right) show the direction and strength of correlations between environmental variables and the nMDS axes. The thick black vectors indicated the best combination variables explaining ostracod species composition (i.e., with the highest adjust R^2) from distance-based redundancy analysis (dbRDA). The yellow labels show abundance-weighted averages scores (WAS) of the top 5 species with the highest occurrence (B) or highest mean relative abundance (D, F) in each biofacies. WAS indicates where a particular species had a high abundance on the ordination space. Note that the color schemes are independent among panels A, B and C; thus, the biofacies are not related. See Fig. 1 for site details. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

Table 2

List of the top 10 species with the highest occurrence (for Sørensen dissimilarity) or mean relative abundance (for Horn and Morisita-Horn dissimilarity) from biofacies 1–4. The species code numbers (No.) correspond to the yellow labels in Fig. 4. Blank indicates that the species are not in the top 10 list but still present in the particular biofacies.

Dissimilarity	No.	Species	1	2	3	4
Sørensen	88	<i>Neomonoceratina delicata</i>	100.00%	100.00%	100.00%	100.00%
	108	<i>Pistocythereis bradyi</i>	96.70%	100.00%	100.00%	100.00%
	132	<i>Sinocytheridea impressa</i>	96.70%	100.00%	100.00%	100.00%
	13	<i>Bicornucythere bisanensis</i> s.l.	93.30%	100.00%	87.50%	83.30%
	37	<i>Copypus posterosulcus</i>	90.00%			66.70%
	65	<i>Keijella kloempritisensis</i>	90.00%			
	75	<i>Loxoconcha malayensis</i>	90.00%	100.00%	87.50%	100.00%
	118	<i>Propontocypris</i> spp.	90.00%		87.50%	100.00%
	145	<i>Xestoleberis</i> spp.	90.00%	100.00%		100.00%
	81	<i>Munseyella japonica</i>	86.70%			
	107	<i>Pistocythereis bradyformis</i>		100.00%		
	8	<i>Aurila</i> cf. <i>disparata</i>		90.90%		
	23	<i>Callistocythere</i> aff. <i>Undulatifacialis</i>		90.90%		83.30%
	38	<i>Cornucoquimba</i> cf. <i>gibboidea</i>		90.90%		
	134	<i>Spinileberis quadriaculeata</i>			100.00%	83.30%
	110	<i>Pistocythereis subovata</i>			87.50%	
	1	<i>Aglaioocypris</i> spp.			75.00%	
	46	<i>Cytherois</i> spp.			75.00%	
Horn	132	<i>Sinocytheridea impressa</i>	19.80%	6.30%	37.70%	44.40%
	88	<i>Neomonoceratina delicata</i>	19.40%	9.50%	10.60%	15.80%
	108	<i>Pistocythereis bradyi</i>	9.00%	4.20%	3.40%	5.60%
	65	<i>Keijella kloempritisensis</i>	4.70%			2.20%
	118	<i>Propontocypris</i> spp.	4.10%	6.50%	3.10%	14.60%
	91	<i>Neosinocythere elongata</i>	3.00%			
	13	<i>Bicornucythere bisanensis</i> s.l.	2.80%	3.50%	15.80%	1.10%
	92	<i>Nipponocythere delicata</i>	2.70%			2.80%
	110	<i>Pistocythereis subovata</i>	2.50%			1.70%
	139	<i>Stigmatocythere roesmani</i>	2.40%			
	89	<i>Neonesidea</i> spp.		6.30%		
	75	<i>Loxoconcha malayensis</i>		5.00%		1.00%
	145	<i>Xestoleberis</i> spp.		4.70%	2.20%	
	8	<i>Aurila</i> cf. <i>disparata</i>		4.00%		
	49	<i>Cytheropteron miurense</i>		3.20%		
	134	<i>Spinileberis quadriaculeata</i>			7.10%	2.60%
	78	<i>Loxoconcha zhejiangensis</i>			2.60%	
	73	<i>Loxoconcha japonica</i>			2.20%	
	77	<i>Loxoconcha</i> sp.			2.00%	
Morisita-Horn	88	<i>Neomonoceratina delicata</i>	23.50%		14.70%	8.80%
	132	<i>Sinocytheridea impressa</i>	16.20%		40.10%	8.90%
	108	<i>Pistocythereis bradyi</i>	9.30%	3.90%	5.40%	5.40%
	65	<i>Keijella kloempritisensis</i>	4.60%		2.40%	
	13	<i>Bicornucythere bisanensis</i> s.l.	4.20%	3.20%	4.80%	4.80%
	139	<i>Stigmatocythere roesmani</i>	2.70%			
	118	<i>Propontocypris</i> spp.	2.60%		9.40%	7.80%
	145	<i>Xestoleberis</i> spp.	2.60%	5.70%		4.30%
	110	<i>Pistocythereis subovata</i>	2.50%		1.70%	
	91	<i>Neosinocythere elongata</i>	2.50%			
	8	<i>Aurila</i> cf. <i>disparata</i>		11.20%		
	49	<i>Cytheropteron miurense</i>		7.00%		
	107	<i>Pistocythereis bradyformis</i>		5.70%		
	89	<i>Neonesidea</i> spp.		5.30%		5.00%
	84	<i>Neocytheretta faceta</i>		5.10%		
	75	<i>Loxoconcha malayensis</i>		5.10%	1.50%	4.20%
	72	<i>Loxoconcha epeterseni</i>		3.60%		
	134	<i>Spinileberis quadriaculeata</i>			3.20%	
	92	<i>Nipponocythere delicata</i>			1.90%	2.60%
	77	<i>Loxoconcha</i> sp.				2.80%

