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**FUNCTIONAL DIVERSITY OF LIVING BENTHIC FORAMINIFERA OF THE SOUTHEASTERN  
PACIFIC OCEAN AND THEIR RELATIONSHIP WITH ENVIRONMENTAL VARIABLES**

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## **STATEMENTS**

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## RESUMEN

Los foraminíferos bentónicos son indicadores clave que proporcionan información sobre el funcionamiento de los ecosistemas a través de sus adaptaciones a los gradientes ambientales. En el presente estudio se evalúa la diversidad taxonómica y funcional de las comunidades de foraminíferos bentónicos vivos a lo largo del margen continental occidental de América del Sur ( $1^{\circ}$ - $50^{\circ}$ S) y a profundidades de entre 79 y 4500 m. Se identificaron un total de 365 especies, siendo Rotalida, Lituolida y Astrorhizida los órdenes más frecuentes.

La riqueza y diversidad de especies mostraron una pronunciada variabilidad espacial estructurada no por un gradiente latitudinal continuo, sino por umbrales ambientales discretos, asociados principalmente a la disponibilidad de oxígeno. Los análisis de diversidad funcional revelaron que las comunidades más ricas desde el punto de vista taxonómico ocupaban un espacio de rasgos funcionales más amplio, aunque de forma desigual en toda la zona de estudio, lo que indica una redundancia funcional considerable. Este patrón se asocia a un fuerte proceso de selección ambiental de especies (filtrado ambiental), junto con asociaciones de alta diversidad taxonómica en las que se observó una saturación parcial del espacio funcional.

Por el contrario, los sectores heterogéneos promovieron una mayor diferenciación de nichos y diversificación funcional. Los rasgos medios ponderados por la comunidad respaldaron cambios latitudinales consistentes en la morfología, las estrategias tróficas y la tolerancia al oxígeno. Los modelos aditivos generalizados identificaron la temperatura, la salinidad, el oxígeno, las concentraciones de fosfato y silicato y las características de los sedimentos como factores determinantes de la diversidad

taxonómica y funcional, siendo el oxígeno el factor estructurante dominante. Los análisis de nicho confirmaron un fuerte efecto de este filtrado ambiental en condiciones de bajo oxígeno, donde las comunidades estuvieron dominadas por especialistas que ocupaban nichos marginales.

Nuestros resultados demuestran que la disponibilidad de oxígeno regula la distribución de las especies, la configuración de los nichos y la organización funcional, modulando el equilibrio entre la redundancia y la diferenciación en los ecosistemas bentónicos del Pacífico sudoriental. Este estudio establece una base de referencia regional para evaluar las consecuencias ecológicas de la continua expansión de la OMZ en el Pacífico sudoriental.

**Palabras clave:** Foraminíferos bentónicos, diversidad funcional, rasgos funcionales, nicho, Pacífico Sur Oriental.

## **ABSTRACT**

Benthic foraminifera are key indicators that provide insights into ecosystem functioning through their adaptations to environmental gradients. This study assesses the taxonomic and functional diversity of living benthic foraminiferal communities along the western South American continental margin (1°–50°S) and water depths from 79 to 4500 m. A total of 365 species were identified, with Rotalida, Lituolida, and Astrorhizida as the most frequent orders.

Species richness and diversity exhibited pronounced spatial variability structured not by a continuous latitudinal gradient, but by discrete environmental thresholds, primarily associated with oxygen availability. Functional diversity analyses revealed that taxonomically richer communities occupied broader functional trait space, though unevenly across the study area, indicating substantial functional redundancy. This pattern emerged under environmental filtering, particularly in highly diverse assemblages where partial saturation of trait space occurred.

In contrast, heterogeneous sectors promoted greater niche differentiation and functional diversification. Community-weighted mean traits revealed consistent latitudinal shifts in morphology, trophic strategies, and oxygen tolerance. Generalized additive models identified temperature, salinity, oxygen, phosphate, silicate concentrations, and sediment characteristics as drivers of taxonomic and functional diversity, with oxygen emerging as the dominant structuring factor. Niche analyses were consistent with environmental filtering under oxygen-depleted conditions, where communities were dominated by specialists occupying marginal niches.

Our results demonstrate that oxygen availability regulates species distributions, niche configuration, and functional organization, modulating the balance between redundancy and differentiation in benthic ecosystems of the Southeastern Pacific. This study establishes a regional baseline to assess ecological consequences of continued OMZ expansion in the Southeastern Pacific.

**Keywords:** Benthic foraminifera, functional diversity, functional traits, niche, Southeastern Pacific

# 1. INTRODUCTION

## 1.1. Marine biodiversity

Marine ecosystems exhibit unique characteristics compared to terrestrial ones, as organisms have evolved under distinct physical and chemical constraints to ensure survival in the marine environment (Grassle *et al.*, 1991). The marine realm hosts approximately 35 Phyla, of which 14 are exclusive to this environment (Briggs, 1974; Gray, 1997). Over the past decades, research has advanced our understanding of marine biodiversity, including its composition, distribution and the processes that regulate community structure (Cardinale *et al.*, 2006; Costello *et al.*, 2010). In recent decades, the Southern Hemisphere has been shown to harbor several biodiversity "hotspots," largely associated with major current systems. Notably, the Benguela Current System (35°S - 15°S) (Kirkman *et al.*, 2013; Sakko, 1998; Zettler *et al.*, 2013) and the Humboldt Current System, which extends from 45°S off southern Chile to 4°S near the Peru–Ecuador border, stand out as key regions of high marine diversity (Montecino & Lange, 2009; Ramírez *et al.*, 2017; Wagner *et al.*, 2020).

In South America, extensive coastlines, heterogeneous habitats, and diverse oceanographic conditions strongly influence the richness and composition of marine invertebrate and fish assemblages across the Pacific and Atlantic coasts (Miloslavich *et al.*, 2011). The Pacific Ocean, in particular, is characterized by a high diversity of benthic macroinvertebrates groups, including Arthropoda, Crustacea, Decapoda, Echinodermata, Mollusca, Polychaeta, Bryozoa among others (Häussermann & Försterra, 2005; Hooker *et al.*, 2013; Lancellotti &

Vásquez, 1999; Mutschke *et al.*, 2017; Tasso *et al.*, 2018) as well as diverse seaweed species (Cárdenas-Calle *et al.*, 2020). This diversity reflects the wide range of habitats present, from rocky coasts to sandy substrates, spanning both intertidal and subtidal zones, and underscores the importance of environmental heterogeneity in shaping marine biodiversity patterns (Tarazona *et al.*, 2003)(Miloslavich *et al.*, 2016)

## **1.2. Southeastern Pacific Ocean (SEP)**

The Southeastern Pacific Ocean (SEP) is strongly influenced by the Humboldt Current System (HCS), one of the most productive eastern boundary current systems in the world. The HCS originates from the northern branch of the Antarctic Circumpolar Current (ACC), driven by westerly wind drift (WWD,  $\sim 42^{\circ}\text{S}$ – $48^{\circ}\text{S}$ ), and transports cold Subantarctic Surface Waters (SAAW) northward toward the Equator (Silva *et al.*, 2009; Strub *et al.*, 1998). As it flows along the coast of Peru and Chile, it divides into two currents, the Peru Ocean Current (POC) and the Peru-Chile Current (PCC) (Figure 1). On the other hand, from the tropical zone, the Equatorial Subsurface Current (EUC) flows southeastward, reaching the Peruvian coast at  $5^{\circ}\text{S}$ , influencing the formation of southward-flowing currents: the Peru-Chile Countercurrent (PCCC) and the Peru-Chile Undercurrent (PCUC) (Figure 1a, b). The PCCC divides the PCC and POC (Chaigneau *et al.*, 2013; Montes *et al.*, 2010; Morales *et al.*, 1996; Strub *et al.*, 1998). While the PCUC transports Equatorial Subsurface Waters (ESSW), which are mainly characterized by being masses of water with high salinity, rich in nutrients and poor in oxygen,

properties that are diluted along their path due to the northward advection of Antarctic Intermediate Waters (AAIW) (Figueroa & Moffat, 2000; Fuenzalida *et al.*, 2009; S & Neshyba, 1979).

A defining characteristic of the HCS is the occurrence of wind-driven coastal upwelling, which brings nutrient-rich ESSW to the euphotic zone. This process sustains extremely high primary productivity rates, estimated at approximately  $1.2 \text{ kg C m}^{-2} \text{ y}^{-1}$  (Gutiérrez *et al.*, 2016; Montecino & Lange, 2009). Consequently, the region supports highly productive marine ecosystems and diverse benthic communities. Another feature of the SEP is the presence of an extensive Oxygen Minimum Zone (OMZ), which is closely associated with the PCUC and characterized by very low dissolved oxygen concentrations (Chavez *et al.*, 2008; Montecino & Lange, 2009; Thiel *et al.*, 2007). Upwelling intensity varies along the latitudinal gradient of the HCS. Permanent upwelling occurs north of  $\sim 32^\circ\text{S}$ , where high nutrient availability sustains large phytoplankton populations dominated by diatoms (Abrantes *et al.*, 2007). While between  $\sim 33^\circ\text{S}$  and  $40^\circ\text{S}$ , coastal upwelling becomes seasonal, occurring mainly during austral spring and summer and often interacting with riverine inputs that further enhance coastal productivity (Rosales Quintana *et al.*, 2021; Strub *et al.*, 1998)

These strong gradients in oxygen availability, nutrient concentration, and productivity create heterogeneous environmental conditions that significantly influence the composition and diversity of benthic communities in the Southeastern Pacific.

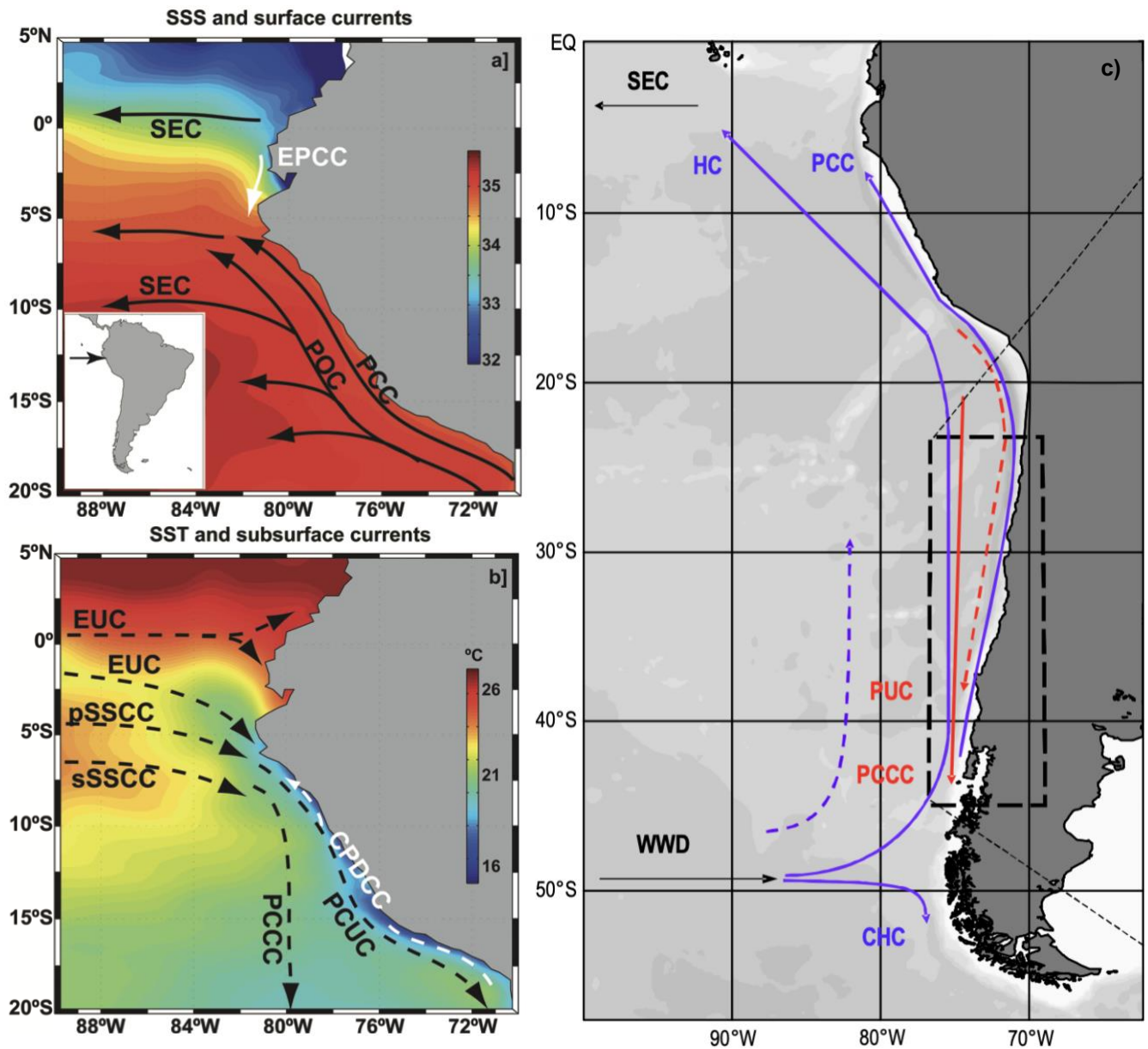


Figure 1. Main ocean currents and water masses in the Southeastern Pacific. a) Sea-surface salinity and surface circulation. b) Sea-surface temperature and subsurface circulation. Surface currents. SEC: South Equatorial Current; EPCC: Ecuador-Peru Coastal Current; POC: Peru Oceanic Current; PCC: Peru Coastal Current. Subsurface currents. EUC: Equatorial Undercurrent; pSSCC: primary (northern branch) Southern Subsurface Countercurrent; sSSCC: secondary (southern branch) Southern Subsurface Countercurrent; PCCC: Peru-Chile Countercurrent; PCUC: Peru-Chile Undercurrent. c) Map of the eastern South Pacific with schematic arrows indicative of the patterns of motion of the dominant water masses. The lines sketch the dominant currents (HC: Humboldt Current, PCCC: Peru-Chile Counter Current, PUC: Poleward Undercurrent, PCC: Peru-Chile Coastal Current, CHC: Cape Horn Current, SEC: South Equatorial Current, and WWD: West Wind Drift), the red/purple lines indicating waters of tropical/subpolar origin. The dashed lines correspond to intermediate waters. The solid and dashed lines are suggestive of the main direction of propagation of the upper and intermediate waters. Figures a) and b) were taken from Chaigneau *et al.* (2013). Figure c) was taken from Llanillo *et al.* (2013).

### **1.3. Environmental gradients shaping marine biodiversity**

Biodiversity patterns are not randomly distributed but are shaped by a variety of environmental and ecological drivers operating across spatial and temporal scales. Factors such as temperature, productivity, oxygen availability, habitat heterogeneity and disturbance have been widely recognized as major determinants of species distribution and community structure (Gaston, 2000; Huston, 1999; Tittensor *et al.*, 2010). In addition, species richness in marine ecosystems appears to be shaped by the interaction between species' environmental tolerances and both seasonal and interannual variability in environmental conditions (Rombouts *et al.*, 2012; Valdés & Moral, 1998; Williams, 1993).

In marine ecosystems, biodiversity patterns often show substantial spatial variability and do not always follow simple or universal gradients. Classical macroecological paradigms, such as the latitudinal diversity gradient, have been widely documented across terrestrial ecosystems, yet marine systems frequently exhibit more complex and region-dependent patterns (Danovaro, 2024; Piacenza *et al.*, 2015). These deviations arise because marine biodiversity is influenced by the interaction of multiple environmental drivers, including temperature, food availability, habitat heterogeneity and oceanographic processes, which vary across spatial and temporal scales (Danovaro, 2024). Furthermore, because marine ecosystems operate in a three-dimensional environment, depth plays a fundamental role in shaping species distributions and biodiversity patterns,

particularly for organisms inhabiting subsurface and benthic habitats (Owens & Rahbek, 2025).

In the Southeastern Pacific, environmental gradients associated with the Humboldt Current System strongly influence benthic community structure. One of the most prominent features of this region is the presence of an extensive Oxygen Minimum Zone (OMZ), which creates pronounced gradients in oxygen availability along the continental margin (Montecino & Lange, 2009; Thiel, 2007). Low oxygen conditions have been shown to negatively affect macrobenthic diversity and species richness on the continental shelf and slope off Chile, often leading to simplified communities dominated by tolerant taxa (Palma *et al.*, 2005). Studies conducted along bathymetric gradients have also demonstrated that benthic diversity patterns vary with depth and oxygen availability, with lower diversity typically observed within the OMZ and higher diversity below it (Gómez *et al.*, 2024). In addition to oxygen, food supply derived from high primary productivity and the physical characteristics of sediments, such as grain size and organic matter content, have been identified as important drivers shaping macrobenthic biodiversity and distribution in this region (Soto *et al.*, 2017).

