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Macrobenthic assemblage structure and organismal stoichiometry control faunal processing of particulate organic carbon and nitrogen in oxygen minimum zone sediments

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Abstract

The Arabian Sea oxygen minimum zone (OMZ) impinges on the western Indian continental margin between 150 and 1500 m, causing gradients in oxygen availability and sediment geochemistry at the sea floor. Oxygen availability and sediment geochemistry are important factors structuring macrofaunal assemblages in marine sediments. However, relationships between macrofaunal assemblage structure and sea-floor carbon and nitrogen cycling are poorly understood. We conducted in situ ¹³C:¹⁵N tracer experiments in the OMZ core (540 m [$[O_2] = 0.35 \mu\text{mol l}^{-1}$]) and lower OMZ boundary (800–1100 m, [$[O_2] = 2.2\text{--}15.0 \mu\text{mol l}^{-1}$]) to investigate how macrofaunal assemblage structure, affected by different oxygen levels, and C:N coupling influence the fate of particulate organic matter. No fauna were present in the OMZ core. Within the OMZ boundary, relatively high abundance and biomass resulted in the highest macrofaunal assimilation of particulate organic carbon (POC) and nitrogen (PON) at the lower oxygen 800 m stations ($[O_2] = 2.2\text{--}2.65 \mu\text{mol l}^{-1}$). At these stations the numerically dominant cirratulid polychaetes exhibited greatest POC and PON uptake. By contrast, at the higher oxygen 1100 m station ($[O_2] = 15.0 \mu\text{mol l}^{-1}$) macrofaunal C and N assimilation was lower, with POC assimilation dominated by one large solitary ascidian. Macrofaunal POC and PON assimilation were influenced by changes in oxygen availability, and significantly correlated to differences in macrofaunal assemblage structure between stations. POC and PON assimilation was characterised by carbon accumulation within the macrofauna, suggesting the importance of anaerobic metabolism at all stations. However, macrofaunal feeding responses were ultimately characterised by preferential organic nitrogen assimilation, relative to their internal C:N budgets.

1 Introduction

Episodic depositions of dead phytoplankton (phytodetritus) represent a major input of particulate organic matter (POM) to the deep-sea floor, stimulating the feeding and

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reproductive responses of the faunal and microbial assemblages that control sea-floor carbon cycling (reviewed by Smith et al., 2010). The bathyal continental margins (200–3000 m) account for a mere 7% of the global sea floor but recycle ~30% of oceanic sedimentary POM (Middelburg et al., 1997). Oxygen minimum zones (OMZs) are stable bodies of poorly oxygenated water ($<22 \mu\text{mol l}^{-1}$) persisting along the continental margins of the eastern Pacific Ocean, Arabian Sea and southwest Africa (Levin, 2003). OMZs are maintained by high primary productivity at the oceans surface and poor advective mixing of the water column, resulting in high organic matter flux and biological oxygen demand within the water column; and accumulation of organic matter in OMZ-impacted sediments (Devol and Hartnett, 2001; Hartnett et al., 1998). OMZs currently impinge upon ~6% of the continental margin sea floor (Helly and Levin, 2004) but are predicted to expand as consequences of anthropogenic climate change and ecosystem degradation (Bakun and Weeks, 2004; Stramma et al., 2008).

Metazoan macrofauna (size range: 250–1000 μm) are important contributors to ecosystem processes in deep sea sediments. Macrofauna contribute directly to OM recycling through ingestion, assimilation and respiration (e.g. Aberle and Witte, 2003; Witte et al., 2003a). Indirectly, macrofaunal subduction of POM provides an important route for deposition of organic matter deeper into the sediment (e.g. Levin et al., 1997, 1999), whilst macrofaunal bioturbation enhances sediment oxygenation; stimulates aerobic metabolism by the sediment community (e.g. Kristensen and Holmer, 2001); and provides microhabitats for microbes and meiofauna (reviewed by Giblin et al., 1995). Macrofaunal also influence sediment microorganisms through predation and competition for resources (e.g. Aller, 1994; Wieltchnig et al., 2008; Hunter et al., in preparation). At OMZ-impacted continental margins the macrofaunal assemblages are influenced by depth-dependent changes in ambient oxygen availability and differences in organic matter (OM) availability within the sediments (e.g. Levin and Gage, 1998). Macrofauna are often absent in the OMZ core, where oxygen levels are lowest; whilst high abundance, low diversity assemblages are typical at the OMZ boundaries. By contrast, seafloor outside the OMZ's influence are characterised by

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low-density, macrofaunal assemblages (Levin et al., 1991, 2000, 2009; Ingole et al., 2010). High macrofaunal abundances in the OMZ boundaries are driven primarily by hypoxia-tolerant polychaete genera, such as the ampheretid *Linopherus* spp., the spinid *Prionospio* spp., or the cirratulid *Monticellina* spp., (e.g. Levin and Edesa, 1997; Levin et al., 2000, 2009; Ingole et al., 2010). Macrofaunal diversity is positively correlated to oxygen availability, increasing concomitantly with decreases in macrofauna abundance across the OMZ boundary (e.g. Levin et al., 2000, 2009).