(#132), *Bicornucythere bisanensis* s.l. (#13) and *S. quadriaculeata* (#134) had WAS within biofacies 3 and contributed 37.7%, 15.8%, and 7.1% of abundance, respectively (Table 2). Biofacies 4 mainly occurred in SS and MS regions and was associated with higher *Mud*. *Sinocytheridea impressa* (#132) and *Propontocypris* spp. (#118) were the characteristic species (i. e., WAS fall within the biofacies 4, Fig. 6D) and contributed to 44.4% and 14.6% of abundance, respectively (Table 2).

3.3.3. Composition of dominant (or very abundant) species

The nMDS analysis based on dominant (or highly abundant) species (i.e., Morisita-Horn or 1–C_{2N}; Fig. 6E and F) showed discrete separation of biofacies 2 from other biofacies. Biofacies 2 was characterized by

higher *Cu*, and by the abundance of *Aurila* cf. *disparata* (#8) and *Neonesidea* spp. (#89) (Fig. 6F). These two species contributed 11.2% and 5.3% of abundance in the biofacies 2 but had relatively low abundance in other biofacies (Table 2). Despite their dissimilarity in the nMDS (Fig. 6F), biofacies 1, 3, and 4 are geographically clustered (Fig. 6E). Biofacies 1 is located in more southern locations, more subject to oceanic influence with deeper *Dep*. *Neomonoceratina delicata* (#88), *Pistocythereis bradyi* (#108), and *Keijella kloempritisensis* (#65) contributed higher relative abundance in biofacies 1 than other biofacies (Table 2, Fig. 6F). Biofacies 3 also had broad geographic extent but included more sheltered locations where *Mud* and *Pb* were higher. The characteristic species (i.e., based on WAS) in biofacies 3 was *Sinocytheridea impressa*