#### **1.4. Foraminifera as a study model**

Foraminifera are predominantly single-celled microorganisms with tests that inhabit mainly marine environments (Murray, 2006; Sen Gupta, 2002). Globally, more than 1000 species of benthic foraminifera have been described (Debennay, 2012; Loeblich & Tappan, 1988). The classification of this group is largely based on the composition and structure of the test wall (Blackmon & Todd, 1959). Three

main types are recognized: 1) **Tectinaceous species**, characterized by an organic or proteinaceous wall, lacking a hard test. 2) **Agglutinated species**, with a proteinaceous or mineralized wall; they construct their tests by binding sediment particles together. 3) **Calcareous species**, with mono- or bilamellar walls, forming their tests from precipitated calcite, either with high or low Mg content (Kaminski, 2014; Lipps, 1973; Murray, 2007).

Benthic foraminifera constitute approximately 50% of the meiofauna in marine sediments (Zarghami *et al.*, 2019). They exhibit high microhabitats specificity and are widely recognized as potential environmental bioindicators (Jima *et al.*, 2022; Murray, 2001; Sen Gupta, 2002), as their tests can record evidence of environmental stress. Their short reproductive cycles and rapid growth make their community structure sensitive to environmental fluctuations (Scott *et al.*, 2005; Yanko *et al.*, 1994). Due to their sensitivity to environmental change and their excellent preservation in marine sediments, benthic foraminifera have become one of the most widely used biological indicators in environmental and paleoceanographic studies (Mackensen, 2008). Furthermore, benthic foraminifera play a key role in biogeochemical cycles, since they are capable of absorbing algal carbon, removing nitrogen through respiration and contributing to denitrification processes (Koho & Piña-Ochoa, 2012; Moodley *et al.*, 2000). Consequently, benthic foraminifera represent an important functional component of carbon and nitrogen cycling in marine ecosystems.

In the Southeastern Pacific, the taxonomic diversity of benthic foraminifera has been extensively investigated over the past decades (Diz *et al.*, 2023; Fernández & Zapata, 2010; Figueroa *et al.*, 2005; Figueroa *et al.*, 2006; Mayor Pastor, 1998). These organisms have been widely used in paleoceanography studies to infer temporal variations in sediment redox conditions (Cardich *et al.*, 2019; Erdem *et al.*, 2020); as environmental bioindicators (Tavera Martínez *et al.*, 2023; Tavera Martínez *et al.*, 2022), and for their potential role in biogeochemical processes (Cardich *et al.*, 2012; Glock *et al.*, 2013; Merma-Mora, 2016). Oxygen availability is considered one of the most important environmental factors controlling the distribution and diversity of benthic foraminiferal communities, as variations in oxygen concentration can strongly influence species composition, abundance and microhabitat distribution within marine sediments (Gooday, 2003; Jorissen *et al.*, 2007). Several species exhibit physiological adaptations that allow them to tolerate hypoxic or even anoxic conditions, including the ability to store and respire nitrate when oxygen availability is limited (Glock *et al.*, 2013; Piña-Ochoa *et al.*, 2010). The latter aspect is particularly relevant in the Southeastern Pacific since its extensive Oxygen Minimum Zone (OMZ). In such environments, benthic communities commonly exhibit reduced diversity and are dominated by taxa capable of tolerating hypoxic conditions (Gooday, 2003; Levin *et al.*, 2002). Benthic foraminifera displays both community-level and physiological responses to varying hypoxic gradients (Cardich, 2012; Cardich *et al.*, 2012). At approximately 12°S, species such as *Bolivina costata*, *Nonionella auris* and *Virgulinitella fragilis* are characteristics of anoxic, organic-rich coastal sediments

(Cardich, 2012). Similarly, at 13°S, hypoxic conditions are associated with low diversity in benthic foraminiferal assemblages, which are strongly dominated by hyaline calcareous species such as *Buliminella elegantissima* and *B. costata*. These species are considered indicators of environmental stress because they exhibit high tolerance to low interstitial pH and tend to dominate under conditions that promote carbonate dissolution within the sediment (Merma-Mora, 2016).

Morphological characteristics, such as pore size and density, have been widely employed as paleoceanographic proxies to reconstruct past oxygenation and redox conditions (S Garrido *et al.*, 2025). These features are often associated with physiological adaptations to low-oxygen environments. Pore characteristics have been linked to the capacity of some benthic foraminifera to store and respire nitrate, allowing them to perform denitrification under hypoxic conditions (Glock *et al.*, 2012; Rathburn *et al.*, 2018; Richirt *et al.*, 2019). Evidence from the Oxygen Minimum Zone off Peru indicates that foraminiferal denitrification can account for up to ~38–50% of the total benthic nitrate loss between 79 and 248 m depth (Glock *et al.*, 2013; Piña-Ochoa *et al.*, 2010). Additionally, test morphology has been linked to the microhabitat of benthic foraminifera. Asymmetrical tests with pores concentrated on one side are typically associated with epifaunal species attached to hard substrates, whereas symmetrical tests with bilateral pore distribution are more characteristic of infaunal taxa living within the sediment (Dubicka *et al.*, 2015).

## 1.5. Functional Diversity

Community diversity studies have been extensively conducted to evaluate the impact of biodiversity loss on ecosystem functioning. The decline in biodiversity has been shown to negatively affect key ecosystem processes, including productivity, trophic interactions, and biogeochemical cycles (Loreau *et al.*, 2001). However, evidence of a consistent and universal relationship between species richness and ecosystem functioning remains elusive, as similar levels of ecosystem performance can be maintained by communities with different taxonomic compositions (Cadotte *et al.*, 2011; Cardinale *et al.*, 2000).

In this context, functional diversity has emerged as a key framework to better understand how biodiversity influences ecosystem processes. Functional diversity focuses on the characteristics of species that influence their ecological roles rather than solely on the number of species present (Cadotte *et al.*, 2011; Díaz *et al.*, 2007; Tilman *et al.*, 1997). By incorporating species' functional traits, this approach provides a more mechanistic understanding of how communities contribute to ecosystem functioning.

The concept of functional diversity has evolved over time and has undergone several conceptual refinements (Figure 2), integrating both the effects of species on ecosystem functioning and their responses to environmental change (Hooper *et al.*, 2005; Laureto *et al.*, 2015; Loreau *et al.*, 2001). In this study, we follow the framework proposed by Loreau *et al.* (2001), where ecosystem functioning is influenced by the diversity and distribution of functional traits within communities. This framework is commonly quantified using complementary indices that capture

different aspects of functional structure, such as the volume of functional space occupied by species (FRic) and the regularity of species distribution within that space (FEve) (Laliberte & Legendre, 2010; Vileger *et al.*, 2008). A crucial step in this framework is the selection and assessment of functional traits. Functional traits are defined as morphological, physiological, or ecological characteristics that determine how organisms interact with their environment and influence ecosystem processes (Francisco & de la Cueva, 2017; Kaminsky *et al.*, 2018; Violle *et al.*, 2007). By quantifying these traits, functional diversity approaches provide insights into community assembly processes, species responses to environmental gradients, and the mechanisms underlying ecosystem functioning. Functional diversity therefore provides a mechanistic framework linking species traits, environmental gradients, and ecosystem functioning. Applying this framework to benthic ecosystems allows the evaluation of how environmental variability shapes functional trait distributions and ecological strategies within benthic communities.

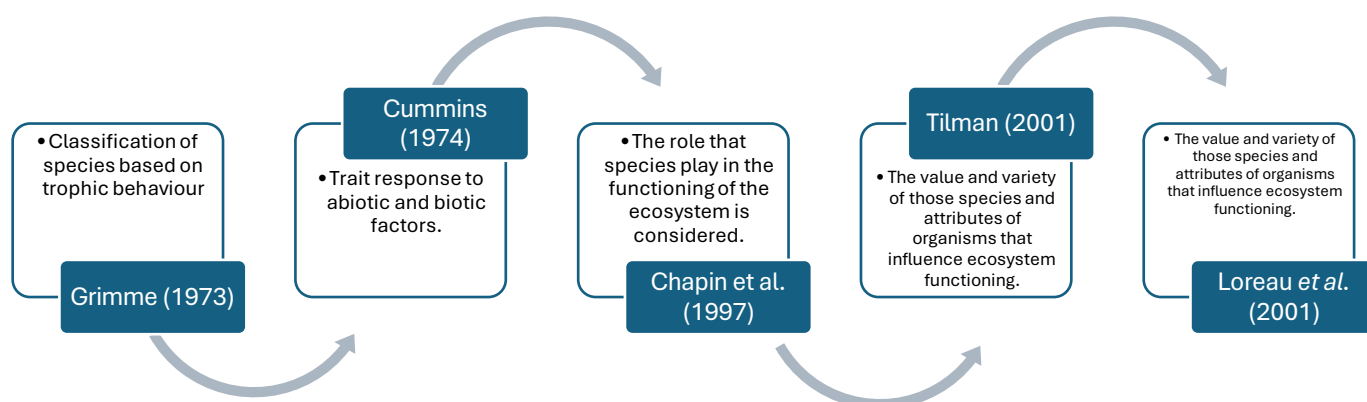


Figure 2. Concepts and definitions in the last decades of functional diversity.

## 1.6. Functional Diversity in the Benthic Ecosystem

The study of functional diversity in benthic ecosystems is essential for understanding how the distribution of functional traits relates to environmental variables, thereby providing insights into biodiversity-ecosystem functioning relationships (Bremner *et al.*, 2006; Kaminsky *et al.*, 2018). Functional traits commonly assessed in benthic organisms include morphological characteristics (*e.g.* body size, mobility), habitat preferences, feeding strategies (*e.g.* feeding mode or prey type), and life-history traits such as larval development or longevity (Brandt *et al.*, 2024; Ghodrati Shojaei *et al.*, 2015; Weigel *et al.*, 2016).

Species distribution in benthic ecosystems is strongly influenced by ecological niche differentiation along environmental gradients, which ultimately shapes the distribution of functional traits within communities (D'Andrea & Ostling, 2016; Kearney *et al.*, 2010). The ecological niche describes the range of environmental conditions that allow a species to persist in each habitat (Kearney, 2006). Differences in niche position and niche breadth can determine patterns of species occurrence, specialization, and tolerance to environmental variability (Qi *et al.*, 2025). Functional traits therefore reflect the ecological strategies through which species occupy and respond to these environmental conditions (Kearney *et al.*, 2010; Włodarska-Kowalczyk *et al.*, 2019).

In the case of foraminifera, several morphological traits have been linked to ecological functions. For example, motility has been suggested to influence sediment mixing and small-scale bioturbation (Deldicq *et al.*, 2020). However, the contribution of foraminifera to sediment reworking and bioturbation has often been underestimated in benthic ecosystem studies.

In addition to individual morphological traits, broader functional classifications have been proposed for foraminifera. One widely used approach is the classification of species into “morphogroups”, based on similarities in test morphology, which has been applied to evaluate stratigraphic and paleoenvironmental variations (Reolid & Martínez-Ruiz, 2013; Reolid *et al.*, 2008). Morphogroups integrate several biological characteristics, including test shape, inferred feeding strategies, and microhabitat preferences, under the assumption that test morphology reflects ecological strategies (Caratelli & Archuby, 2023; Corliss & Chen, 1988; Murray *et al.*, 2011). Although this assumption is not universally accepted (Linke & Lutze, 1993), morphogroup approaches have been successfully used to reconstruct environmental conditions in the geological record, including variations in oxygen availability and nutrient supply (Khare *et al.*, 2017; Reolid & Martínez-Ruiz, 2013; Saalim *et al.*, 2019).

Environmental factors such as oxygen concentration, organic matter availability, salinity, and depth strongly influence both the taxonomic composition and morphological traits of foraminiferal communities (Bouchet *et al.*, 2012; Bouchet *et al.*, 2018). For example, oxygen availability can be reflected in morphological

attributes such as test porosity, wall thickness, and mineral composition (Kaiho, 1994). Habitat preferences of benthic foraminifera have also been conceptualized through the TROX model (Jorissen *et al.*, 1995), which proposes that the vertical distribution of foraminifera between epifaunal (surface dwelling) and infaunal (subsurface dwelling) microhabitats is regulated by the interaction between food availability and oxygen concentration.

## **2. STATEMENT OF THE PROBLEM**

Climate change is expected to adversely affect marine ecosystems through multiple stressors including ocean acidification, increased eutrophication, and expansion of the Oxygen Minimum Zone (Allen *et al.*, 2023; Popova *et al.*, 2016).

Foraminifera, which constitute over 50% of the meiobenthos, are particularly sensitive to environmental factors such as organic matter availability, oxygen concentration, and sediment type (Castillo *et al.*, 2021; Kranner *et al.*, 2022; Miljutin *et al.*, 2012). Their sensitivity, combined with their fossilization potential, makes them indispensable for reconstructions past marine environment (Manasa *et al.*, 2016; S. dos S. de Jesus *et al.*, 2020) and for biomonitoring of potentially toxic elements such as heavy metals (Tavera Martínez *et al.*, 2023; Tavera Martínez *et al.*, 2022).

Although the distribution and taxonomic diversity of benthic foraminifera in the Pacific have been studied in specific regions (Mayor Pastor, 1998; Tavera Martínez *et al.*, 2022; Zapata *et al.*, 1995), comprehensive studies covering multiple surface sediment

stations across a latitudinal broad latitudinal gradient remain scarce (Diz *et al.*, 2023). In particular, studies evaluating the functional diversity of benthic foraminiferal communities in the Southeastern Pacific are scarce. Most previous research in this region has focused primarily on taxonomic diversity or paleoenvironmental applications (Cardich *et al.*, 2012; Erdem *et al.*, 2020; Mallon *et al.*, 2011).

Functional diversity analyses integrate information on species occurrence and abundance with their functional traits, providing insights into how communities respond to environmental gradients and how ecosystem functioning may vary across space (Laureto *et al.*, 2015; Legras *et al.*, 2020). Because benthic foraminiferal species exhibit distinct ecological strategies and physiological adaptations, the analysis of their functional traits provides a valuable framework to understand how environmental variability shapes community structure and ecosystem functioning in marine environments.

Assessing the functional diversity of benthic foraminifera in the Southeastern Pacific will provide novel insights into ecosystem functioning and reveal whether patterns of functional diversity differ from those inferred from taxonomic diversity alone. This study represents one of the first attempts to characterize the functional diversity of benthic foraminiferal communities across this region using ecological and biological traits such as test wall-type, morphology, feeding strategies, and habitat preferences. In addition, the study examines the environmental variables that influence these traits,

contributing to a better understanding of how benthic ecosystems respond to environmental variability.

### **3. RESEARCH QUESTION**

What environmental variables modulate functional and taxonomic diversity in the Southeastern Pacific?

### **4. HYPOTHESIS**

H1: In the Southeastern Pacific Ocean, zones with the highest diversity of living benthic foraminiferal species have a high number of morphological and ecological traits, resulting in the most functional diversity.

H2: The main environmental factors influencing the distribution of functional traits are sediment geochemistry and bottom water oxygen levels.

### **5. GENERAL OBJECTIVE**

To evaluate the functional and taxonomic diversity of living benthic foraminifera, and analyze their relationship with environmental variables in the Southeastern Pacific Ocean

#### **5.1. Specific Objectives**

- 1) To characterize the functional traits of benthic foraminifera to identify their ecological roles at the spatial scale.

- 2) To determine the environmental variables that modulate the functional and taxonomic diversity of benthic foraminifera in the Southeastern Pacific.

## 6. MATERIALS AND METHODS

### 6.1. Data collection

#### 6.1.1. Study area

This study focusses on the western South American continental margin extending from Ecuador (1°S) to the South off Chile (50°S) and spanning water depths from 79 to 4500 m across the shelf and slope, which are influenced by different water masses (Figure 1). For the purposes of this study, the South Pacific is split between the Southeastern Pacific (SEP) and the Equatorial Southeastern Pacific (ESEP) regions. In the case of the ESEP, the analysis is based on three sampling stations located off the coast of Ecuador, and this region is treated as being representative of the equatorial sector of the Southeastern Pacific, rather than as a comprehensive characterization of the province.

#### 6.1.2. Environmental variables

Bottom-water environmental data corresponding to each the sampling stations were utilized compiled from Krahnemann (2012), Krahnemann (2018) and Reyes-Macaya *et al.* (2022) The environmental variables from the compiled dataset (Susaya-Vicente *et al.*, 2025) considered here include temperature, salinity, dissolved oxygen, phosphate, nitrate, nitrite, and silicate. Original hydrological data were interpolated using DIVA (Data-Interpolated Variational Analysis) and the non-outlier Weighted Average Interpolation gridding method

implemented in Ocean Data View (ODV, R. Schlitzer, 2021) View, <https://odv.awi.de/>, 2021), applying maximum grid scales of 30:X and 20:Y nautical per mile.

To match environmental data with sampling locations, a custom Python script was developed to compute the nearest distance neighbour in latitude, longitude, and water depth ensuring site-specific (Python code is available in Reyes-Macaya *et al.* (2024)).

In addition, a sediment geochemistry dataset was compiled, including biogenic opal, total organic carbon, total nitrogen, the C/N ratio and CaCO<sub>3</sub> content, obtained from previously published studies (Sebastián Garrido *et al.*, 2025; Mollier-Vogel *et al.*, 2012; Prahel *et al.*, 2006; Romero & Hebbeln, 2003; Sommer & Dale, 2019) (Appendix 1).