Quantifying the role of the fauna within ecosystem processes is important in order to develop accurate models of sea floor carbon and nitrogen cycling. Stable-isotope labelling techniques allow pathways and rates of organic matter processing to be traced over short time periods. So far, these methods have primarily been used to trace the pathways and processing of organic carbon in deep sea sediments, using ^{13}C labelled phytodetritus. Following deposition of labelled phytodetritus, macrofauna rapidly ingest the ^{13}C label and are responsible for its subduction below the sediment surface (e.g. Levin et al., 1997; Aberle and Witte, 2003; Witte et al., 2003a). Macrofauna play an important role in the initial community response, both directly consuming OM and mediating its availability to other organismal groups (e.g. Witte et al., 2003b; Moodley et al., 2005). At the OMZ-impacted Pakistan margin, macrofauna significantly influenced the fate of OM within the sediments. The polychaete *Linopherus* spp., which occurred in high densities, consumed between 45–80% of the isotopically-labelled OM deposited between 850 and 1800 m, in the OMZ lower boundary (Woulds et al., 2007). However, the relationship between macrofaunal assemblage structure and OM processing remain largely unexplored.

Organic nitrogen availability limits secondary production in marine sediments and may limit carbon cycling at the seabed (Vitousek and Howarth, 1991). Dual-labelling (^{13}C : ^{15}N) studies simultaneously traced the utilisation of organic C & N in intertidal (Rossi, 2007) and shallow subtidal sediments (Evrard et al., 2010), demonstrating that faunal and microbial N demand are important drivers of OM recycling. However, C & N are physiologically decoupled by individual organisms, which use C-rich (e.g.

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where I_{phyto} is the concentration of ${}_{\text{phyto}}\text{C}$ or ${}_{\text{phyto}}\text{N}$, I_{iso} is the amount of ${}^{13}\text{C}$ or ${}^{15}\text{N}$ incorporated, and $\text{at}\%X_{\text{phyto}}$ is the isotopic content (${}^{13}\text{C}$ or ${}^{15}\text{N}$) of the phytodetrital substrate, in atom %. Sediment C and N concentrations, and isotopic data were determined as described for the fauna, against the natural isotopic signatures of background sediment samples. Sediment carbon and nitrogen content were used to calculate the molar C:N ratio, as a proxy for sediment organic matter quality (Hedges and Keil 1995) and the percentage of ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ recovered in the sediment samples estimated for each experiment.

2.4 Statistical analysis

Environmental data for each station (Table 1) shows that alongside depth-dependent oxygen gradients, localised variations in sediment organic carbon and nitrogen content occur both within and between experimental stations. A scatterplot matrix (Fig. A1) displays the relationships between environmental variables and quantifies correlations using Spearman's rank correlation co-efficient. This matrix shows high levels of collinearity between the environmental variables. As such only ambient oxygen availability and sediment C:N ratio, were used as descriptors of each station during analysis.

Macrofaunal assemblage structure (C & N biomass) and macrofaunal assimilation of ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ were modelled as responses to changes in sediment C:N ratios and oxygen availability, between stations by Constrained Analysis of Principle Components (CAPSCALE) (Anderson and Willis, 2003). T1 540 m was excluded from analyses because macrofauna were absent at this station. The solitary ascidian at T2 1100 m was excluded from these analyses as an extreme outlier. Biomass C and biomass N were transformed by $\sqrt{(x + 0.01)}$ to reduce the influence of highly abundant taxa. ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ assimilation data were transformed by $\sqrt[3]{(x + 0.01)}$, to control for the effects of differences in incubation time at T2 800 m and T2 1100 m and reduce the influence of highly abundant taxa. Ordination models were constructed from Bray-Curtis dissimilarity matrices, with the multivariate normality of each matrix tested using beta-dispersion

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tests (1000 permutations, $p < 0.05$) (Anderson, 2006). Model selection was carried out by minimizing the Aikike Information Criterion (Bedrick and Tsai, 1994). Significance of the final ordination models were tested by permutational analysis of variance (1000 permutations, $p < 0.05$) (Anderson and Willis, 2003). Multivariate correlations between changes in ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ assimilation to changes in macrofaunal assemblage structure (biomass C and biomass N) were tested by Procrustes rotational tests (PROTEST) between final ordination models. This method compares data matrices using a rotational-fit algorithm to minimise their sum-of square residuals and calculate a goodness-of-fit statistic (m^2). The significance of the m^2 statistic was tested using a randomisation test (1000 permutations; $p < 0.05$) (Jackson, 1995). All analyses were carried in R2.9.2 (R Development Core Team 2009) using VEGAN (Oksanen et al., 2009) and MASS (Venables and Ripley, 2002) packages.

3 Results

3.1 Macrofaunal assemblage

Description of the macrofauna at each station is based upon both four and seven day experiments, with each spreader treated as a replicate. Sample sizes were $n = 3$ at stations T1 540 m and T1 800 m, $n = 5$ at T2 800 m and $n = 4$ at T2 1100 m. Metazoan macrofauna were absent at T1 540 m, where oxygen levels were $\sim 0.36 \mu\text{mol l}^{-1}$. Between 800 and 1100 m, total abundance, biomass C and biomass N were higher at T1 800 m and T2 800 m, compared with T2 1100 m (Fig. 2). In the present study the definition of macrofauna included large nematodes ($>250 \mu\text{m}$), which contributed $\sim 30\%$ of macrofaunal abundance at station T2 800 m, and $\sim 60\%$ at T2 1100 m. Nematodes contributed little C or N biomass (Fig. 63). The polychaetes were the dominant macrofaunal taxa, by biomass. However, at T2 1100 m the presence of a single large ascidian in one spreader contributed $\sim 45\%$ of C biomass and $\sim 20\%$ of N biomass. Cirratulids and sabellids were the most abundant polychaete families at T1 800 m and

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oweniids and cirratulids were abundant at T2 800 m and T2 1100 m. Cirratulids and oweniids contributed most to polychaete C & N biomass. The cirratulids were characterised by three genera: *Cirratulus* spp.; *Tharyx* spp.; and the mud-ball forming *Monticellina* spp. (Levin and Edesa 1997), and the oweniids represented by the genus *Owenia* spp. According to Fauchald and Jumars (1979) polychaete feeding group classifications, stations T1 800 m, T2 800 m and T2 1100 m were dominated by surface deposit feeding polychaetes, with sub-surface deposit feeding families playing only a minor role.