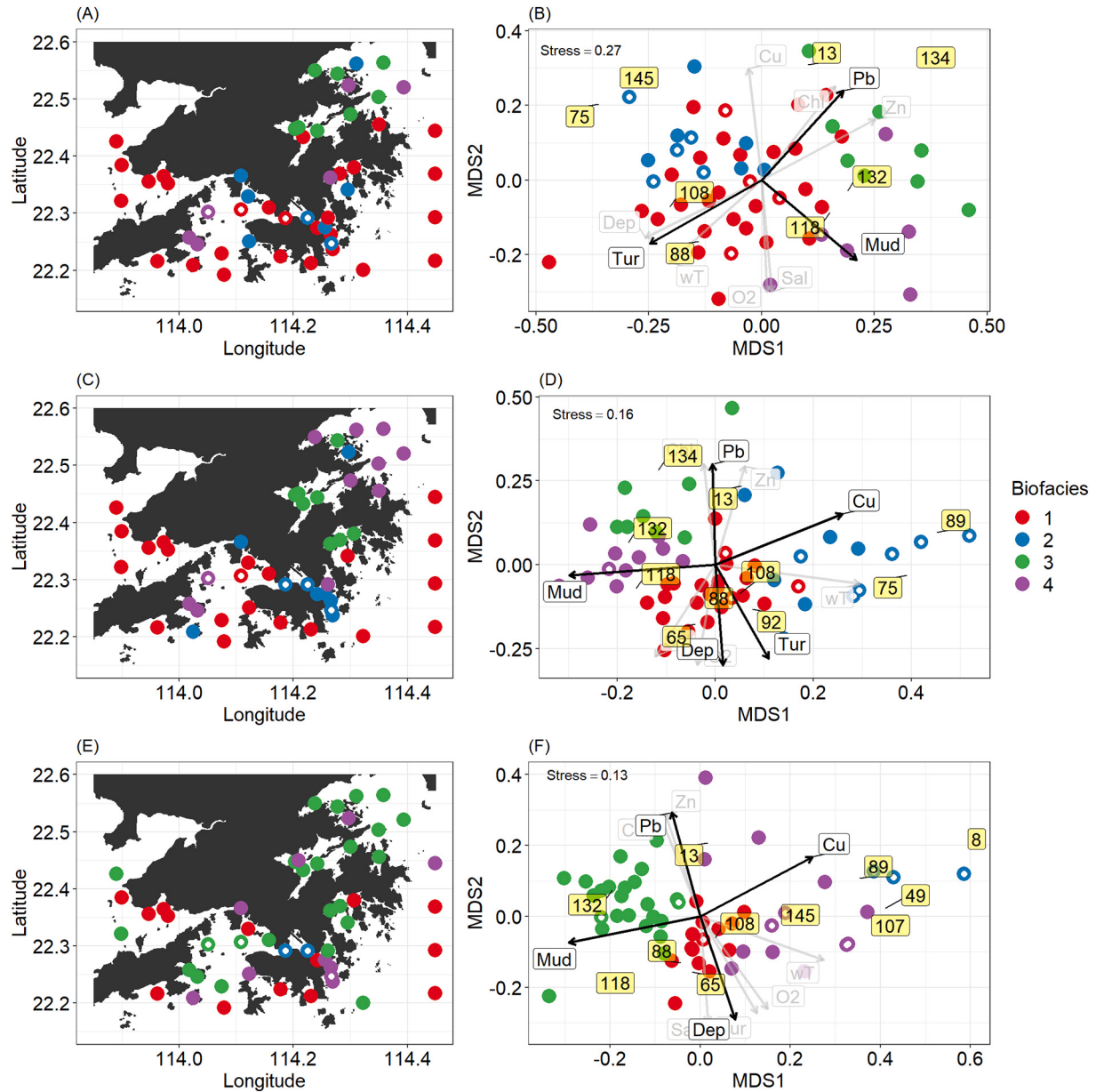


Fig. 7. Same as Fig. 6 except for vectors showing the direction and strength of correlations of all environmental variables with the same nMDS axes. The missing environmental values were excluded and thus, the correlations were only based on the 45 samples from Hong et al. (2021) (i.e., the solid circles). The thick black vectors indicated the best combination of predictors from all available environmental variables explaining ostracod species composition (i.e., highest adjust R^2) from distance-based redundancy analysis (dbRDA). dbRDA was also only based on the 45 samples from Hong et al. (2021). See Fig. 1 for site details.

Table 3

Results of distance-based redundancy analysis (dbRDA) between environmental variables and ostracod dissimilarities. *Variable*, R^2_{adj} and *AIC* show the subsequent addition of the variables, the increase of explanatory power and the decrease of Akaike's Information Criterion of the dbRDA model, respectively. *Score* shows the square root of the sum of the square of the variable loading on the first two dbRDA axes.

Dissimilarity	Variable	R^2_{adj}	Df	AIC	F	Pr(>F)	Score
Sørensen	+ Pb	0.07	1	111.9	4.9	0.001	0.95
	+ Cu	0.12	1	109.7	4.2	0.001	0.61
	+ Zn	0.17	1	107.4	4.1	0.001	0.56
	All	0.19					
Horn	+ Cu	0.19	1	59	13.7	0.001	0.94
	+ Pb	0.41	1	42.6	20.7	0.001	0.92
	+ Dep	0.48	1	36.6	8	0.001	0.71
	+ Zn	0.52	1	32.7	5.6	0.003	0.76
	All	0.54					
Morisita-Horn	+ Cu	0.13	1	89.5	9.2	0.001	0.89
	+ Zn	0.36	1	73.9	19.6	0.001	0.71
	+ Dep	0.45	1	66.6	9.4	0.001	0.73
	+ Pb	0.51	1	61	7.4	0.003	0.90
	All	0.51					

(#132), which accounted for 40.1% of abundance (Table 2). Biofacies 4 resembled biofacies 2 (i.e., they share the same root on dendrogram), and exhibited a narrower geographic extent than biofacies 1 and 3. Biofacies 4 had higher *Cu*, and occurred closer to the northeast and southwest shores of the Kowloon Peninsula and Hong Kong Island (Fig. 6F). The WAS of *Bicornucythere bisanensis* s.l. (#13) and *Xestoleberis* spp. (#145) overlapped with biofacies 4 and contributed 4.8% and 4.3% of abundance. (Table 2, Fig. 6F)

3.3.4. Environmental correlates of compositional dissimilarity

Distance-based redundancy analysis (dbRDA) showed that *Dep*, *Mud*,

Cu, *Pb*, and *Zn* could explain 19%, 54%, and 51% respectively of the total variation in Sørensen dissimilarity ($1-C_{0N}$), Horn dissimilarity ($1-C_{1N}$), and Morisita-Horn dissimilarity ($1-C_{2N}$) (Table 3). The best supported subsets of environmental variables were *Pb*, *Cu* and *Zn* for Sørensen dissimilarity ($R^2_{adj} = 0.17$), *Cu*, *Pb*, *Dep* and *Zn* for Horn dissimilarity ($R^2_{adj} = 0.52$), and *Cu*, *Zn*, *Dep* and *Pb* for Morisita-Horn dissimilarity ($R^2_{adj} = 0.51$, Table 3, Fig. 6B, D, and F). When the ten additional samples were removed, the dbRDA based on the full environmental data showed that *Pb* and *Mud* were selected for all three dissimilarities (Table S2, Fig. 7). *Tur* was also selected for Sørensen and Horn dissimilarity. *Cu* and *Dep* were also selected for Horn and Morisita-Horn dissimilarities. In general, the results of dbRDA and model selection were consistent between the reduced and full ostracod datasets.