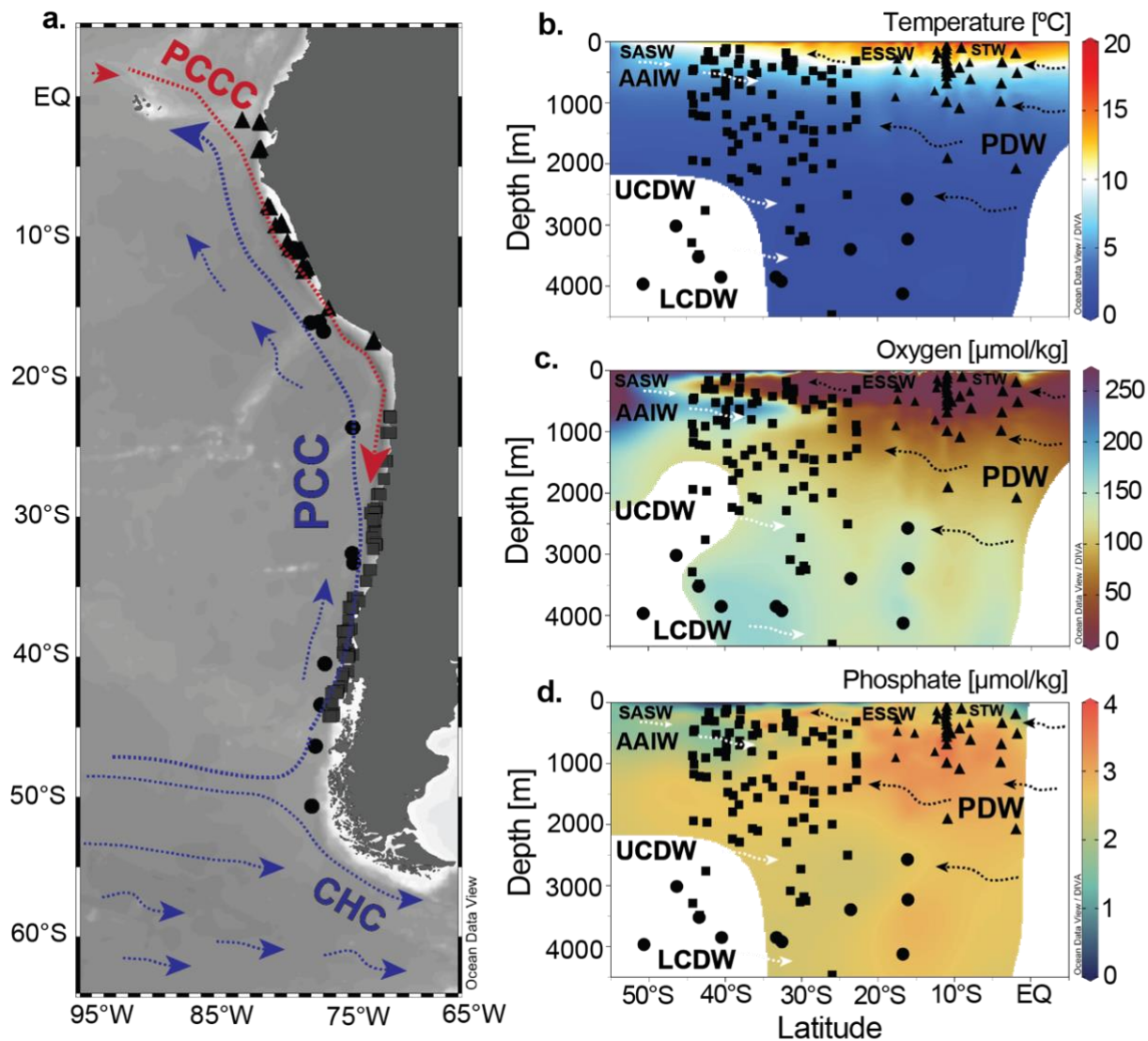


Figure 3. Sampling stations along the western South American margin. (a) Sampling locations: triangles = coasts of Ecuador and Peru (expedition M77 of the R/V Meteor, Mallon (2011)); grey squares = north and south-central Chile margin (PUCK expedition, Hebbeln *et al.* (2001)); dots = Nazca Ridge and Chile Rise (R/V Roger Revelle RR9702A). (b-d) Depth distribution of (b) temperature, (c) oxygen, and (d) phosphate concentrations of the main water masses influencing the sampling stations in the Southeastern Pacific. Currents: PCCC: Peru-Chile Countercurrent, PCC: Peru-Chile Current, CHC: Cape Horn Current. Water masses: ESSW: Equatorial Subsurface Water, STW: Subtropical Water, SASW: Subantarctic Surface Water, AAIW: Antarctic Intermediate Water, PDW: Pacific Deep Water, UCDW: Upper Circumpolar Deep Water, LCDW: Lower Circumpolar Deep Water.

### 6.1.3. Surface sediments

Benthic foraminiferal data were obtained from 35 surface sediment samples collected during legs 1 and 2 of the R/V Meteor cruise M77, conducted

between October 22 and December 22, 2008 (Pfannkuche, 2011). Samples were retrieved using a video-guide along the margin between 1°S (Ecuador) and 17°S (Peru), at depths ranging from 79 to 2092 m. The multicorer tubes had an inner diameter of 10 cm, and one tube with an even surface was selected from each deployment for analysis. Sampling was carried out along the continental margin between 1°S (Ecuador) and 17°S (Peru) at water depths ranging from 79 to 2,092 m. Sediments were retrieved using a video-guided multicorer with a 10 cm inner diameter. From each deployment, the core with the most undisturbed sediment surface was selected for analysis. Surface sediments were sliced into intervals of 2, 3, 5, or 10 mm intervals following standard procedures. Benthic foraminiferal assemblages, from the 63-2000  $\mu\text{m}$  size fraction, following of (Mallon, 2011).

Additionally, 132 sediment samples were collected during leg 2 of the R/V Sonne cruise SO156 (PUCK expedition), conducted between April 24 and May 14, 2001, from 22°S (Iquique) to 44°S (Guaitecas Islands), at depths between 110 to 4470 m. As with R/V Meteor, the samples were collected, with a multicorer (10 cm inner diameter), and the selected cores were sliced into 1 cm-thick intervals (Hebbeln *et al.*, 2001). Benthic foraminiferal assemblages were analyzed. from >150  $\mu\text{m}$  size fraction

Finally, 10 surface sediment samples were obtained during the R/V Roger Revelle cruise RR9702A (GENE03RR), conducted between February 25 and March 16, 1997, along Chilean continental margin between 16°S and 50°S,

at depths ranging from 2575 to 4124 m. Data from this expedition are available at (<https://www.pangaea.de/expeditions/events/RR9702A>).

#### 6.1.4. Sample processing

Sediment subsamples were preserved in a solution of 2 g L<sup>-1</sup> Rose Bengal prepared in 96% ethanol and stored at 4°C in dark conditions until laboratory processing. This procedure is commonly used to distinguish living foraminifera at the time of collection

Foraminifera Specimens showing bright pink staining in 50% of their chambers were classified as living individuals (Fontanier *et al.*, 2016; McCorkle *et al.*, 1990; Murray & Bowser, 2000; Rathburn & Corliss, 1994). The faunal analysis was performed on the >150 µm. Size fractions to ensure comparability among sampling sites.

## 6.2. Taxonomic Diversity

After sieving and drying, a comprehensive taxonomic analysis of all samples was conducted. Separation, census, and identification of living benthic foraminifera were performed under a stereomicroscope. Samples from cruise RR9702A (16°S - 50°S) were analyzed at the Woods Hole Oceanographic Institution (WHOI, USA), whereas those from cruise M77 (1°S - 17°S) were analyzed at GEOMAR, Helmholtz Centre for Ocean Research, Kiel, Germany (Mallon, 2011), and the samples from the PUCK cruise (22°S and 44°S) were processed at the Paleoceanography Laboratory of the Lyell Centre, Heriot-Watt University, Edinburgh, Scotland.

Species determination was performed using standard taxonomic references and identifications keys including Cushman & Wickenden (1929), Bandy & Rodolfo (1968), Brady (1884), Ellis & Messina catalogue (1942-2006), Figueroa et al. (2005, 2006), Ingle et al. (1980), Loeblich & Tappan (1988), Resig (1981, 1990), Smith (1963), Uchio (1960), Whittaker (1988), Zapata et al. (1995,1999).

To ensure taxonomic consistency and avoid overestimating species richness due to synonyms, the current generic classification, valid species names, and potential synonymies were verified using the World Register of Marine Species (WoRMS database <https://www.marinespecies.org/foraminifera/>).

### 6.3. Estimation of the taxonomic diversity of benthic foraminifera

After taxonomic identification, approximately 100 individuals per sample were counted to ensure statistical significance (Murray, 2007; Patterson & Fishbein 1989; Schönfeld *et al.*, 2012). Samples containing fewer than five individuals were excluded from subsequent analyses.

Given the heterogeneous sampling techniques and faunal census procedures along the western South American Margin, the population density of living foraminifera was standardized to the number of individuals per 50cm<sup>3</sup> of near-surface sediment, acknowledging that the top 1 cm was not consistently sampled across all sites (Schönfeld *et al.*, 2012).

Relative abundances were calculated to determine the percentage of contribution of each species to the total assemblage. Species present in more than 20% of the total samples were classified as dominant taxa.

Finally, community diversity indexes (species richness, Shannon–Wiener diversity index, and Pielou’s evenness) were calculated using the PASTv4 software (Hammer *et al.*, 2001). Consequently, the iNEXT (iNterpolation/EXTrapolation) package using R v4.5 (Chao & Hsieh, 2016) was used to calculate the rarefaction curves. This technique allows direct comparison of species richness across samples with unequal abundances, thus enabling the evaluation of whether sampling effort was sufficient. When the rarefaction curve approaches an

asymptote, it will indicate that the sampling effort adequately captured the diversity (Gotelli & Colwell, 2011).

## **6.4. Estimation of the functional diversity of benthic foraminifera**

### 6.4.1. Functional traits matrix

A total of five functional traits, subdivided into 25 categories, were selected to characterize the ecological roles of benthic foraminifera within the ecosystem (Table 1). According to (Palacio *et al.*, 2022), a minimum of three traits is recommended for a reliable assessment of functional diversity without substantial loss. The selected traits included: 1) Test wall type, referring to the composition of the shell. 2) Morphology, corresponding to the overall shape or coiling pattern of the test. 3) Microhabitat, describing the lifestyle position of the species within the benthic environment. 4) Oxygen condition, reflecting the tolerance or requirement of species to different oxygenation levels. 5) Trophic behavior, indicating the feeding strategy of the species. (Kaiho, 1994; Linke & Lutze, 1993; Lipps & Valentine, 1970; Murray *et al.*, 2011).

Table 1. Functional traits and associated categories of benthic foraminifera considered in this study, adapted from Guy-Haim and Bouchet (2025).

<b>Traits</b>	<b>Category</b>	<b>Function and processes in the ecosystem</b>
Wall Type	Agglutinated	Related to the sediment grain size, carbonate production, gas exchange
	Porcelaneous	
	Hyaline	
Morphology	Planispiral	Foraging mode, migration in the sediment.
	Trochospiral	
	Uniserial	
	Biserial	
	Triserial	
	Miliolide	
	Spheric	
Microhabitat	Epifaunal, 0 cm	Role in vertical sediment flow, i.e. bioturbation, biogeochemistry, fluxes and the benthic boundary
	Shallow infaunal, 0 – 0.5 cm	
	Intermedia infaunal, 0.5 – 1.5 cm	
Oxygen preference	Oxic, >50 $\mu\text{mol/l}$	Respiration, denitrification ability, contribution to the nitrogen cycle
	Hypoxic, < 50 – 5 $\mu\text{mol/l}$	
	Anoxic, < 5 $\mu\text{mol/l}$	
Trophic behavior	Phytodetritivore	Food acquisition, metabolism, endobionts, growth requirements
	Detritivore	
	Bacterivore	
	Herbivore	
	Omnivore	
	Carnivore	

Traits of information were compiled from published literature, preferentially at the species level. When species-specific data were unavailable, trait values were assigned from the closest relative within the same taxonomic group. Classification criteria followed: test wall type and chamber arrangement after Loeblich & Tappan (1988); microhabitat depth ranges as proposed by Corliss and van Weering (1993) and Fontanier *et al.* (2002); and oxygenation requirements according to Kranner *et al.* (2022) and Mallon *et al.* (2011).

After compiling the traits data, a fuzzy coding approach was applied to assign affinity value ranging from 0 (no affinity) to 1 (high affinity) for each species within each trait category (Chevene *et al.*, 1994). This method allows species to exhibit partial membership across multiple trait modalities, thereby capturing ecological plasticity. To ensure comparability across traits and species, the scores assigned to each species were subsequently standardized so that the sum of affinities equaled one

## 6.5. Functional diversity analysis

Functional diversity indexes (FRic, FDiv, FEve, FDis, Rao's Q) and community-weighted means (CWM) were calculated using the fuzzy coded functional traits matrix together with the relative abundance matrix of benthic foraminifera, employing the *dbFD* function in the "FD" R package (Laliberté *et al.*, 2014).

- **Functional richness (FRic)** quantifies the proportion of the functional space filled by the species within community, regardless of abundance, and is computed as the volume of the minimum convex hull that encloses all species in a multidimensional traits space (Villegger *et al.*, 2008).
- **Functional divergence (FDiv)** describes how abundance is distributed within the occupied functional space, with higher values indicating greater divergence from the community centroid (Mason *et al.*, 2005; Villegger *et al.*, 2008).
- **Functional evenness (FEve)** measures the regularity of species' abundance across the traitspace, with highervalues indicating greater divergence from the community (Villegger *et al.*, 2008).
- **Functional dispersion (FDis)** corresponds to the abundance-weighted mean distance of species from the community in centroid in trait space ((Laliberte & Legendre, 2010).
- **Rao's quadratic entropy (Rao's Q)** integrates both species abundances with functional distances (or dissimilarities) between all pairs of species (Botta-Dukát, 2005).

Before computing these indices, the density matrix (ind/50 cm<sup>3</sup>) was transformed into relative abundances. This step avoids overrepresentation of stations with high densities and ensures that functional indices reflect proportional contributions rather than absolute abundances (Mouillot, Graham, *et al.*, 2013).

## 6.6. Statistical analysis

### 6.6.1. Correlation between indices

To examine the relationship between functional and taxonomic diversity, a Spearman correlation analysis was conducted. Taxonomic diversity was quantified using Shannon index, Fisher's  $\alpha$ , Simpson's dominance, Pielou's evenness, abundance, and species richness, while functional diversity was represented by Rao's Q, FRic, FEve, FDis, and FDiv. This approach tests whether higher taxonomic diversity correlates with increased functional diversity, or if functional redundancy is present.

### 6.6.2. Effect of the environmental variables in the diversity indexes

Environmental variables 6.6.1 were standardized using the 'z-score' method (scale function in R), which normalizes values by subtracting the mean and dividing the standard deviation (Becker *et al.*, 1988). To reduce multicollinearity, pairwise Spearman correlations and variance inflation factors (VIF) were evaluated. Variables with  $|\rho| > 0.7$  or  $VIF > 10$  were excluded from further analyses, ensuring model parsimony and minimizing overfitting.

The influence of the selected environmental predictors on taxonomic and functional diversity indices was assessed using Generalized Additive Models (GAMs), fitted with the mgcv R package (Wood & Wood, 2015). Distributions for each index were chosen according to their statistical properties and best model fit (Table 2). To account for spatial structure, two-dimensional smoothers of latitude and longitude were included using a Gaussian Process base function (bs = "gp").

Additionally, the same environmental variables were tested against the functional traits of benthic foraminifera to evaluate how environmental gradients modulate specific ecological roles within communities. The final dataset used for the GAM and OMI analyses is provided in Appendix 2.

Table 2. Generalized Additive Models (GAMs) applied, including distribution of families, basis functions, and smoothing parameters.

Response Variable	Predictor Variables	Family
Shannon	$s(\text{Temp}) + s(\text{Salinity}) + s(\text{Depth}) + s(\text{Oxygen}) + s(\text{Phosphate}) + s(\text{Nitrite}) + s(\text{Silicate}) + s(\text{TOC}) + s(\text{TN}) + s(\text{Opal}) + s(\text{CaCO}_3) + s(\text{C/N}) + s(\delta^{15}\text{N}) + s(\text{Longitude, Latitude, bs} = \text{"gp"})$	Gaussian
Taxa		Tweedie
Simpson		Quasibinomial
Pielou		Gaussian
Abundance		Gaussian
FRic		Tweedie
FDis		Gaussian
FDiv		Beta
FEve		Beta
RaoQ		Gaussian

### 6.6.3. OMI analysis

The influence of environmental variables on the functional traits of benthic foraminifera was assessed using an Outlying Mean Index (OMI) analysis (Dolédec *et al.*, 2000), implemented with the “niche” function in the ‘ade4’ package (Dray & Dufour, 2007). The analysis was based on a compiled dataset consisting of 1) the standardized abundance of the benthic foraminifera per station, 2) environmental variables at each station, and 3) the functional traits for each species.

It is important to note that CaCO<sub>3</sub> concentration was excluded this analysis, as its inclusion would have reduced the dataset by more than half (n = 38) (Appendix 2). In OMI analysis, the realized niche of species is represented in a multidimensional space defined by their distribution and trait composition. Three key parameters are obtained:

- 1) OMI (marginality index): measured the distance between the mean environmental conditions of a species and the mean environmental conditions of the study area.
  