Natural abundance stable isotope signatures of the macrofauna are summarised in figure 4. The data exhibits considerably variation across all taxa. An harpacticoid copepod exhibited the most depleted $\delta^{13}\text{C}$ signature (-26‰) and an ophiuroid displayed the heaviest $\delta^{13}\text{C}$ signature (-11‰). $\delta^{15}\text{N}$ signatures typically ranged from 2 to 12‰. However, harpacticoid copepods, nyphtid polychaetes and bivalves all exhibited depleted $\delta^{15}\text{N}$ ($<2\text{‰}$). Heavier $\delta^{15}\text{N}$ signatures ($>12\text{‰}$) were observed in the turbellaria, as well as phoxocephalid crustacea and goniadid polychaetes. The natural isotopic signatures provide background data for the quantification of the ^{13}C and ^{15}N enrichments of macrofauna in each spreader experiment.

3.2 Macrofaunal uptake of phytodetrital C and N

Between 800 and 1100 m, mean phytoC and phytoN processed by the metazoan macrofauna ranged from $5540.46 (\pm 2482.74)$ to $1511.66 (\pm 1862.28) \mu\text{g C m}^{-2}$ and $355.50 (\pm 334.82)$ to $21.96 (\pm 20.04) \mu\text{g N m}^{-2}$, after 4 days; and from $2150.56 (\pm 1463.19)$ to $373.79 (\pm 414.89) \mu\text{g C m}^{-2}$ and $97.41 (\pm 84.42)$ to $31.64 (\pm 5.44) \mu\text{g N m}^{-2}$, after 7 days. Between 4 and 7 days decreases in both phytoC and phytoN assimilation were observed at T2 800 and T2 1100 m. Assimilation of phytoC and phytoN was dominated by the cirratulids at both 800 m stations (Fig. 5). However, at T2 800 m, there is a shift in phytoC and phytoN assimilation between four and seven days from the cirratulids to other taxonomic groups. At T2 1100 m, differences in

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phytoC and phytoN assimilation were generally small between taxa. However, the solitary ascidian accounted for $\sim 76\%$ of the faunal biomass and $\sim 75\%$ of phytoC assimilation in one of the four day experiments at this station.

Biomass specific phytoC and phytoN assimilation was relatively constant between macrofaunal taxa (Fig 6). During the 4 day incubations, cirratulid polychaetes exhibited higher biomass specific assimilation at both T1 800 m and T2 800 m. However, greatest biomass specific assimilation of both phytoC and phytoN were observed in a eunicid polychaete at T2 800 m. Likewise, at T2 1100 m, relatively high biomass specific assimilation of phytoC and phytoN were observed in a solitary flabelligerid polychaete. Fewer macrofauna were recovered from the 7 day experiments and biomass specific phytoC and phytoN assimilation exhibited no taxon-specific changes between four and seven days.

3.3 Multivariate ordination models

CAPSCALE ordination models tested the significance of environmental changes upon macrofaunal assemblage structure and phytoC and phytoN assimilation, between stations (Fig. 7). The influence of sediment C:N ratios upon assemblage structure (biomass C and biomass N) and feeding responses (phytoC and phytoN assimilation) could not be isolated from stochastic variations in the data and were excluded during model selection (Supplement). CAPSCALE ordinations demonstrated significant relationships between ambient oxygen availability and macrofaunal assemblage structure (Fig. 7a, b). In addition, spatial variability within the ordination models indicates unconstrained heterogeneity in macrofaunal assemblage structure at all three stations. CAPSCALE ordinations showed that changes in oxygen availability to have significant effects upon macrofauna phytoC & phytoN assimilation (Fig. 7c, d). Unconstrained spatial variability was again a feature of the ordinations, suggesting that phytoC and phytoN assimilation by the macrofaunal assemblage is influenced by assemblage structure. Procrustes rotational analysis confirms this by demonstrating significant correlations between

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Tharyx spp.; *Monticellina* spp.) and the oweniids (*Owenia* spp.) between 800 and 1100 m. This contrasts with Ingole et al.'s (2010) study at the Indian margin, highlighting the influence of spatial and temporal heterogeneity upon benthic macrofaunal assemblages. Within OMZs, zonation of the macrofaunal assemblages is associated with taxon-specific differences in tolerance to dysoxia, causing the abundance of annelids, molluscs, crustacea and echinoderms to increase sequentially (Levin et al., 2000, 2009). The spatial-scale of present study is too small to observe this zonation. However, family-level changes in macrofaunal distribution occurred across the lower OMZ boundary exhibiting a significant relationship to differences in oxygen availability. Previously, changes in benthic macrofaunal assemblages have been linked to changes in both oxygen availability and sediment geochemistry, on OMZ-impacted margins (Levin and Gage, 1998; Levin et al., 2009). The present study could not isolate the effects of sediment geochemistry upon macrofaunal assemblage structure from the spatial heterogeneity at each station. Nevertheless, it is feasible that small-scale changes in sediment geochemical parameters, at each station, may be an important factor driving the spatial variability of the macrofaunal assemblages.