4. Discussion

4.1. Diversity

Our modelling results revealed that metal concentrations, *Mud*, and *Dep* were the dominant environmental factors structuring meiobenthic diversity in Hong Kong's coastal waters (Tables 1 and S1). *Mud* had a greater effect on species richness (0D) (i.e., less species in higher *Mud* sites), whereas *Zn* and *Dep* affected the diversity of abundant and dominant species (1D and 2D) (i.e., lower diversities in sites with elevated *Zn* concentrations and shallower depths). Most importantly, *Cu* affected all diversity measures strongly; higher diversities were observed in sites characterized by elevated *Cu* concentrations. *Cu* was especially high in Victoria Harbour (Figs. 5 and 8), where *Cu* was greater than the ERL (Effect Range-Low: 34 mg/kg) and close to the ERM (Effect Range-Median: 270 mg/kg) of the sediment guidelines of the U.S. Environmental Protection Agency (USEPA) regarding toxicological effects on marine organisms (Long et al., 1995). In contrast, *Zn* and *Pb*

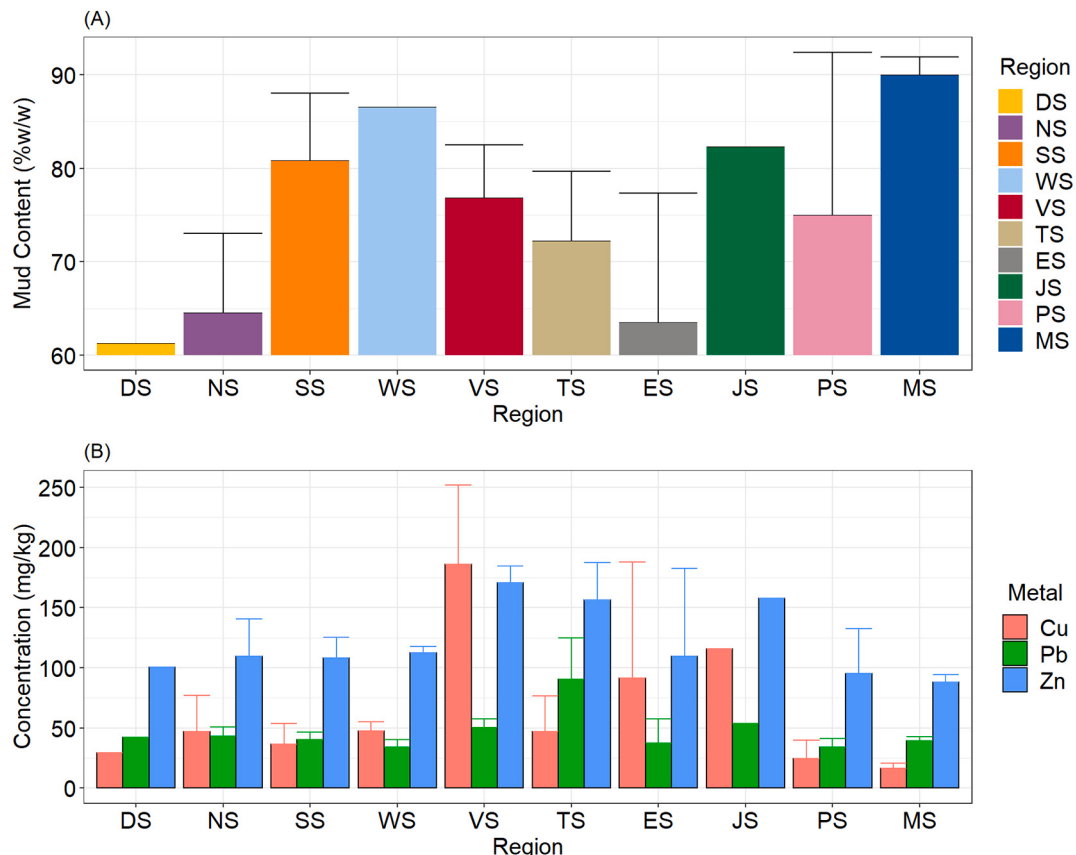


Fig. 8. Mean mud contents (A) and metal concentrations (B) in each EPD region. Error bars show the upper 95% confidence interval.