- 2) Tolerance (Tol): quantifies niche breadth, representing the ranges of environmental conditions under which a species occurs. High values indicate generalist species, while low values denote specialists restricted to narrow environmental ranges.

3) Residual tolerance (RTol): represents the proportion of variance not explained by the analysis (Dolédec *et al.*, 2000).

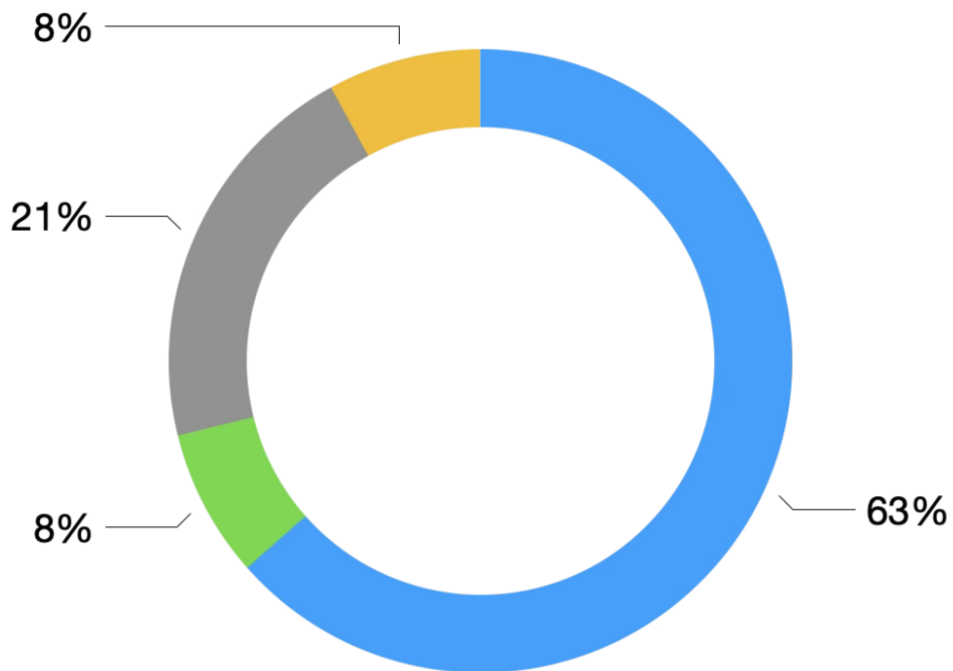
## 7. RESULTS

### 7.1. Taxonomic diversity of benthic foraminifera

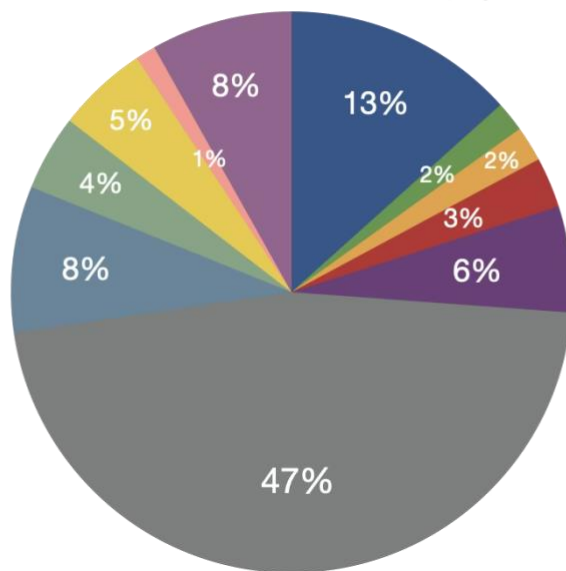
A total of 365 benthic foraminiferal species were recorded along the western South American Margin (Appendix 3). These species belong to the Classes Globothalamea, Nodosariata, Tubothalamea and Monothalamea, encompassing 11 orders. The most frequent orders were Rotalida (47%), Lituolida (13%), Polimorphynida (8%), and Astrorhizida (8%) (Figure 4).

Additionally, several species recorded in this study represent new occurrences for the Southeastern Pacific (SEP), including *Astrorhizoides cornutus*, *Crithionina pisum*, *Galwayella trigonoornata*, and *Liebusella goesi* (Figure S2).

● Globobulimina ● Tubobulimina ● Nodosariata ● Monobulimina



● Lituolida ● Loftusiida ● Spirillinida ● Vaginulinida  
● Miliolida ● Rotaliida ● Astrorhizida ● Textulariida  
● Nodosariida ● Robertinida ● Polimorphynida



**Figure 4.** Percentage contribution of benthic foraminifera classes and orders recorded for the Southeastern Pacific Ocean.

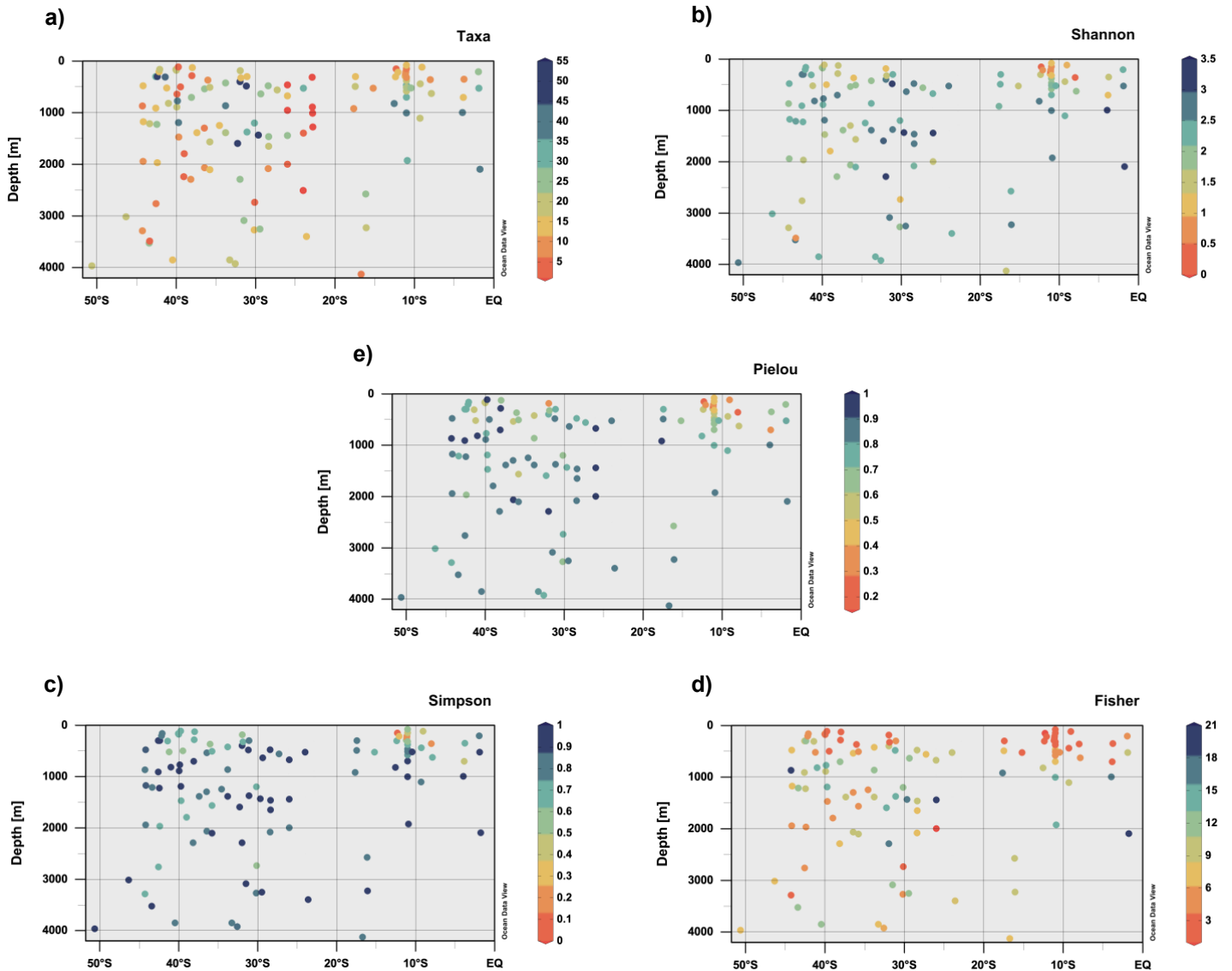
The species richness and indices of Shannon-Wiener diversity, Simpson's Dominance, Fisher's- $\alpha$  and Pielou's evenness were estimated (Figure 5), along with the percentage abundance of benthic foraminifera species (Figure 6). Species richness exhibited marked variability, ranging from 1 to 54 species per sample (Appendix 4). Notably conspicuous peaks in species richness (>40 species) were recorded, at stations located between latitudes 1°–3°S, 29°–32°S, and 39°–42°S, corresponding to depths greater than 900 m, 400–1600 m, and 300–1100 m, respectively (Figure 5a).

Simpson's Dominance value ranged from 0.1–0.96, excluding stations with very low counts (1–3 specimens; Appendix 4), where the index could not be reliably calculated. The values ranged between 0.50 and 0.95 with most of the samples (75%) exhibiting values greater than 0.70 (Appendix 4).

The Shannon index varied from 0 and 3.30 (Appendix 4), with only four stations (Site\_1\_2092, Site\_3\_995, Site\_29\_1432, and Site\_31\_481) recording values above 3. In contrast, stations with index values below 1 were primarily located between latitudes 7° and 12°S and at depths of 100 -250 meters. These stations were directly influenced by the ESSW, characterized by low oxygen concentrations (1 - 2  $\mu\text{mol/kg}$ ) (Figure 5c). The Fisher- $\alpha$  index ranged between 10 and 20 across the Southeastern Pacific (Figure 5d), with higher values typically observed at depths greater than 900 m.

Pielou's equitability was generally high (>0.7) at sites off the coast of Ecuador and at depths >500 m off Peru, with the highest values observed along the Chilean margin (Figure 4e). In contrast, low equitability (0.4–0.1) was recorded between latitudes 3°–

12°S at sites shallower than 400 m (Figure 5e). An exception was the site observed at 3°S and 700 m, where the value was 0.39 (Appendix 4). Along the Chilean coast, between latitudes 31° and 41°S, most stations shallower than 550 m exhibited average equitability values 0.5, except for one station (Site\_31\_183) which recorded a lower value of 0.3.



**Figure 5.** Variation of taxonomic richness along the Southeastern Pacific (SEP). (a) Species richness. (b) Shannon-Wiener Index. (c) Simpson Dominance Index. (d) Fisher's  $\chi^2$  Index. (e) Pielou Evenness.

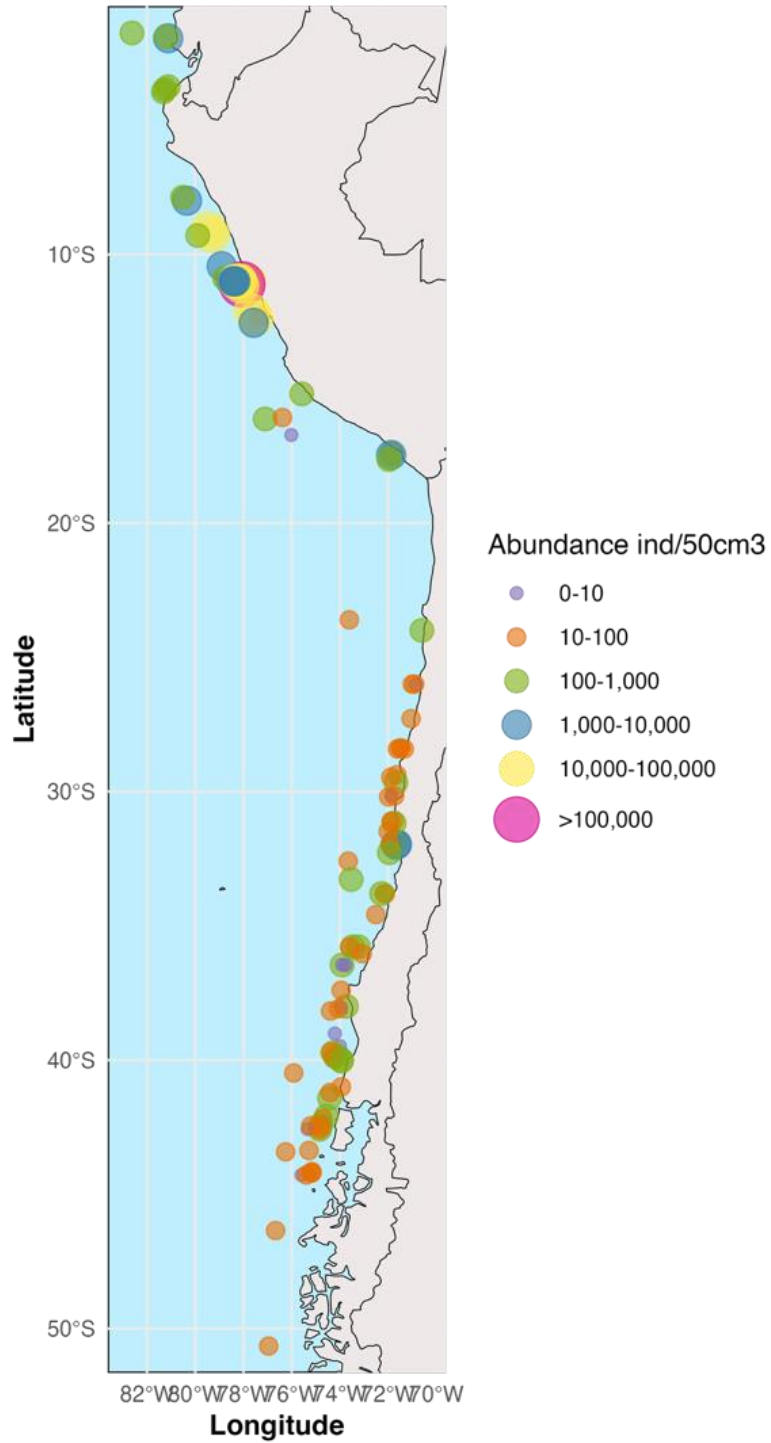
Benthic foraminifera abundance (ind/50 cm<sup>3</sup>) showed a pronounced increase between 1°S and 17°S (Figure 6). Within this latitudinal range, however, the species' richness and diversity were markedly lower at 9°S and 12°S, coinciding with the low-oxygen conditions characteristic of this region. These sites recorded the highest abundances across the study area (>10,000 ind/50 cm<sup>3</sup>), with values ranging from 15,370 to 105,558 ind/50 cm<sup>3</sup>. In contrast, abundances south of 23°S generally did not exceed 1000 ind/50 cm<sup>3</sup>, except at two sites (Site\_31\_183 and Site\_31\_398), which yielded 5580 and 1010 ind/50 cm<sup>3</sup>, respectively (Figure 6; Appendix 5). Of the 77 sites situated between 23°S and 50°S, most (n = 55) contained between 10 and 100 ind/50 cm<sup>3</sup>.

The most abundant taxa across all sites belonged to the genus *Bolivina*, which accounted for 79.54% along the Southeastern Pacific (Table 3). Species present in more than 20% of the samples included: *Bolivina costata*, *B. interjuncta*, *B. plicata*, *B. seminuda*, *B. spissa*, *Cassidulina carinata*, *Cyclammina cancellata*, *Glaphyrammina americana*, *Globobulimina pacifica*, *Hansenisca soldanii*, *Haplophragmoides pussillus*, *Hoeglundina elegans*, *Lobatula wuellestorfi*, *Melonis affinis*, *Nodulina dentaliniformis*, *Nonionoides grateloupii*, *Pullenia bulloides*, *Reophax scorpiurus*, *Stainforthia fusiformis*, *Trifarina angulosa*, *Uvigerina auberiana*, *U. peregrina* and *Valvulineria glabra* (Figure S3–S6). Notable changes were observed in the abundance and relative contribution of *Bolivina seminuda* and *Uvigerina peregrina* across Southeastern Pacific locations (Figure S3–S6). Between latitudes 1° and 17°S, abundances were overwhelmingly dominated by genus *Bolivina* (Figure S3). South of

23°S, although *Bolivina* persisted, its contribution declined markedly, particularly at depths greater than 500 m.