4.2 Linking assemblage structure to macrofaunal feeding responses

Traditionally, the feeding ecology of deep-sea macrofauna has been investigated using specimen morphology and comparison with the feeding modes of shallow-water analogues (e.g. Jumars et al., 1990; Pagliosa, 2005). The use of both natural stable-isotopic signatures and stable isotope-labelling experiments provide powerful tools to test these assumptions and trace the flow of carbon and nitrogen through macrofaunal assemblages (Aberle and Witte, 2003; Witte et al., 2003a; Sweetman and Witte, 2008a, b; Gontikaki et al., 2011). In the present study, the macrofaunal assemblages at both 800 and 1100 m were dominated by polychaetes, such as the cirratulids and oweniids, defined as surface deposit feeders by Fauchald and Jumars (1979). Natural stable-isotopic signatures of macrofauna at the Indian margin support this assumption,

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with faunal $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ signatures (Fig. 4) similar to the natural $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of sediment OM at each station (Appendix B). Natural abundance $\delta^{13}\text{C}$ values were similar to those of marine phytoplankton (Fry and Sherr, 1984) indicating the importance of phytodetritus as an OM source for these macrofauna.

The present study compares macrofaunal feeding responses across the OMZ-impacted Indian margin, tracing the uptake $^{13}\text{C}:^{15}\text{N}$ labelled phytodetritus doses (650 mg C m^{-2} ; 160 mg N m^{-2}) at four stations. Feeding responses were observed at the 800 and 1100 m stations, with $_{\text{phyto}}\text{C}$ and $_{\text{phyto}}\text{N}$ uptake highest at 800 m, where macrofaunal biomass was greatest. Macrofaunal uptake decreased between four and seven days, similar to the trend observed by Witte et al. (2003b) at the Porcupine Abyssal Plain. However, in the present study decreases in assimilation of $_{\text{phyto}}\text{C}$ and $_{\text{phyto}}\text{N}$ may be primarily driven by differences in faunal biomass between experimental incubations. Macrofaunal feeding responses at the 800 and 1100 m stations exhibit a significant relationship to station-specific differences in oxygen availability. These strongly correlate to changes in macrofaunal assemblage structure observed between 800 and 1000 m. Thus, indicating that oxygen-driven changes in macrofaunal assemblage structure across the lower OMZ boundary are important drivers of macrofaunal processing of POC and PON.

It is hypothesised that family-level differences in trophic ecology and individual variation in feeding behaviour influence the feeding responses of deep-sea macrofaunal assemblages (Witte et al., 2003a; Sweetman and Witte 2008a, b). In deep sea sediments, bulk OM uptake by the macrofauna is primarily driven by the feeding responses of the dominant taxonomic groups to POM sedimentation. In particular, deposit feeding polychaetes dominate macrofaunal uptake of POM at the bathyal continental margins (Sweetman and Witte 2008a; Gontikaki et al., 2011.). In the present study $_{\text{phyto}}\text{C}$ and $_{\text{phyto}}\text{N}$ assimilation were dominated by the cirratulid polychaetes at the 800 m, whilst a solitary large ascidian dominated assimilation of $_{\text{phyto}}\text{C}$ in one experiment at 1100 m. This is consistent with observations that consumption of POM is proportional to consumer biomass (e.g. Middelburg et al., 2000; Woulds et al., 2007).

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Overall, higher somatic C:N ratios may indicate accumulation of lactate within faunal tissues. Lactate is an end product of anaerobic metabolism whose accumulation in animal tissue indicates persistent oxygen-stress (Seibel, 2011). Subsequently, we hypothesise that modest energy yield from anaerobic metabolism increases faunal demand for organic C, but limits its mineralisation to dissolved inorganic carbon, resulting in the observed preferential assimilation of ${}_{\text{phyto}}\text{C}$.

Animals are characterised by stoichiometric homeostasis, adapting their feeding responses to maintain nutrient consumption at an optimum level (Frost et al., 2002, 2005). The ${}_{\text{phyto}}\text{C}:\text{}_{\text{phyto}}\text{N}$ ratio for biomass specific uptake (Fig 8c) supports the proposed hypothesis. These data show relationships between ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ assimilation were similar across the major taxa, driven by a strong N preference. This indicates that macrofaunal feeding maximises organic N uptake relative to an internal nutrient budget, approximated by individual somatic C:N ratios. Macrofauna contribute to OM remineralisation through respirations (DIC) and ammonification (DIN), and so assimilation of ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ into their tissues accounts for only a fraction of their activity. In marine sediments macrofaunal are ammonotelic, mineralising organic N directly to ammonia which is rapidly excreted (Wright, 1995). Therefore, much of the ${}_{\text{phyto}}\text{N}$ processed by fauna is excreted, whilst the low-oxygen conditions resulted in ${}_{\text{phyto}}\text{C}$ accumulation within their tissues. In a complimentary study, Hunter et al. (in preparation) observed microbial activity was limited by the presence of fauna across the Indian margin OMZ. Although faunal assimilation of ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$ were relatively small, they were concomitant with losses of labile OM and ${}_{\text{phyto}}\text{N}$ from the sediments. This indicates that faunal activity is an important control upon C and N cycling in low-oxygen sediments.

5 Conclusions

Within the lower boundary of the Indian margin OMZ, changes in oxygen availability directly influence the composition of the macrofaunal assemblage and macrofaunal

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assimilation of ${}_{\text{phyto}}\text{C}$ and ${}_{\text{phyto}}\text{N}$. Differences in short term POC and PON processing by macrofauna were directly related to oxygen driven changes in assemblage structure. At present there is a lack of data on macrofaunal feeding behaviour, with much emphasis placed upon the role of broad functional groups (e.g. Fauchald and Jumars, 1979; Pagliosa, 2005). The present study is the first to use stoichiometric ratios of C and N to investigate macrofaunal feeding responses, identifying the importance of individual nutrient budgets as drivers of the macrofaunal pathways for organic matter recycling.