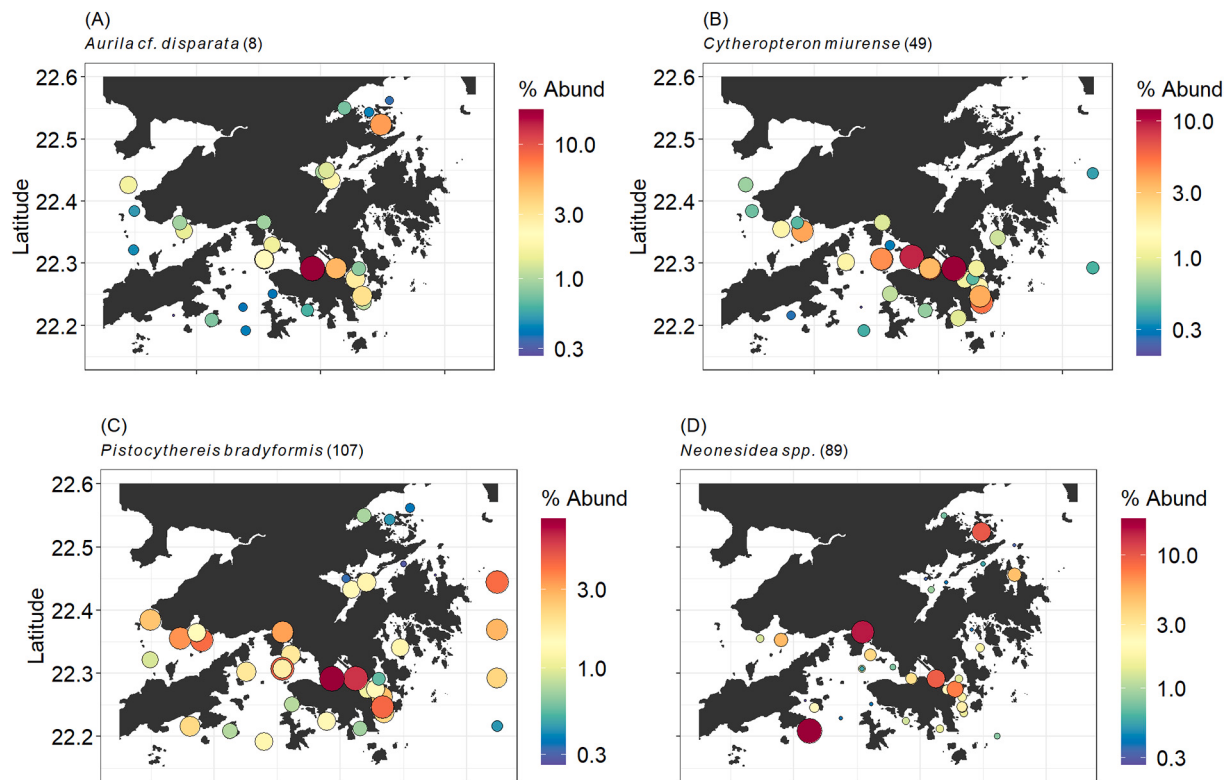


Fig. 9. Distribution and relative abundance of the most dominant species in biofacies 2 (i.e., center of Victoria Harbour) based on Morisita-Horn Dissimilarity. Color key shows relative abundance. Numbers in the bracket (following species name) correspond to the species code numbers in Table 2 and the yellow labels in Fig. 6F and Fig. 7F. *Aurilla cf. disparata* and *Neonesidea* spp. are phytal taxa, and *Cytheropteron miurense* and *Pistocythereis bradyformis* are sand dwellers. See main text for details. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

concentrations (Fig. 8) were much lower than ERM (410 mg/kg and 218 mg/kg, respectively), which was consistent with our modelling results showing weaker effects of Zn and Pb on ostracod diversity than Cu (Tables 1 and S1).

It remains unclear why Cu pollution is positively associated with meiofaunal diversity. Although finer grained sediments tend to show higher metal concentrations (Lakhan et al., 2003), it is unlikely that this relationship is producing the association between Cu and diversity, because grain size is not finer (i.e., Mud is not greater) at the high diversity sites that are affected by Cu pollution (Fig. 8). Many factors (e.g., temperature, salinity, pH, organic matters, sulfide content) are known to affect toxicity of metals (Besser et al., 2004; Campbell et al., 2014; Cooper et al., 2014; Blewett et al., 2016; Mu et al., 2018), but it is unlikely that these factors can change the Cu influence on diversity from negative to positive. Alternatively, perhaps Cu negatively affected dominant species which led to an increase in the diversity of rare species. However, our modelling results indicate that Cu affected all diversities positively, even the diversity of dominant species (i.e., 2D). The last possibility is multicollinearity, i.e., the presence of a confounding environmental factor that is highly correlated with Cu, could produce the observed positive association between Cu and diversities in our multivariate analyses. Indeed, among the environmental factors that we considered, Zn showed moderately high correlation with Cu ($\rho = 0.82$, $p < 0.001$). We assessed this possibility by separately examining the relationships between different environmental factors and diversities within the highly polluted general area of Victoria Harbour (i.e., VS, ES, and WS), and found that both Cu and Zn showed similar negative relationships with meiobenthic diversities (Fig. S2). Hong Kong's coastal waters, especially in the area of Victoria Harbour and Junk Bay (Fig. 2), have been seriously affected by metal pollution via the local discharge of sewage, industrial inputs, and surface runoff from the urban environment (Tanner et al., 2000; Warren-Rhodes and Koenig, 2001; Choi et al.,

2006; Zhou et al., 2007; Tang et al., 2008). The central part of Hong Kong is one of the busiest ports in the world and antifouling paints from ship traffic typically contain these metals (Kwok and Leung, 2005; Choi et al., 2006; Tang et al., 2008). Based on our analyses, we conclude that metal pollution has played an important role in structuring meiobenthic diversity. But note that other pollutants that were not included in our modelling (such as pesticides and polycyclic aromatic hydrocarbons) may also be highly correlated with Cu and other metals because they share similar sources (Liang et al., 2008; Wei et al., 2008), and these could also affect diversity.