The most abundant taxa across all sites belonged to the genus *Bolivina*, which accounted for 79.54%, primarily driven by the contribution of *B. seminuda* (61.37%) along the SEP (Table 3). Species present in more than 20% of the samples included: *Bolivina costata*, *B. interjuncta*, *B. plicata*, *B. seminuda*, *B. spissa*, *Cassidulina carinata*, *Cyclammina cancellata*, *Glaphyrammina americana*, *Globobulimina pacifica*, *Hansenisca soldanii*, *Haplophragmoides pussillus*, *Hoeglundina elegans*, *Lobatula wuellestorfi*, *Melonis affinis*, *Nodulina dentaliniformis*, *Nonionoides grateloupii*, *Pullenia bulloides*, *Reophax scorpiurus*, *Stainforthia fusiformis*, *Trifarina angulosa*, *Uvigerina auberiana*, *U. peregrina* and *Valvulineria glabra* (Figure S2 – S5). The dominance of *Bolivina* was particularly pronounced within the oxygen-minimum zone (7°–17°S) at neritic and upper bathyal depths (Figure S2, S3). Latitudes 7° and 17°S at neritic (0 – 250 m) and upper bathyal (>250 – 850 m) depths, abundances were overwhelmingly dominated by genus *Bolivina* (Figures S2 – S3). However, this pattern is not showed in lower bathyal (>850 – 2000 m) and abyssal samples (>2000 m), where their contributions are less than 10% and only reach until ~10°S (S3 Figure), except for *B. spissa* (Figure S3 – S4). South of 23°S, although *Bolivina* persisted, its contribution declined markedly, and the genus *Uvigerina* and *T. angulosa* rise contributing ~10% or more per site and their contributions increased at sites > 250 m. The contribution of *C. carinata* is notably high mainly on sites at 250 – 850 m (Figure S3), as well, *H. elegans* and *M. affinis* contribute considerably at stations at the upper bathyal to abyssal zones. Finally, agglutinated species as *G. americana*, *R. scorpiurus* and *N.*

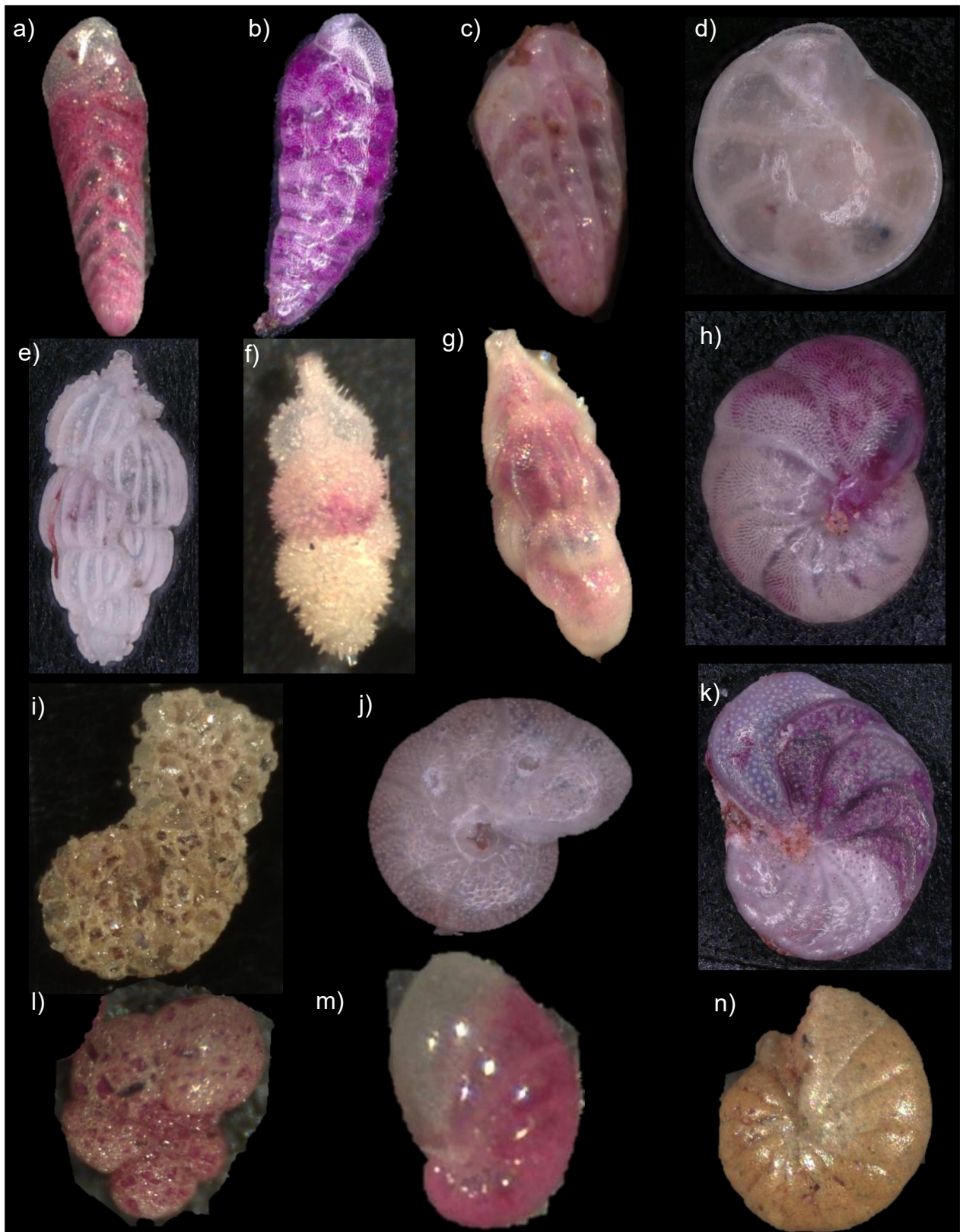
*dentaliniformis* have a contribution mainly in stations > 250 m. In neritic zones *G. americana* contribute ~2% at stations south of 37°S, while *R. scorpiurus* contributions is only notable at stations of 1°S (Figure S2). South of 23°S *G. americana* contribution increases to 5% at upper bathyal zones (Figure S3) as well as *R. scorpiurus*, which contributes 50% at 39°S while *N. dentaliniformis* only contributes 4% at a station at 31°S (Figure S3). In lower bathyal zones, *G. americana* represent a contribution >15% in stations at 26°S and 39°S, as well as *N. dentaliniformis* (~12%) while *R. scorpiurus* is more representative (4 – 28%) in stations between 10° – 32°S (Figure S4). Finally, station in the abyssal zone, *G. americana* has a contribution of 30% only at 36°S, *N. dentaliniformis* shows representativity of 2 – 14% at stations between 1° - 30°S and *R. scorpiurus* continue with a contribution of 1 – 34% along the sites (Figure S5) South of 23°S, a major shift in assemblage composition occurred. The relative contribution of *Bolivina* declined markedly, giving way to more diverse fauna. The genera *Uvigerina* and *Trifarina angulosa* became significant contributors (~10% or more per site) at depths >250 m (Figure S3, S4). Notably, agglutinated species such as *Glaphyrammina americana*, *Reophax scorpiurus*, and *Nodulina dentaliniformis* increased their contribution with depth, becoming dominant constituents in many lower bathyal and abyssal stations (Figs. S4, S5).



**Figure 6.** Spatial distribution of the total abundance of benthic foraminifera (ind/50cm<sup>3</sup>) of living benthic foraminifera along the western South American Margin.

**Table 3.** Species contributing most to the total number of sampling sites based on their abundance (ind/50cm<sup>3</sup>) in the Southeastern Pacific. Ocean.

% Contribution	Species
10.27%	<i>Bolivina costata</i>
2.86%	<i>Bolivina interjuncta</i>
5.04%	<i>Bolivina plicata</i>
61.37%	<i>Bolivina seminuda</i>
1.13%	<i>Buliminella elegantissima tenuis</i>
1.48%	<i>Cancris carmenensis</i>
1.26%	<i>Epistominella obesa</i>
2.66%	<i>Nonionella stella</i>
1.33%	<i>Stainforthia fusiformis</i>
1.54%	<i>Uvigerina peregrina</i>
2.53%	<i>Valvulineria glabra</i>
8.51%	Others



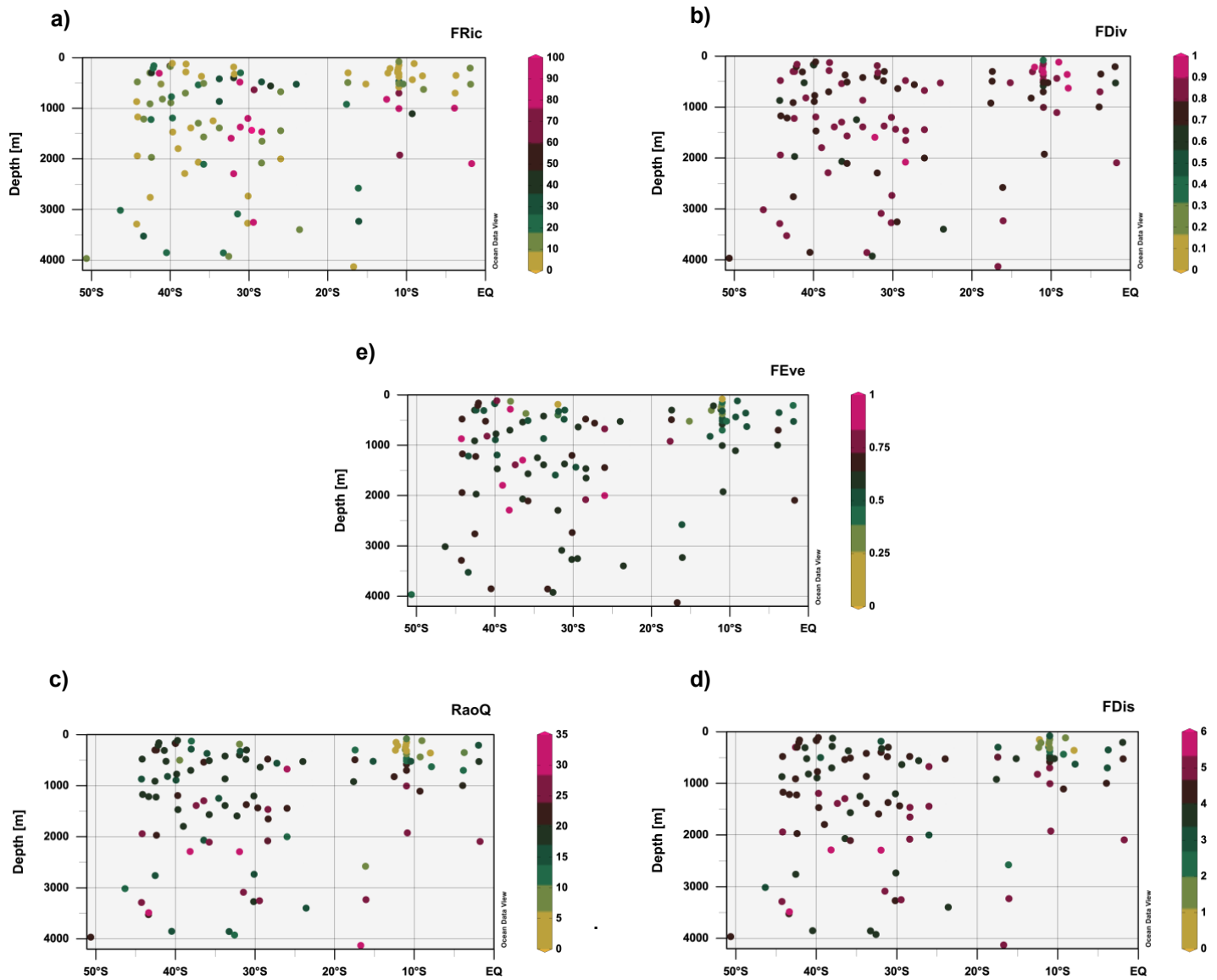
**Figure 7.** Most abundant taxa of living benthic foraminifera across the Southeastern Pacific. a) *Bolivina spissa*. b) *B. seminuda*. c) *B. plicata*. d) *Hoeglundina elegans*. e) *Uvigerina peregrina*. f) *U. auberiana*. g) *Trifarina*

*angulosa*. h) *Valvulineria glabra*. i) *Glaphyrammina americana*. j) *Melonis affinis*. k) *Lobatula wuellestorfii*. l) *Haplophragmoides pussillus*. m) *Nonionoides grateloupii*. n) *Cyclammina cancellata*.

Regarding the rarefaction curves (Figure S3), only a subset of sites reached the asymptote, indicating an optimal sampling effort. These included sites “1\_207”, “3\_350”, “7\_559”, “9\_115”, “9\_437”, belonging to the group 1; “10\_521”, “11\_291”, “11\_317”, “11\_319”, “11\_375”, “11\_495”, “11\_511”, (group 2); “11\_697”, “12\_214”, “12\_302”, “12\_823”, “17\_298”, (group 3); all sites in Group 4 except for “17\_918” and “26\_1996”; “31\_183”, (Group 5); all sites in Group 6; “37\_124” and “39\_174” (group 7); “39\_190”, “40\_165”, “41\_308”,(group 8); “42\_371”, “43\_1212”, “43\_3523”, “44\_1771”, “44\_1939”, (group 9), and , “46\_3014” and “50\_3964”. In contrast, sites where the curves did not stabilize suggest the potential presence of higher, unrecorded diversity.

## 7.2. Assessment of Functional Diversity

Functional richness (FRic) showed low values, with only 11.8% of sites exhibiting values  $>70$  (Figure 7a). In contrast, functional divergence (FDiv) was generally high, ( $>0.7$ ) at 80% of sites (Table S4), being noticeable at sites between  $7^{\circ}\text{S}$  –  $12^{\circ}\text{S}$ , where FDiv values approached 1 (Figure 7b). The RaoQ index, measuring functional dissimilarity, increased with depth ( $>500$  m) across most samples (Figure 7c). Conversely, functional dispersion (FDis) revealed low values ( $<3$ ) near  $11^{\circ}\text{S}$  (Figure 7d). Functional evenness (FEve) was relatively low to intermediate (0.4 – 0.65) across the study area (Figure 7e). The uniform distribution of traits (FEve  $> 0.7$ ) was observed in a mere 14% of the sites, predominantly situated south of  $26^{\circ}\text{S}$ .



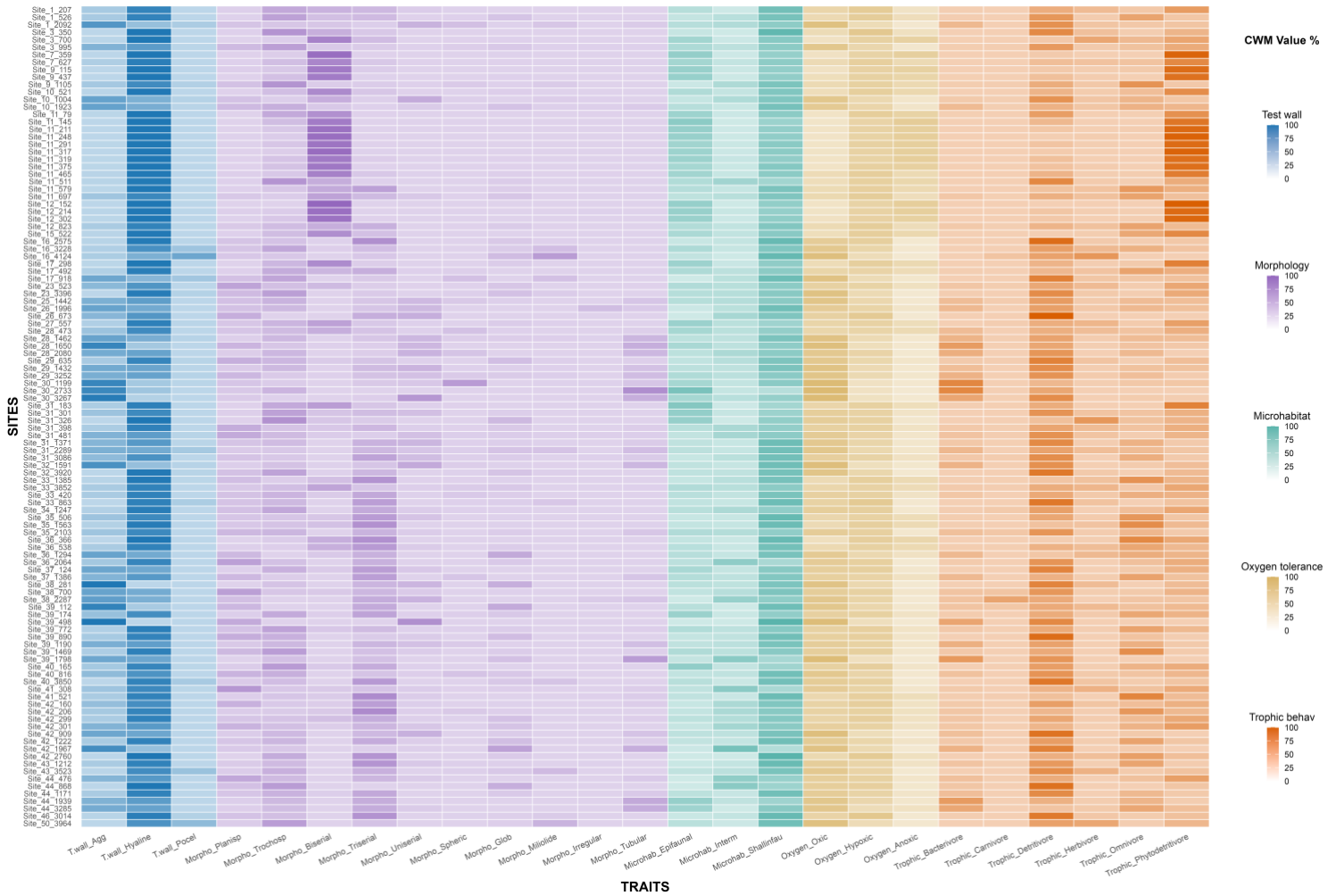
**Figure 8.** Functional diversity index of benthic foraminifera along the Southeastern Pacific. (a) Rao Quadratic Functional Diversity Index. (b) Functional Richness. (c) Functional Divergence. (d) Functional Dispersion. (e) Functional Evenness

### 7.3. Characterization of Functional Traits

Community-weighted means (CWM) of traits revealed clear spatial patterns along the Southeastern Pacific, particularly in morphology, oxygen preference, and trophic behavior. Morphological traits (Figure 9) exhibited more distinct spatial variation. Biserial forms, mainly composed by *Bolivina* (15 sp), were found to predominate (>80%) between 9° and 11°S, but in less percentage south of 17° (Figure 8), where the assemblages exhibited a transition towards a predominance of trochospiral and triserial morphologies (collectively 30% – 80%). This change is given by *Trifarina angulosa*, *Hoeglundina elegans*, *Cibicidoides* and *Uvigerina* (Table S6). Oxygen and trophic preferences also exhibited clear latitudinal shifts. Hypoxia-tolerant species consistently accounted for 40–60% of assemblages, with peaks (>60%) at bathyal depths south of 17°S. Detritivores were the dominant feeding strategy throughout, while phytodetritivores were particularly abundant between 1°–15°S and 23°–36°S. Conversely, omnivores gained prevalence south of 33°S, and bacterivores contributed substantially (35–60%) at specific bathyal sites (~1500 m, near 30°S). Finally, test wall type and microhabitat preference showed less pronounced spatial variation. Hyaline tests occur across the study area contributing mostly with >50% in the assemblages, as well, the less contribution (<25%) was at 1000–3000 m off 28°–32°S and coincide with the highest values of agglutinated (>70%). Agglutinated test presented values <25% in most of the sites (73) exhibiting a minimum or zero contribution in neritic to upper-bathyal zones (79–700 m) between 3°–17°S and some deepest sites off 42°, 43°, 46° and 50°S, showing no strong latitudinal gradient. Despite the

contribution of porcelaneous tests being small across the stations, its remarkable contribution is notable (8–28%) at some sites >900 m (Figure 9). Shallow infaunal taxa were ubiquitously dominant (50–100%) in all the sites, while epifaunal taxa showed highest contribution (>70%) between 3°–12°S, and intermediate infaunal taxa showed dominance <30% at most of the sites.