Supplementary material related to this article is available online at:

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Table 1. Environmental conditions. Oxygen, Temperature and Salinity data are as reported in Hunter et al. (2011). Porosity values represent the means (\pm SD), between 00 and 05cm sediment depth, of three background sediment cores at each station. Sediment Organic Matter and Unprocessed Algal material data are presented as the mean values (\pm SD) for the upper 5 cm sediment, within each spreader experiment.

Station	Duration (Days)	O2 Conc. (μ M)	Temp ($^{\circ}$ C)	Salinity ‰	Porosity θ	Sediment Organic Matter		Unprocessed Phytodetrital Material		
						%POC	%TN	C:N	% C	%N
T1 540m	4	0.35	11.86	35.20	0.63 (\pm 0.08)	3.11 (\pm 0.11)	0.29 (\pm 0.02)	10.67 (\pm 0.17)	1.00 (\pm 0.85)	1.00 (\pm 1.00)
						3.15 (\pm 0.16)	0.33 (\pm 0.03)	9.70 (\pm 0.35)	1.76 (\pm 2.44)	1.96 (\pm 2.62)
						2.99 (\pm 0.42)	0.30 (\pm 0.04)	9.94 (\pm 0.25)	0.56 (\pm 0.83)	0.55 (\pm 0.98)
T1 800m	4	2.09	9.83	35.08	0.73 (\pm 0.02)	3.75 (\pm 0.12)	0.43 (\pm 0.02)	8.77 (\pm 0.30)	1.19 (\pm 2.31)	0.40 (\pm 0.77)
						3.84 (\pm 1.15)	0.44 (\pm 0.09)	8.57 (\pm 0.92)	0.15 (\pm 0.28)	0.01 (\pm 0.01)
						3.89 (\pm 0.36)	0.44 (\pm 0.06)	8.90 (\pm 0.52)	0.25 (\pm 0.47)	0.05 (\pm 0.09)
T2 800m	4	2.63	9.81	35.08	0.70 (\pm 0.03)	3.37 (\pm 0.17)	0.36 (\pm 0.02)	9.45 (\pm 0.82)	0.14 (\pm 0.09)	0.04 (\pm 0.08)
						3.73 (\pm 0.45)	0.39 (\pm 0.04)	9.52 (\pm 0.97)	0.25 (\pm 0.23)	0.02 (\pm 0.04)
	7	3.61 (\pm 0.45)	0.39 (\pm 0.02)	9.34 (\pm 0.59)	0.16 (\pm 0.21)	0.03 (\pm 0.05)				
		3.31 (\pm 0.25)	0.37 (\pm 0.05)	8.99 (\pm 0.78)	0.07 (\pm 0.04)	0.00 (\pm 0.01)				
		3.29 (\pm 0.12)	0.35 (\pm 0.03)	9.49 (\pm 0.76)	0.16 (\pm 0.22)	0.01 (\pm 0.03)				
T2 1100m	4	17.46	7.60	34.67	0.74 (\pm 0.05)	3.28 (\pm 0.19)	0.35 (\pm 0.01)	9.30 (\pm 0.52)	0.15 (\pm 0.20)	0.04 (\pm 0.04)
						3.21 (\pm 0.55)	0.36 (\pm 0.06)	8.87 (\pm 1.00)	0.07 (\pm 0.10)	0.03 (\pm 0.04)
	7	3.10 (\pm 0.71)	0.35 (\pm 0.03)	8.87 (\pm 1.46)	0.13 (\pm 0.17)	0.02 (\pm 0.03)				
		3.25 (\pm 0.33)	0.36 (\pm 0.04)	9.04 (\pm 0.42)	0.13 (\pm 0.18)	0.03 (\pm 0.03)				

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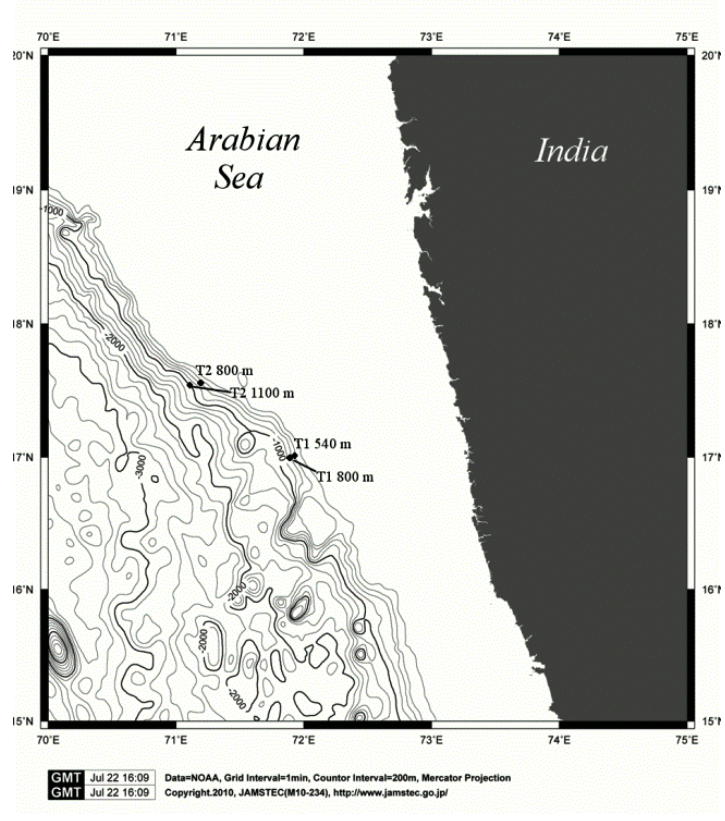


Fig. 1. Map of the study area indicating the location of experimental stations.