The explanation for Mud's effect on diversity is much more straightforward. Low Mud, and consequently greater grain-size heterogeneity than purer mud, could foster greater meiobenthic diversities (Tables 1 and S1; Fig. 8). For benthic organisms, grain size heterogeneity may represent habitat heterogeneity which could sustain species with different ecological preferences (e.g., both sand and mud dwellers). Coarser grained sediment may also foster more phytal habitat, and consequently phytal-associated ostracod species, compared to muddy environments. Muddy substrates are often associated with lower diversities because of associated eutrophic and hypoxic conditions (e.g., Yasuhara et al., 2007), but that is not the case here, because eutrophication and deoxygenation factors (*Chl* and O_2) were each included independently in our models and neither showed any significant associations with diversities (Table S1).

Regarding geographical variation, Victoria Harbour and the adjacent Eastern Buffer, the center of Hong Kong metropolis, showed the highest meiobenthic diversity in Hong Kong (Figs. 3 and 4). In contrast, both polluted Tolo Harbour and "pristine" (i.e., far from the Pearl River, which is one of the main sources of pollution) Mirs Bay and Port Shelter showed the lowest diversities (Figs. 3 and 4). These unexpected trends can be explained, in part, by metal pollution and Mud, as discussed above. For example, the sample with the lowest diversity in Mirs Bay

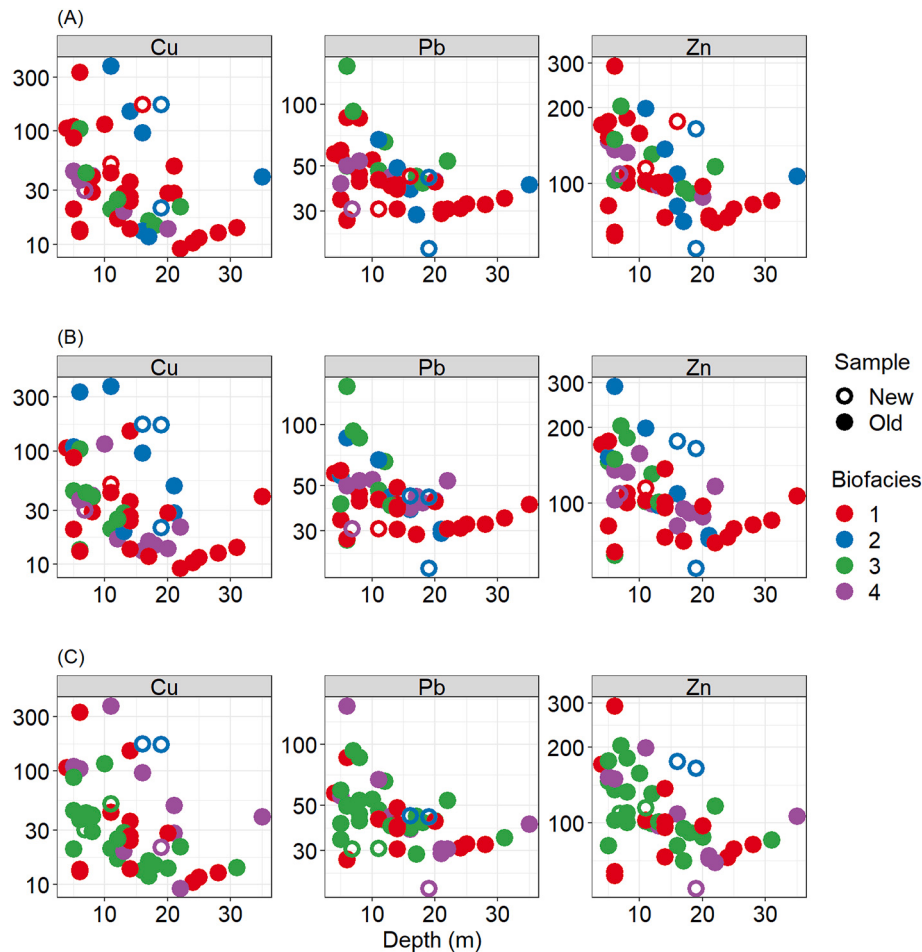


Fig. 10. Scatter plot showing the variations of copper (Cu), lead (Pb), and zinc (Zn) concentration with depth. The biofacies were based on (A) Sørensen, (B) Horn, and (C) Morisita-Horn dissimilarity and Ward minimum variance cluster analysis. Solid circles: data from Hong et al. (2021); open circles: new data in this study.