Spearman's correlation analysis revealed that species richness (taxa) is the main driver of functional richness in the benthic foraminiferal community, given the positive and significant correlation ( $\rho = 0.841$ ,  $p < 0.01$ ; Appendix 6). Similarly, species richness exhibited a positive correlation with various other taxonomic diversity indices, except for Pielou's evenness index ( $-0.04$ ,  $p \geq 0.05$ ), indicating that species richness is not dependent on abundance uniformity (Appendix 6). FEve and abundance ( $\rho = -0.667$ ,  $p < 0.001$ ), indicating that sites with high individual's densities, are dominated by species with generalist traits. Unexpectedly, a negative correlation between the abundance and RaoQ ( $\rho = -0.367$ ,  $p < 0.001$ ) was observed, reflecting that sites with many individuals are often dominated by a few species. Finally, FDiv showed no correlation with species richness ( $\rho = -0.057$ ,  $p > 0.05$ ), therefore a greater number of species does not necessarily correspond to dominance of species with specialized traits.



**Figure 9.** Community Weighted Means (CWM) of functional traits of living benthic foraminifera along the southeastern Pacific. Latitudinal order is shown from bottom to top. Colors indicate trait categories: blue = test wall type; purple = morphology; green = microhabitat preference; yellow = oxygen tolerance; orange = trophic behavior. T.wall = test wall type, Agg = agglutinated, Porcel = Porcelanaceous, Planisp = planispiral, Trochosp = trochospiral, Glob = Globose/subglobose/ovoid or streptospiral.

#### 7.4. Environmental variables which modulate the taxonomic and functional indices

The GAM analyses identified temperature, salinity, oxygen, phosphate, and silicate as the primary environmental drivers affecting both the functional and taxonomic diversity indexes (Table 4). Abundance, species richness (taxa), and functional richness (FRic) of benthic foraminifera were also influenced by sediment properties, specifically the C/N ratio, opal, and CaCO<sub>3</sub>, respectively (Table 5). Latitude, longitude, and d15N (δ15N) were significant predictors only for species richness and FRic. Oxygen was the sole variable affecting functional dispersion (FDis) and functional evenness (FEve). No significant relationships were detected between environmental variables and functional divergence (FDiv) or Rao's Q.

**Table 4.** GAM models for the taxonomic and functional indexes, including response variables, predictor variables, AIC, R<sup>2</sup>, and deviance explained.

Response Variable	Predictor Variables	AIC	R-sq	Deviance explained
Shannon	Temperature, Salinity, Depth, Oxygen, Phosphate, Nitrite, Silicate, TOC, TN, Opal, CaCO <sub>3</sub> , C/N, d15N, Longitude, Latitude, bs = "gp"	42.195	0.787	89.3%
Taxa		337.142	0.828	97.3%
Simpson		NA	0.771	84.8%
Pielou		-90	0.648	81.3%
Abundance		164.231	0.806	90%
FRic		417.919	0.922	91.9%
FDis		-55.425	0.622	74.4%
FDiv		-55.983	0.227	64.3%
FEve		-26.442	0.091	55.7%
RaoQ		85.299	0.47	63.8%

**Table 5.** Summary of GAM results showing the effects of environmental variables on taxonomic and functional diversity indices ( $p < 0.05$ ).

Index	Significant variables	Degrees of Freedom	p-value
Shannon	Temp	3.75	p<0.01
	Salinity	1.00	p<0.05

	Oxygen	3.08	p<0.01
	Phosphate	1.00	p<0.01
	Silicate	1.00	p<0.01
<b>Taxa</b>	Temp	1.00	p<0.01
	Salinity	1.00	p<0.01
	Oxygen	1.00	p<0.01
	Phosphate	1.00	p<0.01
	Silicate	1.00	p<0.01
	Opal	7.19	p<0.01
	CaCO <sub>3</sub>	3.36	p<0.05
	Long, Lat	12.64	p<0.01
<b>Simpson</b>	Temp	1.00	p<0.01
	Salinity	1.00	p<0.05
	Oxygen	2.83	p<0.01
	Phosphate	3.97	p<0.01
	Silicate	1.00	p<0.01
<b>Pielou</b>	Oxygen	1.00	p<0.05
	Phosphate	3.82	p<0.05
	Silicate	1.00	p<0.05
<b>Abundance</b>	Salinity	3.70	p<0.05
	C/N	1.00	p<0.05
<b>FRic</b>	Temp	1.00	p<0.01
	Salinity	1.00	p<0.05
	Oxygen	2.62	p<0.01
	Phosphate	1.00	p<0.01
	Silicate	1.00	p<0.01
	Opal	6.72	p<0.01
	CaCO <sub>3</sub>	1.69	p<0.05
	δ <sup>15</sup> N	1.00	p<0.01
<b>FDis</b>	Oxygen	1.00	p<0.05
<b>FDiv</b>	N.A	-	-
<b>FEve</b>	Oxygen	3.13	p<0.05
<b>RaoQ</b>	N.A	-	-

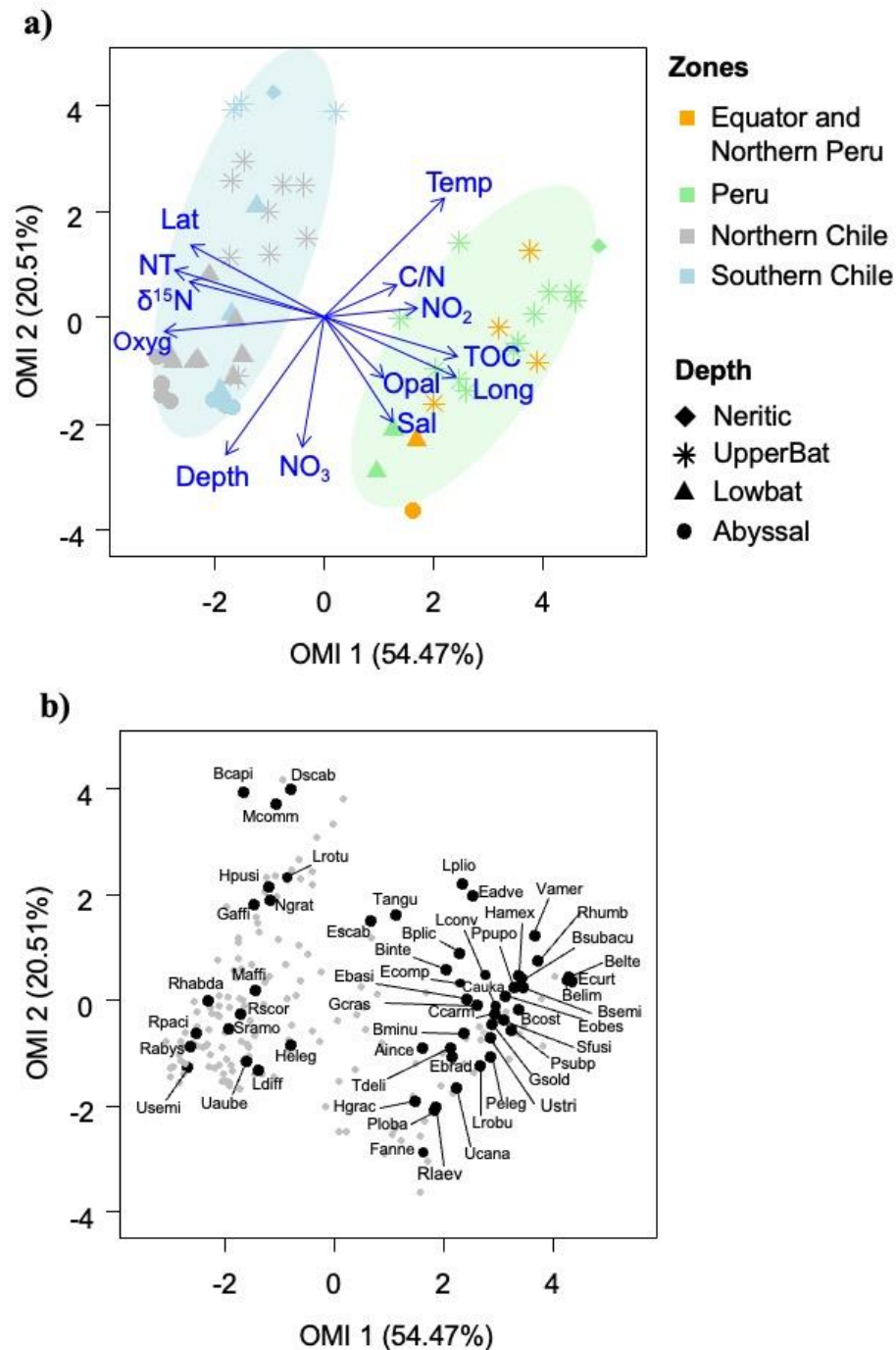
## 7.5. Niche analysis

The realized niche analysis (OMI) revealed a strong structure of the benthic foraminiferal community along the environmental conditions (Figure 10a). The first and second axis of the OMI analysis encompassed 74.98% (OMI 1, 54.47% variance; OMI 2, 20.51% variance) characterizing two groups of stations. One group is formed by the Ecuador and Peru zones which are associated with low dissolved oxygen, high TOC, nitrite, opal and ratio C/N. While the second group, formed by Northern and Southern Chile, is characterized by greatest oxygen values, total nitrogen,  $\delta^{15}\text{N}$  and depth. Additionally, OMI 2 clearly shows a bathymetric gradient from neritic to abyssal stations, with high to low temperature values, respectively. Similarly, Figure 10b exhibited a pattern of species grouping, which tends to organize themselves in relation to the same environmental variables as shown in Figure 10a.

Regarding to the distribution of the functional traits across the two OMI axes (Figure S8), we observed that species with hyaline test were strongly associated with Peru and Ecuador zone characterized by high TOC, and nitrite, at neritic and upper bathyal sites. Species with agglutinated test walls are related to sites with more oxygen and greater depth, mainly in Chile zone (Figure S8a). In this region, morphologies such planispiral, irregular, uniserial, and triserial shells dominate mainly at upper and lower bathyal zones (Figure S8b). On the other hand, the biserial morphology was more common at sites off Peru at neritic and upper bathyal depths, influenced mainly by the high temperature, C/N and  $\text{NO}_2^-$ . Finally, the microhabitat and trophic behavior traits (Figure S8c, S8d) did not show a grouping pattern.

In addition, marginality was assessed for 281 species (Table S7). The classification of niche strategies for species was based on the absolute values of OMI and TOL (Table 6). Both metrics were conducted using quartiles, which were calculated from the distribution of values for all species analyzed. *Vaginulina americana* (Vamer - abbreviations used in Table 6 and Fig. 9b), *Ebuliminella curta* (Ecurt), *Buliminella elegantissima limbosa* (Belim), and *Buliminella elegantissima tenuis* (Belte) had the highest Marginality values of the group (OMI

= 35.27, 31.67, 31.02, and 30.50, respectively), indicating marked specialization (Low Tolerance, Tol = 0.00, 0.03, 0.03, and 0.02, respectively). Agglutinated species *Bathysiphon capillare* (Bcapi), which is present in southern Chile at upper bathyal depth, also exhibited analogous values (OMI = 24.45, Tol = 0.00; Figure 9 a, b). Species as *Eggerelloides advena* (Eadve), *Hanzawaia mexicana* (Hamex), *Bolivina subadvena acuninata* (Bsubacu), *Bulimina pagoda* (Bpago), *Bolivina seminuda* (Bsemi) and *Protoglobobulimina pupoides* (Ppupo) exhibited moderate to high marginality (OMI = 23.27, 21.2, 19.77, 17.99, 15.28 and 13.40, respectively) and higher tolerance values than strict specialists (Tol = 2.7, 3.09, 1.01, 0.49, 5.22 and 4.35, respectively). Therefore, these species are interpreted as flexible specialists. Most of the species showed moderate marginality (OMI = <13.38–6.72) and variable tolerance values (Tol = 0.06–2.83), representing intermediate strategies. Finally, the most generalist species are *Hoeglundina elegans* (Heleg) and *Melonis affinis* (Maffi), which present the lowest marginality value (OMI = 1.65 and 2.25, respectively) with a high tolerance (Tol = 1.42 and 1.17, respectively).



**Figure 10.** Outlying mean index (OMI) of the influence of the environmental variables on the functional traits of the benthic foraminifera species. a) Plot representing the environmental variables tested. The length of the blue vectors represents the importance of each variable. The zone refers to the area to which the sample sites belong. Ecuador and Northern Peru are the sites off 1°S–3°S, Peru: 7°S–17°S, North-Chile: 23°S–39°S, South-Chile: 39°S–50°S. Depth refers to the water depth the sample has been taken. Neritic: 0–250 m, UpperBat: >250–850 m, Lowbat: >850–2000 m, Abyssal: >2000 m. b) Realized

niches of benthic foraminifera species in the Southeastern Pacific. •□ Species that showed significant marginality ( $p < 0.05$ )

**Table 6.** Niche parameters of the species of benthic foraminifera and their traits are estimated by the Outlying Mean Index (OMI). Niche parameters are given for each species as both absolute values and percentages of variability. Inertia = total variability, OMI = Outlying Mean Index (i.e., marginality), Tol = Tolerance, RTol = Residual Tolerance. Only species with a significant OMI index ( $p < 0.05$ ) are shown.