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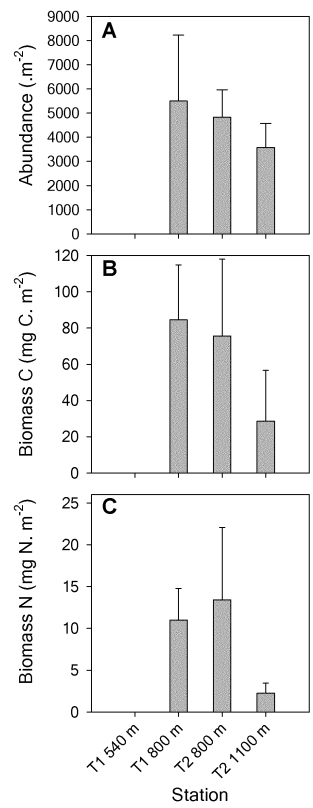


Fig. 2. Mean (\pm standard deviation) macrofaunal (A) abundance, (B) biomass C and (C) biomass N at the four experimental stations.

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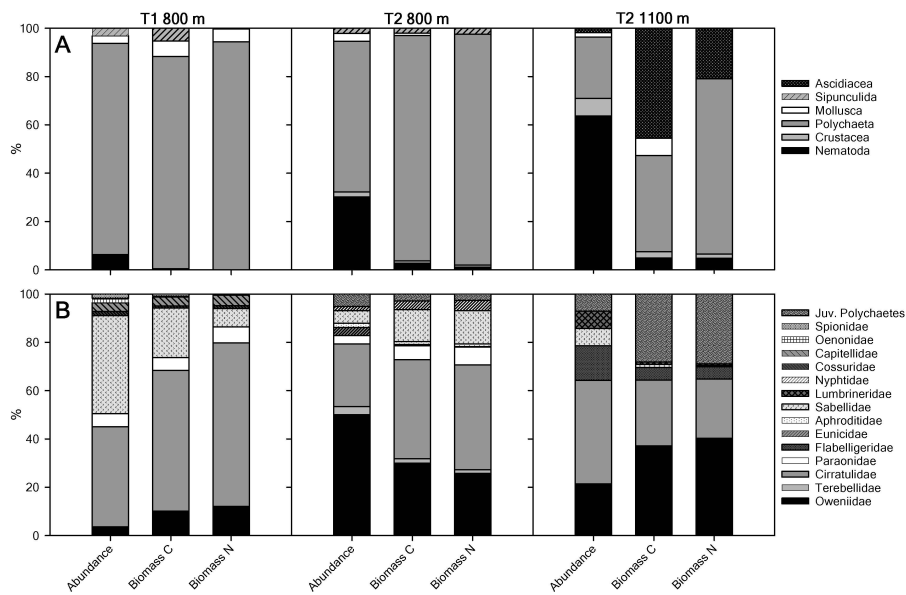


Fig. 3. Relative abundance, biomass C and biomass N of (A) the major macrofaunal taxa and (B) the polychaetes families at stations T1 800 m, T2 800 m and T2 1100 m.

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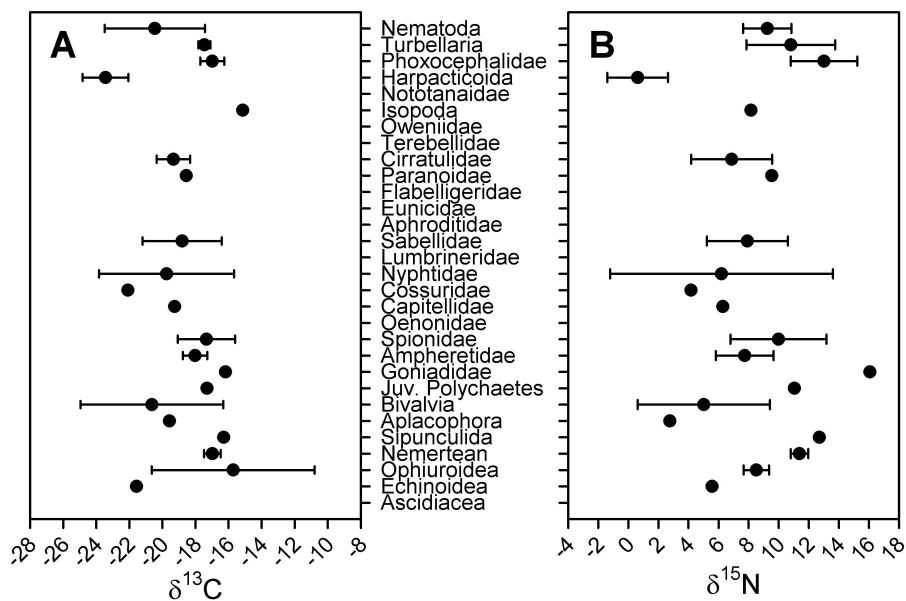


Fig. 4. Natural Stable-Isotopic signatures of macrofauna at the Indian continental margin. Data is from a series of 13 push cores taken between 800 and 1000 m, by *Shinkai 6500* during YK08-11.