shows very high *Mud* (Fig. 5). However, these factors do not explain all of the diversity variation among regions. For example, why is diversity the highest in the Eastern Buffer? The Eastern Buffer does not have particularly high Cu values or low *Mud*, yet showed the highest diversity in Hong Kong (Figs. 3–5). A possible explanation may be surface productivity (=Chl). Diversity showed a unimodal relationship with Chl with the diversity peak at the intermediate productivity level represented by the Eastern Buffer sites (Fig. S1). Intermediate level of eutrophication may be beneficial to soft sediment benthos as observed in other embayments affected by eutrophication, such as Osaka Bay (Yasuhara et al., 2007). When primary productivity is too low, there may be insufficient food to sustain a diverse benthic fauna (Yasuhara et al., 2007). Conversely, too much food (i.e., excess food supply) can lead to the dominance of a few opportunistic species and, consequently low diversity (Yasuhara et al., 2007; Yasuhara et al., 2012a). Intensive eutrophication also causes bottom water deoxygenation, as seen in Tolo Harbour (Fig. S1), although our diversity modelling does not show any significant relationship between diversities and O_2 (Fig. S1; Table S1). In fact, lowest productivity regions of Mirs Bay and Port Shelter and highest productivity region of Tolo Harbour both show lowest level of diversities (Fig. S1). *Abs(Chl)* (for the potential unimodal relationship) showed significant model support when metal concentrations were not included in our modelling (Table S2).

4.2. Biogeography

Faunal composition tends to be distinct in Victoria Harbour and Eastern Buffer. Cluster analysis based on dominant (or very abundant)

species (i.e., Morisita-Horn) shows a distinct Victoria Harbour biofacies, while that based on rare species (i.e., Sørensen) does not show such distinct biofacies (Figs. 6 and 7). For abundant species (i.e., Horn), we observed a distinct “Victoria Harbour + Eastern Buffer” biofacies (Figs. 6 and 7). These Victoria Harbour and “Victoria Harbour + Eastern Buffer” biofacies are characterized by sand dwellers (*Cytheropteron miurense*, *Pistocythereis bradyformis*, *Loxoconcha epeterseni*) and phytal taxa (*Aurila*, *Neonesidea*, *Xestoleberis*) (Fig. 9; Table 2) (Yasuhara and Irizuki, 2001; Yasuhara et al., 2002). So, *Mud* or intermediate-level productivity may play a role not only in diversity as discussed above but also in faunal composition. But the reason behind this is elusive, since these biofacies do not show any clear trends with any environmental variables including *Mud* and *Chl* (Figs. 10, 11, and S3).

Hong Kong ostracod biogeography tends to be characterized by a vast southern water biofacies, evident in cluster results of rare (Sørensen) and abundance species (Horn) (Figs. 6 and 7). Environmental coverage is more limited for the full dataset (Fig. 6), but analyses of the restricted sample dataset indicated that high *Mud* and high *Tur* are indicative of the southern water fauna (Figs. 7 and 11), which is characterized by *Sinocytheridea impressa*, *Neomonoceratina delicata*, *Pistocythereis bradyi* (Fig. 12; Table 2). These species (especially, *Sinocytheridea impressa* and *Neomonoceratina delicata*) are known as typical mud dwellers, preferring high *Tur* environments like the East and South China Seas (Zhao and Wang, 1988; Irizuki et al., 2005; Irizuki et al., 2006; Yasuhara and Seto, 2006; Irizuki et al., 2009; Tanaka et al., 2011; Hong et al., 2019). Tolo Harbour (and Port Shelter) shows distinct fauna in rare (and abundant) species (biofacies 3; Figs. 6 and 7). The fauna is characterized by low diversity (Figs. 3 and 4). Tolo Harbour and Port