Code	Specie	OMI	Tol	Rtol	Classification
Vamer	<i>Vaginulina americana</i>	35.27	0	0	Strict Specialist
Ecurt	<i>Eubuliminella curta</i>	31.67	0.03	0.35	Strict Specialist
Belim	<i>B. elegantissima limbosa</i>	31.02	0.03	0.40	Strict Specialist
Belte	<i>B. elegantissima tenuis</i>	30.5	0.02	0.45	Strict Specialist
Rhumb	<i>Rhumblerella humboldti</i>	29.35	0.68	2.45	Strict Specialist
Bcapi	<i>Bathysiphon capillare</i>	24.45	0	0.16	Strict Specialist
Mcomm	<i>Martinottiella communis</i>	22.03	0.03	2.40	Strict Specialist
Dscab	<i>Dorothia scabra</i>	20.27	0.01	1.98	Strict Specialist
Eadve	<i>Eggerelloides advena</i>	23.27	2.7	5.93	Flexible Specialist
Lplio	<i>Lenticulina pliocaena</i>	21.77	1.25	7.76	Flexible Specialist
Hamex	<i>Hanzawaia mexicana</i>	21.2	3.09	5.56	Flexible Specialist
Bsubacu	<i>Bolivina subadvena acuminata</i>	19.77	1.01	5.79	Strict Specialist
Bpago	<i>Bulimina pagoda</i>	17.99	0.49	1.95	Strict Specialist
Bsemi	<i>Bolivina seminuda</i>	15.28	5.22	7.65	Flexible Specialist
Lconv	<i>Lenticulina convergens</i>	14.5	2.22	5.05	Flexible Specialist
Psubp	<i>Pseudoparrella subperuviana</i>	14.16	1.56	3.43	Flexible Specialist
Ppupo	<i>Protoglobobulimina pupoides</i>	13.4	4.35	5.48	Flexible Specialist
Ucana	<i>Uvigerina canariensis</i>	13.38	0.12	2.66	Intermediate Low Tol.
Ploba	<i>Pseudobrizalina lobata</i>	13.24	0.33	8.45	Intermediate
Bcost	<i>Bolivina costata</i>	12.67	2.5	9.24	Intermediate
Fanne	<i>Fissurina annectens</i>	12.05	0.34	3.58	Intermediate
Cauka	<i>Cassidulina auka</i>	11.19	1.51	6.96	Intermediate
Eobes	<i>Epistominella obesa</i>	11.02	0.54	7.97	Intermediate
Ustri	<i>Uvigerina striata</i>	10.96	0.55	4.84	Intermediate
Sfusi	<i>Stainforthia fusiformis</i>	10.91	0.84	7.94	Intermediate

Peleg	<i>Pullenia elegans</i>	10.67	0.31	7.54	Intermediate
Lrobu	<i>Labrospira robusta</i>	10.56	0.06	6.52	Intermediate Low Tol.
Rlaev	<i>Reussoolina laevis</i>	10.48	1.42	5.94	Intermediate
Ccarm	<i>Cancris carmenensis</i>	10.12	2.83	6.09	Intermediate High Tol.
Gsold	<i>G. soldanii multilocula</i>	10.11	0.54	6.93	Intermediate
Usemi	<i>Uvigerina semiornata</i>	9.84	0.12	0.89	Intermediate Low Tol.
Bminu	<i>Bolivinita minuta</i>	9	0.33	5.88	Intermediate
Rabys	<i>Rhabdammina abyssorum</i>	8.81	0.15	0.83	Intermediate Low Tol.
Ecom	<i>Ehrenbergina compressa</i>	8.68	0.17	4.34	Intermediate Low Tol.
Gcras	<i>Globocassidulina crassa</i>	8.65	1.7	7.04	Generalist
Hgrac	<i>Hyalinonettrion gracillimum</i>	8.28	3.87	5.08	Generalist
Rpaci	<i>Rhabdammina pacifica</i>	7.94	0.70	2.58	Generalist
Bplic	<i>Bolivina plicata</i>	7.7	4.94	8.16	Generalist
Lrotu	<i>Lenticulina rotulata</i>	7.23	2.58	4.67	Generalist
Tdeli	<i>Takayanagia delicata</i>	6.82	2.9	6.57	Generalist
Ebrad	<i>Eggerella bradyi</i>	6.72	0.31	9.28	Generalist
Hpusi	<i>Haplophragmoides pusillus</i>	6.44	3.16	5.60	Generalist
Gaffi	<i>Globobulimina affinis</i>	5.96	2.9	5.46	Generalist
Tangu	<i>Trifarina angulosa</i>	5.75	1.07	12.36	Generalist
Binte	<i>Bolivina interjuncta</i>	5.64	1.92	11.18	Generalist
Ngrat	<i>Nonionoides grateloupii</i>	5.63	4.26	6.13	Generalist
Escab	<i>Eggerelloides scabrum</i>	5.56	0.92	11.14	Generalist
Rhabda	<i>Rhabdammina</i> sp.	5.29	0.48	4.85	Generalist
Aince	<i>Ammodiscus incertus</i>	4.77	3.79	11.53	Generalist
Sramo	<i>Saccorhiza ramosa</i>	4.28	1.51	7.12	Generalist
Ldiff	<i>Lagenammina diffflugiformis</i>	4.20	1.26	8.82	Generalist
Uaube	<i>Uvigerina auberiana</i>	3.9	0.96	4.56	Generalist
Rscor	<i>Reophax scorpiurus</i>	3.27	2.89	7.64	Generalist
Maffi	<i>Melonis affinis</i>	2.25	1.17	8.40	Generalist
Heleg	<i>Hoeglundina elegans</i>	1.65	1.42	8.76	Generalist

## 8. DISCUSSION

### 8.1. Taxonomic diversity patterns

This study recorded a total living benthic foraminifera richness of 365 species along the investigated latitudinal gradient (1°–50°S) in the SEP. This number is greater than previous studies (Bandy & Rodolfo, 1964; Mallon, 2011; Romero *et al.*, 2023; Tapia *et al.*, 2008). Furthermore, we recorded *Astrorhizoides cornutus*, *Crithionina pisum*, *Galwayella trigonoornata*, *Liebusella goesi* in the SEP. These species have been found mainly in the Northwest Atlantic, near fjords or in the Arctic Ocean (Gooday *et al.*, 2008; Hesemann, 2023; Jima *et al.*, 2022; Schönfeld, 2002). However, the genus *Galwayella* has been established a few decades ago (Patterson & Pettis, 1986) and has only been reported in a few instances (Hesemann, 2023; Patterson & Burbidge, 1995). *G. trigonoornata* is described here for the first time in the region. Interestingly, several species found during this study were previously denominated as being extinct (*Fissurina (Fissurina) ovata*, *Haplophragmoides bulloides*, *Lagena parvulipora*, *Reophax duplex*, *R. globosus*). These were reported from the Upper Cretaceous/Lower Tertiary (*H. bulloides*) in Europe (Bugge *et al.*, 1975; Neagu, 1968), Upper Cretaceous-Eocene (*R. duplex*, *R. globosus*) (Silva, 2011; Waśkowska, 2021), or rather vague in description (Seguenza, 1868). *L. parvulipora* was reported wrongly as *Pygmaeoseistrum parvuliporum* for the Miocene in Chile (Finger, 2013). The author cites the original description of Bandy (1949a), which was corrected later (Bandy, 1949b).

The diversity indexes (Shannon, Fisher- $\alpha$ , Pielou's evenness) exhibited a strong latitudinal and bathymetric variation (Figure 5a). The highest values of diversity

and species richness were observed at sites influenced by well-oxygenated water masses as Subantarctic (SAAW) and Antarctic intermediate Water (AAIW), where other authors have identified a similar pattern (Bandy & Rodolfo, 1964; Ingle *et al.*, 1980; Patarroyo & Martinez, 2021). In contrast, areas bathed by the ESSW, characterized by hypoxia and high nutrient concentrations (Figure 2c), showed a low overall diversity, in particular, between 7° and 12°S and at 100 – 50 m (Figure 5b), where severe hypoxia prevailed (1-2  $\mu\text{mol/kg}$ ). A Shannon index  $<1$ , Pielou evenness  $<0.05$ , and the Fisher- $\alpha <3$  reflects an ecosystem with high abundances of a few species only (notably *Bolivina* spp.) as reported in previous studies (Cardich *et al.*, 2012; Glock *et al.*, 2013; Mallon *et al.*, 2012), demonstrating that OMZs are strong environmental filters, favoring the adaptation and development of opportunistic or hypoxia-tolerant species.

The abundance of benthic foraminifera exhibited a remarkable division of high abundances between 1°–17° S, and consistently lower abundances between 22°–50° S (Figure 4). This pattern overlaps with the year and seasonal round upwelling (~4°–35°S), as well as with high biogenic opal content in surface sediments (Figure S1), which reflects primary productivity and export fluxes to the seafloor (Tarazona & Arntz, 2001; Zuta & Guillen, 1970). Within the northern productive sector, the highest abundances ( $>10,000$  ind/50  $\text{cm}^3$ ) were recorded between 9°–12°S, mainly dominated by *Bolivina* species (Figure S3). Similar oxygen deficient environments, such as the Arabian Sea, have documented as dominant taxa to *Bolivina* and *Uvigerina* species (Kaithwar *et al.*, 2020; Schumacher *et al.*, 2007). The dominance of bolivinidae across the equatorial to central Peruvian margin

suggests that opportunistic species that thrives well under low-oxygen conditions in bottom/pore waters associated with high organic matter preservation at shallow infaunal sediment depths. Conversely, more heterogeneous assemblages at southern latitudes indicated that moderate to high oxygen levels and variable sediment composition supported a higher taxonomic evenness and diversity (Castillo *et al.*, 2021; Tapia *et al.*, 2008).

## **8.2. Functional diversity, linking with taxonomic diversity**

Functional richness (FRic) was low (<35). Only 14 sites showed high values (>70). In several of these sites, high FRic coincided with elevated species richness (~40) and low abundances (Figure 7a, 3a; Table S5). The overall positive correlation between species richness and FRic ( $\rho = 0.841$ ,  $p < 0.01$ ) supports the expectation that the functional space increased with the number of species (Petchey & Gaston, 2002). Functional space represents the multidimensional space occupied by species' traits, and the volume or dispersion of this space reflects the functional strategies present in the community (Mason *et al.*, 2003). However, most stations displayed a low FRic regardless of species count, indicating that relationship was not consistent across the region. A decoupling can be explained by “functional redundancy” or “functional similarity”, where multiple species share similar trait combinations and therefore contribute similarly to ecosystem functioning (Eisenhauer *et al.*, 2023; Fischer & de Bello, 2023; Petchey & Gaston, 2006). Consequently, species-rich communities may occupy a relatively narrow functional space. This pattern is complemented by functional

evenness (FEve), especially at sites where its negative correlation with total abundance ( $\rho = -0.667$ ,  $p < 0.01$ ) reveals that these communities reflect a small set of strategies, mainly stress-tolerance (OMZ sites), resulting in an uneven distribution of functions within the functional space (Mason *et al.*, 2005; Vileger *et al.*, 2008). Functional divergence (FDiv) reinforced the interpretation, remaining high at most stations ( $>0.66$  at 79% of sites). This pattern implies that the dominant species occupy the extremes of functional space in these communities and are well adapted to environmental stress (Vileger *et al.*, 2008).

The integration of these indices indicates that the benthic ecosystem does not represent a continuous gradient across the SEP. Only few sites exhibit moderate to high FEve ( $\geq 0.6$ ) and RaoQ ( $\geq 17$ ) high FRic ( $>66$ ), and high FDiv (Figure 7b). This specific combination represents communities where numerous strategies coexist in equilibrium. The traits are supported by balanced abundances of high functional complementarity, species differ markedly in their traits, and by a low probability of functional similarity, which indicated an efficient resource utilization (Mason *et al.*, 2013; Xie *et al.*, 2025). These sites occur mainly in the deepest areas ( $>1000\text{m}$ ) of Ecuador and northern Peru ( $1^{\circ}$ – $3^{\circ}\text{S}$ ), outside the OMZ core but also between  $27^{\circ}$ – $32^{\circ}\text{S}$ . This type of correlation (high species richness, high functional diversity) has been recorded in deep sea ecosystems where the functional similarity is low, and a high number of species has a direct effect on functional diversity (Danovaro *et al.*, 2008). These specific zones could also be influenced by local characteristics, for instance the change from permanent to

seasonal upwelling and the topography (Figueroa & Moffat, 2000; Montecino & Lange, 2009).

Assemblages that present high FEve and RaoQ, moderate to high FDiv (~0.5–0.9) and low FRic (<23), are randomly distributed between 300 – 4000 m. This indicates a community where the small functional space maintains distinct roles with balanced coexistence (Mason *et al.*, 2013). Most sites were south of 16°S and >900 m. Other taxonomic groups revealed that changes in functional diversity components with taxonomic diversity are due to the environmental characteristics of the zone or habitat heterogeneity (Buhl-Mortensen *et al.*, 2012; Carvalho & Barros, 2017; Gomes *et al.*, 2024). In this case of the SEP, the respective sites were exposed to the mixing zone of AAIW, ESSW and PDW, characterized by low temperatures (~3°C), dissolved oxygen (~100–218  $\mu\text{mol kg}^{-1}$ ) and phosphate concentrations increasing from ~1.9 to > 2.4  $\mu\text{mol kg}^{-1}$  (Reyes-Macaya *et al.*, 2022; Silva *et al.*, 2009). Additionally, being outside the OMZ, these characteristics were also mirrored by sediment heterogeneity and thereby constitute a heterogeneous zone with greater species diversity for meio- and macrofauna (Neira *et al.*, 2001; Sellanes *et al.*, 2010), hence influencing the functional structure. Environmental heterogeneity generates an assemblage with low redundancy but high niche complementarity.

On the other hand, sites with lowest FRic (<10), low to intermediate FEve (0.2–0.5) and RaoQ (1–18), high FDiv and high abundances were mainly found in the shallow OMZ (9°, 10°, 11°, 12°, 31°S). The low FRic suggests an environmental filter linked to severe hypoxia in the area, exhibiting a high functional similarity in

the assemblage. It was dominated by a few species (low FEve), which possess similar tolerant traits to anoxia conditions such as elongated morphologies and physiological strategies to overcome the dependence on oxygen (Glock, 2023; Koho & Piña-Ochoa, 2012; Mason *et al.*, 2013). This pattern matches with other macrofaunal and meiofaunal assemblages along Peruvian and Chilean OMZ, where low diversities but high densities of hypoxia-tolerant groups dominate (Gómez *et al.*, 2024; Neira *et al.*, 2001; Palma *et al.*, 2005). Despite the restricted functional space, the high FDiv indicates that few surviving taxa occupy distinct positions within the constrained functional space, representing highly specialized and functionally distinct strategies adapted to the OMZ (Mouillot, Bellwood, *et al.*, 2013). Slight variations of this functional combination extend to sites surrounding the OMZ (1°–15°S and 31°–39°S) and at upper bathyal depths (300–800 m). This suggested that although the functional space increases slightly, the dominance of a small set of tolerant traits persists under the influence of low oxygen levels.

### **8.3. Functional traits distribution**

The community-weighted mean (CWM) trait analyses indicated strong latitudinal shifts in morphology, oxygen tolerance, and trophic strategies (Figure 8). The hyaline test type was dominant across the SEP, as this type is considered the most diverse (Murray, 2007). However, agglutinated species predominated (30%) at deeper sites (>990 m). This pattern coincides with the sites showing elevated FRic, which include both test type wall type and multiple morphologies (Bandy & Rodolfo, 1964; Patarroyo & Martinez, 2021) suggesting that sites at intermediate

and deeper sites harbor more morphological traits. The lower contribution of porcelanaceous or miliolide species could be related to oxygen conditions of the study area, since *Quinqueloculina*, *Pyrgo* and *Triloculina* species prefer cold environments with high oxygenation, low organic carbon content, and sandy to slightly muddy sediments (Anbuselvan & Senthil Nathan, 2019; Narciso-Mezones *et al.*, 2025; Noucoucouk *et al.*, 2023). On the other hand, a distinct pattern was observed in the distribution of agglutinated species (Figure 8), even though they are not well adapted to these conditions (Mallon *et al.*, 2012). None the less, agglutinants such as *Ammoglobigerina globigeriformis*, *Cyclamina cancellata*, *Ammodiscus incertus*, *Eggerella bradyi* was lower (<2%) at certain stations (~3°-12°S). The proportions of arenaceous species slightly increased (~5–17%) south of the OMZ at 15°S, which is displayed by *Haplophragmoides*, *Reophax*, *Portatrochmmina*, *Trochammina*, *Dorothia* and *Bathysiphon*. These taxa have been associated with low oxic conditions (Gooday *et al.*, 2000; Koho & Piña-Ochoa, 2012; Tetard *et al.*, 2024), suggesting that the environmental conditions along the OMZ are not homogeneous. According to Gooday *et al.* (2009), the abundance of agglutinated species starts to increase beyond 500 m, and beneath the OMZ (~1300 m), from where on the dominates the community.

Morphologies revealed a clear distinction between assemblages found within and outside the OMZ core. Inside and surrounding the OMZ there was a predominance of biserial, triserial and trochospiral species, e.g. *Bolivina*, *Uvigerina*, *Bulimina* and *Fursenkoina* (Table S7). These taxa, also designated as rectilinear or tapered/cylindrical morphogroup, have been reported in similar OMZ

ecosystems as a good indicator of these conditions (Kaithwar *et al.*, 2020; Mazumder & Nigam, 2014). Although the prevalence of these morphologies under low oxygen conditions is not fully resolved, it may be related to the elongated and compressed arrangement of the chambers, which provide functional advantages by enhancing gas-exchange efficiency. For instance, species such as *Uvigerina peregrina* increase their chamber size during growth, while *Bolivina* or *Loxostomum* modulate their chamber number (Keating-Bitonti & Payne, 2017). These morphologies may also compensate for the reduced surface area to volume ratios of larger tests. Through increased porosity, they improve gas exchange (Belanger, 2022). This is of particular importance in OMZ ecosystems *Bolivina*, *Buliminella*, *Fursenkoina*, *Nonionella* and *Uvigerina*, need to acquire and store nitrate to perform denitrification, which allows them to respire in the absence of oxygen (Bernhard *et al.*, 2012; Glock *et al.*, 2013; Høgslund *et al.*, 2008). Additionally, those species may rely on kleptoplastidy to supplement their metabolism under oxygen-depleted conditions (Glock, 2023; Meilijson *et al.*, 2015). In the same way, traits such as pore size or density, although not quantified in this study, are recognized as morphological adaptations in the efficiency of gas exchange under low-oxygen conditions (Glock *et al.*, 2019; Glock *et al.*, 2012; Kuhnt *et al.*, 2013). The dominance of species with these adaptations in our sites is consistent with hypoxia and anoxia tolerant traits observed in the CWM analysis, suggesting those traits would be contributing most strongly within the core of the OMZ. Additionally, body size is an important morphological trait because low-oxygen conditions often lead to a higher proportion of small-sized

individuals, thereby altering the diversity and abundance of the assemblage (Palmer *et al.*, 2020).