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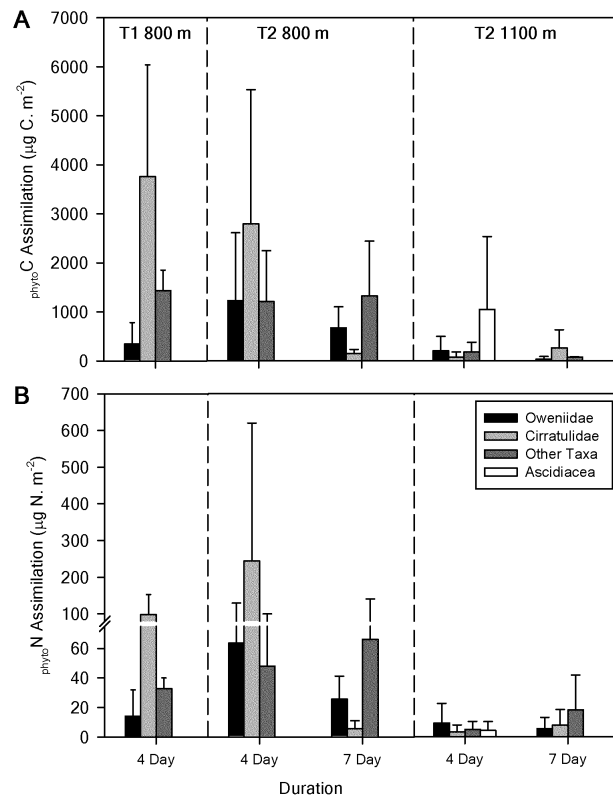


Fig. 5. Total assimilation of (A) phytoC and (B) phytoN by the main macrofaunal taxa, over four and seven day incubation periods.

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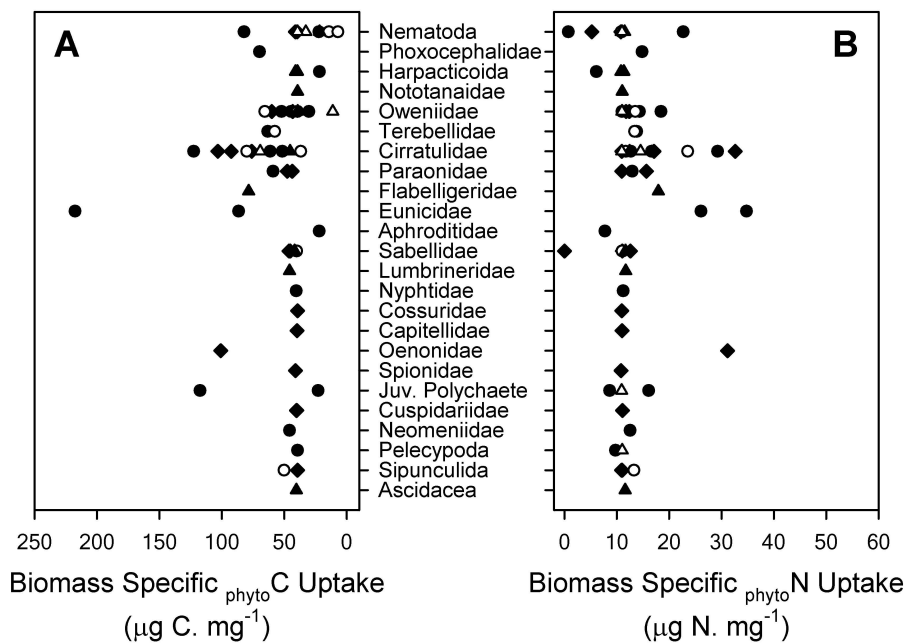


Fig. 6. Biomass specific assimilation of phytoC and phytoN by individual macrofaunal taxa at stations T1 800 m (4 day, \blacklozenge), T2 800 m (4 day \bullet , 7 day \circ) and T2 1100 m (4 day \blacktriangle , 7 day \triangle).

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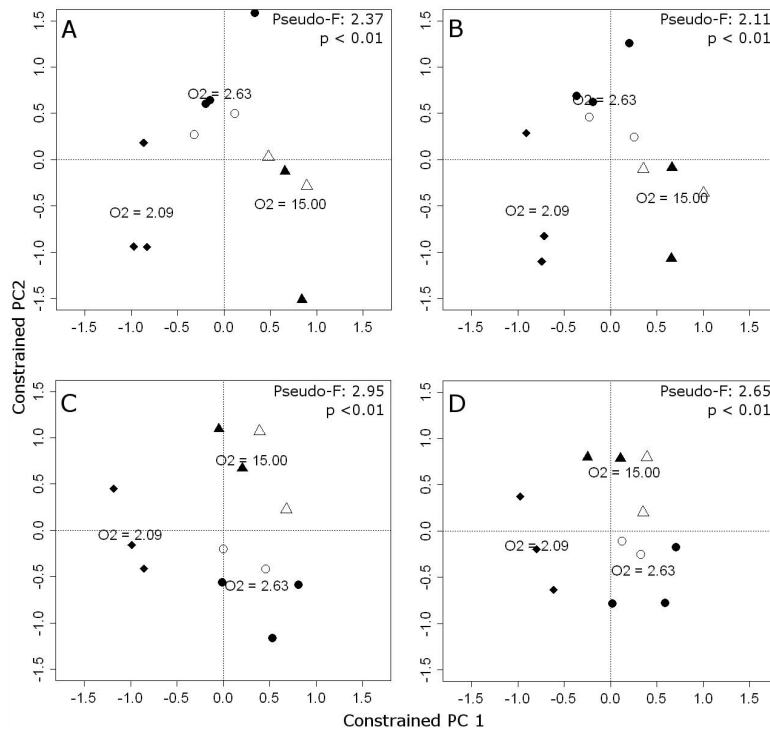


Fig. 7. Constrained ordinations models of macrofaunal assemblage structure, determined by **(A)** biomass C and **(B)** biomass N, and the macrofaunal assimilation of **(C)** phytoC and **(D)** phytoN , at stations T1 800 m (4 day, \blacklozenge), T2 800 m (4 day \bullet , 7 day \circ) and T2 1100 m (4 day \blacktriangle , 7 day \triangle).