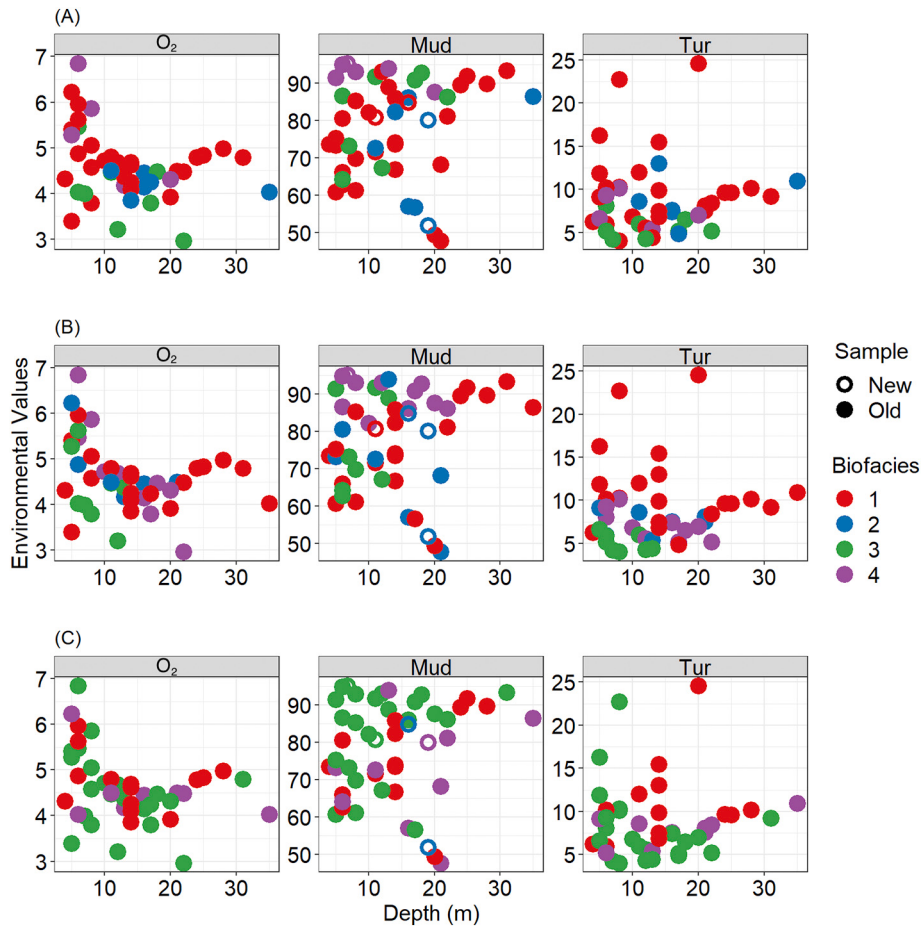


Fig. 11. Scatter plot showing the variations of oxygen contents, mud contents, and turbidity with depth. The biofacies were based on (A) Sørensen, (B) Horn, and (C) Morisita-Horn dissimilarity and Ward minimum variance cluster analysis. Solid circles: data from Hong et al. (2021); open circles: new data in this study.

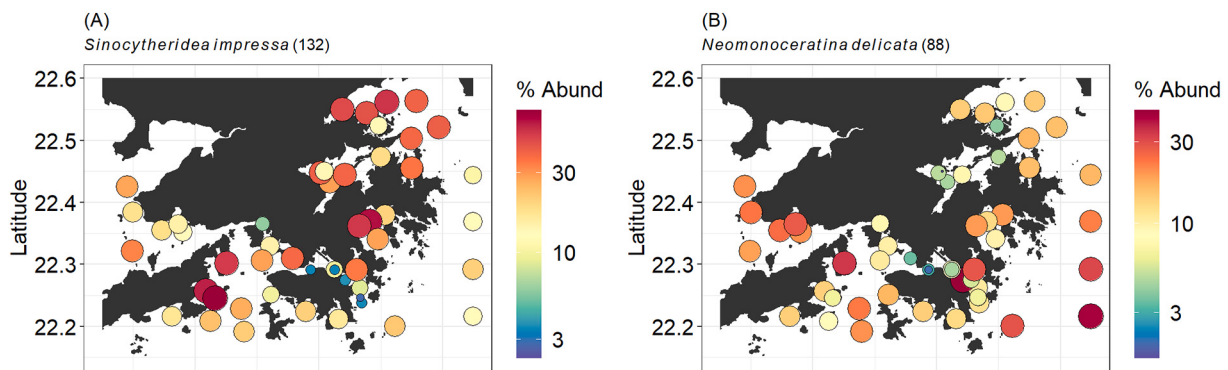


Fig. 12. Relative abundance of the dominant species *Sinocytheridea impressa* and *Neomonoceratina delicata*. Numbers in the bracket (following species name) correspond to the species code numbers in Table 2. These species are typical mud dwellers, preferring high turbidity environments (see main text for more details).

Shelter faunas may be controlled by (Pb) pollution, because the biofacies 3 sites of rare (Sørensen) and abundance species (Horn) are characterized by high Pb concentrations (Fig. 10A and B), although Pb pollution level (Fig. 8) is not high in regard to the USEPA sediment guidelines (ERM: 218 mg/kg).

4.3. Comparison with other taxonomic groups

Hong Kong macrofauna that occur in soft sediment environments also show a diversity peak in the central region, and a vast southern water biofacies, similar to those we observed in our study of meiofaunal

ostracods (Shin and Thompson, 1982; Shin and Ellingsen, 2004; Wang et al., 2017). Thus, geographic variation in the diversity of soft sediment benthos, and the environmental factors responsible for diversity structure, are congruent across different taxonomic groups. In contrast, scleractinian corals (hard coral) exhibit markedly different diversity patterns. Although Hong Kong is labelled as a marginal reef, one characterized by slow coral growth due to marginal environmental conditions for hard corals such as high Tur and low wT, it still harbors high biodiversity with over 85 species of hard coral. Hong Kong scleractinian coral diversity shows strong East to West decline controlled by the productivity gradient, with more eutrophic waters occurring in the west

(Duprey et al., 2016; Ng et al., 2017; Yeung et al., 2021). Their highest diversities are found in low productivity sites in Mirs Bay and Port Shelter (Duprey et al., 2016). This contrast makes sense, because hard corals are predominantly found in oligotrophic (low nutrient) waters and are quite sensitive and susceptible to environmental stressors, especially eutrophication (Cybulski et al., 2020; Duprey et al., 2020). These results suggest that meiofaunal ostracods can be used more broadly as a bioindicator for soft sediment diversity structure.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.marmicro.2021.102067>.

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