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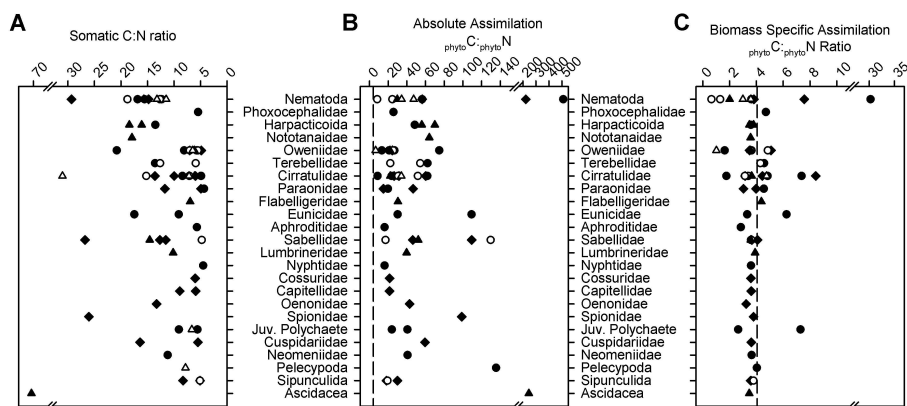


Fig. 8. C:N stoichiometry of the **(A)** somatic biomass **(B)** overall assimilation of phytoC and phytoN and **(C)** biomass specific assimilation of phytoC and phytoN of individual macrofaunal taxa at stations T1 800 m (4 day \blacklozenge), T2 800 m (4 day \bullet , 7 day \circ) and T2 1100 m (4 day \blacktriangle , 7 day \triangle). Dashed lines represent the phytodetrital C:N ratio as a reference.

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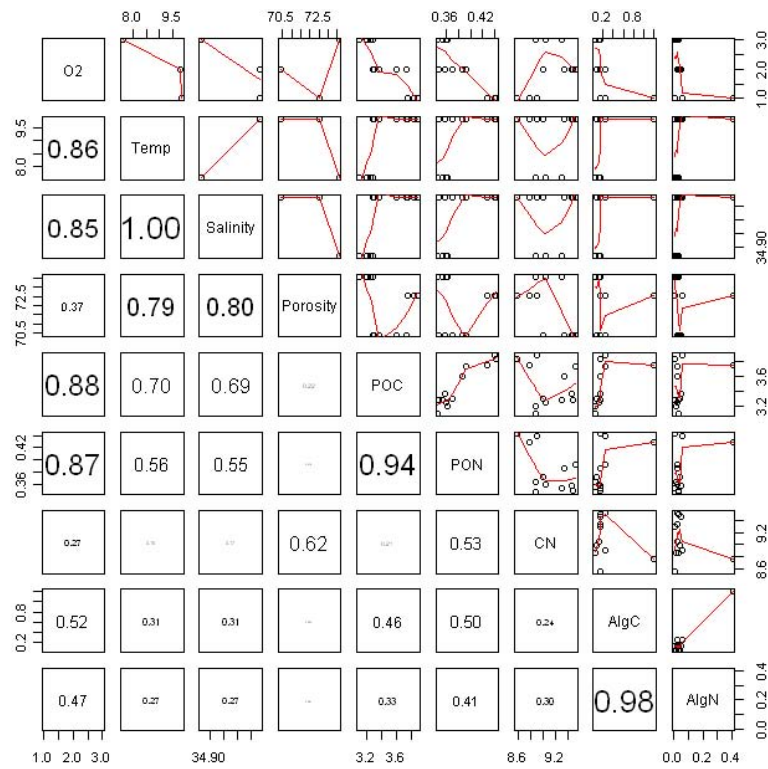


Fig. A1. Paired Scatterplots of Environmental Data, showing the level of correlation between variables as the Spearman's rank correlation coefficient.

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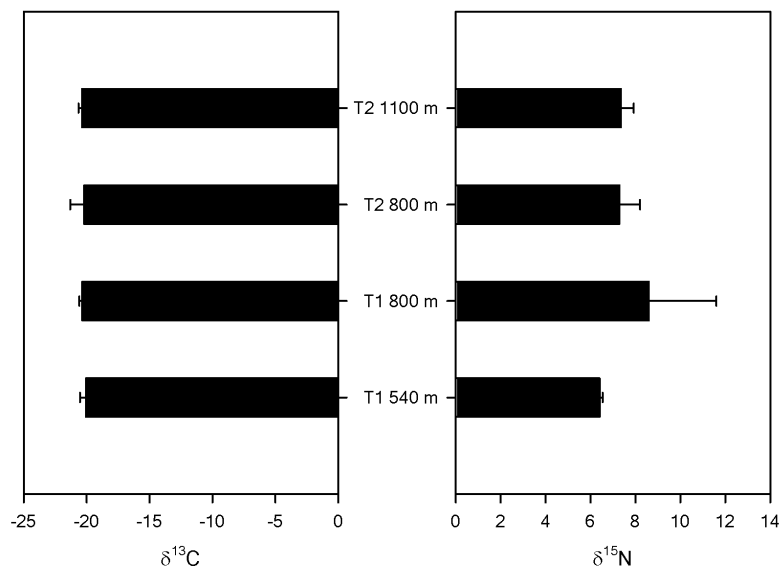


Fig. A2. Natural Stable-Isotopic signatures of sediment organic matter (00–05 cm) at the four experimental stations. Data is derived from 3 push cores taken at each station.

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