The availability of organic matter is another factor that can modulate foraminiferal test shape (Caulle *et al.*, 2015; Keating-Bitonti & Payne, 2016). These above-mentioned strategies coincide with ecological traits linked to organic-matter utilization. Many of the species dominating oxygen depleted sites exhibited high CWM values for the phytodetritivore trait. Although detritivore trophic behavior was widespread across the study area, phytodetritivore species were particularly abundant in the shallow OMZ core sites and between 33°–42°S. Those sites are subjected to a high export flux of organic matter to the sediments (Gutiérrez *et al.*, 2011; Loginova *et al.*, 2020). However, C/N ratios in these sites ranged from moderate to high values (~6.5–9.5), indicating that organic matter reaching the sediment is not exclusively fresh in phytodetritus but includes partially degraded material (Figure S1). Consequently, the dominance of phytodetritivore traits does not necessarily reflect strict dependence on labile organic matter, but rather opportunistic strategies of episodic pulses of fresh material. Such feeding strategy has been reported for some species inhabiting OMZ (Cardich *et al.*, 2015; Nomaki *et al.*, 2006), where ecological success relies more on physiological tolerances. Although most foraminiferal species are catalogued at a lower trophic level (Nomaki *et al.*, 2008), yet their feeding strategies are often flexible. At some sites south of 33°S and below the OMZ core (>600m), where C/N ratios are lower (<7.4) but total abundance decreases, omnivore behavior was more common (>20%) (Figure 8). These settings indicate that the organic matter is fresher, but

the flux is lower and competition for resources increases (Neira *et al.*, 2001). Hence, omnivores likely reflect feeding flexibility, enabling them to exploit a wider range of food resources (Gooday *et al.*, 1992; Gooday *et al.*, 2008; Murray, 2006). At the deepest sites, bacterivore species became more abundant (>25%), particularly agglutinated species. However, foraminifera have been considered to not consume bacteria selectively but to ingest them incidentally (Goldstein & Corliss, 1994; Gooday *et al.*, 2008). The apparent bacterivory probably reflects the general low food availability and degradation of sinking organic matter in bathyal to abyssal environments, where the sediments are dominated by microbial biomass and lipids (Dale *et al.*, 2015; Flores *et al.*, 2022). Consequently, the trophic diversity observed here likely reflects opportunistic, deposit or suspension-feeding strategies, where species classified as omnivores or bacterivores may switch between feeding modes depending on food availability and degradation state (Gooday *et al.*, 1992).

The microhabitat preferences trait did not exhibit a strong spatial pattern. It is controlled by oxygen penetration in the sediment, quality and quantity of organic matter, and bioturbation rates (Corliss, 1985; Geslin *et al.*, 2004). Epifaunal and shallow infaunal (SI) species were present in most shallow and hypoxic sites, in particular calcareous rotaliid taxa with tolerance to low oxygen conditions (Figure 8). The microhabitat selection reflects an adaptation to anoxia and preference for labile organic matter (Cardich *et al.*, 2015; Fontanier *et al.*, 2002). These assemblages, as well as the species that occupy deep or intermediate sediments

depths in higher oxygenated settings within the SEP OMZ follow the TROX model (Jorissen *et al.*, 1995). In this framework, taxa adjust their microhabitat position based on food availability (Cardich *et al.*, 2015), thereby contributing to sediment mixing as bioturbators (Deldicq *et al.*, 2020; Diz *et al.*, 2023). Additionally, it has been evaluated that some epifaunal species could rework particles at the sediment surface, whereas infaunal species, that live within sediments, contribute to particle mixing from the surface down to the deepest sediment layer they can occupy (Deldicq *et al.*, 2020).

Nevertheless, the SI outside the OMZ, particularly in the southernmost part of the SEP (~31°–50°S), while intermediate infaunal (II) exhibits more contribution. These sites demonstrate elevated oxygen levels, intermediate values of organic carbon accumulation, as well as less organic matter degradation (Figure S1). Such conditions are likely to facilitate more flexible vertical distribution, therefore benthic foraminifera can occupy the microhabitat that best suits their trophic preferences, bottom water conditions, or sediment grain type (Anbuselvan & Senthil Nathan, 2019; Enge *et al.*, 2016).

#### **8.4. Environmental drivers**

GAM analyses highlighted that temperature, salinity, oxygen, phosphate, and silicate concentrations were the main environmental drivers shaping both taxonomic and functional diversity (Table 4). These predictors are linked to the water mass structure and act as integrative descriptors of them rather than

independent drivers (Reyes-Macaya *et al.*, 2022; Silva *et al.*, 2009). Temperature and salinity are defined as conservative tracers while the oxygen and nutrients (e.g. nitrate, phosphate, silicate) are non-conservative properties that can change by chemical or biological process in the column water (Talley *et al.*, 2011), while nutrients also represent signals of organic matter remineralization. These bottom water variables have been shown to drive the diversity of other macrobenthic groups (Neira *et al.*, 2001; Ysebaert & Herman, 2002) indicating that the foraminiferal communities respond similarly to environmental gradients. In this context, the effect of this set of variables on taxonomic metrics is captured by the shifts in species dominance and presence/absence patterns among sites, whereas functional richness reflects changes in the extent of functional space, allowing a broader range of ecological strategies to be expanded under specific water mass characteristics (Bandy & Rodolfo, 1964; Ingle *et al.*, 1980). These interactions have been reported in previous studies where some foraminiferal communities adapt to those properties of bottom-water masses (Eichler *et al.*, 2016; Eichler *et al.*, 2008).

Sedimentary and geochemical variables such as opal, CaCO<sub>3</sub> content,  $\delta^{15}\text{N}$  and C/N ratio values contributed to explaining variation in taxonomic indices and functional richness, indicating a relevant role of benthic habitat characteristics. Biogenic opal in surface sediments is widely used as a proxy for past and present primary productivity, reflecting the export of siliceous phytoplankton to the seafloor during upwelling conditions (Mohtadi & Hebbeln, 2004; Romero & Hebbeln, 2003). Therefore, higher opal contents in the study area indicate organic matter fluxes,

which can promote species richness and influence dominance patterns through increased food availability. Similarly,  $\text{CaCO}_3$  content has been shown to contribute to sediment composition and spatial heterogeneity, which has implications for variability in grain-size (Nait-Hammou *et al.*, 2025). In the present study,  $\text{CaCO}_3$  in surface sediments reflects the biogenic production given by the deposition of calcifying organisms (Mohtadi *et al.*, 2005; Saavedra-Pellitero *et al.*, 2013). Hence, the benthic foraminifera community structure could be influenced by the composition or the sediment grain size (Hromic *et al.*, 2006; Milker *et al.*, 2009). From a functional perspective, biogenic opal likely increases the FRic, due to its association with surface productivity, contributing to trophic strategies and microhabitat (Alve, 2010; Enge *et al.*, 2011). While  $\text{CaCO}_3$  content could affect the preservation of their tests as well as allow the coexistence of multiple test wall types promote the expansion of functional space (de Nooijer *et al.*, 2014). Additionally,  $\delta^{15}\text{N}$  is considered a predictor only for functional richness, likely reflecting nitrogen cycling processes and its impact on functional traits. In the SEP, within the OMZ, low  $\delta^{15}\text{N}$  signal reflects the signature of intense sedimentary denitrification, which minimizes isotopic discrimination (Hoogakker *et al.*, 2025). Consequently, functional volume (FRic) is reduced, as only specialized traits adapted to these conditions can persist. Such metabolic strategies would be reflected in species previously mentioned (Glock, 2023; Koho & Piña-Ochoa, 2012).

On the other hand, abundance was associated with C/N ratio, highlighting the importance of organic matter quality in regulating benthic foraminiferal densities.

Zones off Peru and northern Chile, the degradation of nitrogen rich components increases C/N ratios above Redfield values during anaerobic microbial pathways and denitrification process (De Pol-Holz *et al.*, 2009; Flores *et al.*, 2023). These elevated C/N values (>7.5), indicate shifts in the lability and composition of organic matter reaching the seafloor. In this context, in the Peruvian OMZ core, the combined effect of diatom productivity and high organic matter export (Figure S1a, b and d), indicate that C/N values likely reflect the accumulation of relatively fresh phytodetritus, promoting high benthic foraminifera abundances (Cardich *et al.*, 2015). Additionally, those abundances can reflect reproductive responses to pulses of phytodetritus export (Nomaki *et al.*, 2008). The prevalence of *Bolivina* species within the OMZ core (Figure S2-S3) likely implies the ability to exploit a broad spectrum of organic substrates. This is consistent with their documented capacity off the coast of Japan to switch from fresh phytodetritus to degraded organic matter depending on temporal availability (Nomaki *et al.*, 2008), where input of terrigenous material from the continent to the shelf might be like the one in Chile. In contrast, similar C/N values along Chilean margin did not present the same foraminiferal density, suggesting that oxygenation state and organic matter flux regulate the population limiting total foraminiferal abundance ((Neira *et al.*, 2001). This agrees with previous work in the Southeast Pacific, where it has been shown that sedimentary organic matter quality regulates both the distribution and microhabitat of calcareous benthic foraminifera along the central Peruvian margin and northern Chile (Cardich *et al.*, 2015; Castillo *et al.*, 2021).

Among all predictors, oxygen was the only variable that explained variation in FEve and FDis, highlighting its role as environmental filter that impacts community abundance and specialization. Low oxygen concentrations modify food availability by favoring species with specific morphological and physiological adaptations to hypoxia (S Garrido *et al.*, 2025; Glock, 2023; Piña-Ochoa *et al.*, 2010). This limitation results in lower FDis (species with similar traits) and lower FEve (a few tolerant species dominate). Conversely, environments with superior oxygenation exhibit broader and more uniform functional traits.

Finally, our results reveal that FDiv and RaoQ were less sensitive to the measured environmental variables, suggesting that other biotic or unmeasured abiotic factors may influence the distribution of specialized traits. It should be noted that although the GAM analyses were restricted to the 55 stations (Appendix 2) with complete environmental coverage, the patterns detected represent strong relationships for the subset of sites where benthic habitat quality could be fully assessed.

### **8.5. Realized niche**

The The OMI analysis reveals a group of highly specialized taxa within the SEP, characterized by high marginality (OMI) and very low tolerance (Tol) values (Table 6) (Dolédec *et al.*, 2000). Species such as *Vaginulina americana* (Vamer), *Ebuliminella curta* (Ecurt), *Buliminella elegantissima limbosa* (Belim), *Buliminella elegantissima tenuis* (Belte) and *Rhumblerella humboldti* (Rhumb) occupy restricted portions of the environmental gradient, indicating strong ecological

specialization. The specific environmental conditions of Ecurt, Belim, Belte occur primarily in the Peruvian zone, at neritic and upper bathyal depths, which are influenced by high NO<sub>2</sub> and low oxygen (Figure 10 a, b). This coincides with previous reports explaining that *B. elegantissima* is associated with shallower environments characterized by high organic matter accumulation (Merma-Mora, 2020). In contrast, the marginality of Vamer, Bpago and Rhumb are evident in the equatorial region of the SEP (ESEP), at shallow depths characterized by higher C/N and warmer temperature. Previous studies have reported the occurrence of these species in similar geographic and bathymetric settings (Mallon, 2011); however, the OMI analysis applied in this study provides an interpretation of the observed distribution by identifying the specific environmental gradients. Similarly, other taxa such *Bathysiphon capillare*, *Martinotiella communis* and *Dorothia scabra* also exhibits a high marginality, at neritic to upper bathyal depths but in the Southern Chilean region. These results indicate a specific ecological specialization for each region within the SEP. Flexible specialist species combined high marginality with moderate to high tolerance (Table 6), indicating specialization within specific environmental settings but retaining ecological flexibility (Dolédec *et al.*, 2000). A representative example of this pattern is *B. seminuda* (Bsemi), which exhibited elevated tolerance (Tol = 5.22). Its niche optimum (Figure 10a) is consistent with previous reports describing the species associated with elevated TOC and nitrogen in suboxic to postoxic sediments (Cardich *et al.*, 2012). However, its broad latitudinal distribution (3°S–35°S)

suggests that *Bsemi* responds to fluctuating redox conditions rather than being restricted to the persistent OMZ core.

Intermediate species occupy transitional ecological positions along environmental gradients, reflected by moderate marginality and variable tolerance values (Dolédec *et al.*, 2000). *Bolivina costata* (*Bcost*) is within this category, indicating its realized niche located in upper bathyal environments off Ecuador and Peru (Figure 10a, b). Although previous coastal studies have described this species as dominant and tolerant of sulfidic conditions (Cardich *et al.*, 2012; Merma-Mora, 2016), our results suggest a narrower ecological amplitude, preferentially associated with dynamic suboxic environments rather than the persistently anoxic OMZ core. However, within this group, some species exhibited low *Tol* values, indicating that although their niche occurs under common environmental conditions, they occupied a narrow portion of the available gradient. *Uvigerina canariensis* (*Ucana*), *U. striata* (*Ustri*), *Labrospira robusta*, *Bolivinita minuta* and *Ehrenbergina compressa* are primarily associated with areas characterized by elevated TOC, C/N and opal content (Figure 10a, b) (Gutiérrez *et al.*, 2011), suggesting specialization where oxygen fluctuates near hypoxic thresholds. Conversely, *U. semiornata* (*Usemi*) and *Pseudobrivalina lobata* were linked to deeper bathyal settings with higher oxygen concentrations and salinity in Chile and Northern Peru, respectively (Figure 9a), despite occasional occurrences within OMZ influenced depths (Table S5). This suggests that their ecological optimum lies in more stable deep-water conditions rather than oxygen depleted

zones. Overall, taxa within this category reflect specialization in productive, low-oxygen or variable-salinity settings.

In contrast to specialist and intermediate taxa, generalist species occupy the average or "typical" environmental conditions of the study area, their broad tolerance suggests reduced sensitivity to environmental variability (Dolédec *et al.*, 2000; Hernández Fariñas *et al.*, 2015). Examples of this include *Hoeglundina elegans* (Heleg) and *Melonis affinis* (Maffi), present in lower bathyal environments off Chile (Figure 10a, b). Maffi is associated with elevated  $\delta^{15}\text{N}$  and total Nitrogen in the sediment, whereas Heleg is linked to high oxygen, greater depth and lower C/N ratios. Both species have been widely reported from well-oxygenated slope and deep-water settings (Bandy & Rodolfo, 1964; Gooday, 2003; Ingle *et al.*, 1980), supporting the low marginality and broad tolerance observed here. Additionally, Heleg is described as a shallow infaunal (SI) specie capable of responding to moderate variations in organic matter supply under oxic or suboxic conditions (Venturelli *et al.*, 2018), while Maffi is catalogued as transitional infaunal specie (Rathburn & Corliss, 1994). These traits support their classification as generalists and explain their wide latitudinal and bathymetric distribution (Figure S3, S4), including occurrences at OMZ-influenced depths atypical of their OMI optimum. It should be noted that the ecological categories (Table 6) represent relative positions along continuous marginality–tolerance gradients rather than fixed classifications.

Beyond the ecological categories, OMI results reveal niche differentiation within *Bolivina*. A direct comparison of Bsemi and Bcost further emphasizes species-

specific ecological strategies within the genus, given their occurrence in such conditions (Cardich *et al.*, 2012; Merma-Mora, 2016). While *B. interjuncta* and *B. plicata* are associated with suboxic conditions and organic-rich sediments (Cardich, 2012; Cardich *et al.*, 2019; Ingle *et al.*, 1980; Resig, 1981). This coincides with their generalist category and optimal niche in bathyal zones, as well as their restriction to deeper environments (Figure S2, S3), suggesting ecological flexibility constrained by depth. Overall, this partitioning suggests that species within *Bolivina* exploit similar redox–productivity systems through distinct ecological amplitudes rather than identical niche optima. Similar patterns have been documented in other oxygen-depleted systems (Fontanier *et al.*, 2016; Gooday *et al.*, 2008; Tetard *et al.*, 2024). A comparable pattern is observed within *Uvigerina*, where species display differentiation primarily along depth and oxygen gradients (Ingle *et al.*, 1980; Resig, 1981). Recent findings identify *Ucana* associated with dysoxic conditions, whereas *Uaub* and *U. peregrina* are linked to suboxic environments (Tetard *et al.* 2024). *U. peregrina* has also been described as part of transitional assemblages outside the OMZ core (Erdem *et al.*, 2020), which may explain the absence of a significant OMI signal in the present study. Finally, OMI analysis of functional traits revealed a distinct niche among ecological strategies (Figure S6). Most of the functional traits have been discussed in the previous section; however, here we can highlight the optimum of microhabitat traits. Many SI taxa (uvigerinids, bolivinids) are associated with the utilization of low labile organic matter available (Mojtahid *et al.*, 2010). Epifaunal taxa (e.g. *Cancris*, *Cibicidoides*, *Eponides*, *Hoeglundina*, *Nutallides*) are linked by low C/N,

ratios, well-oxygenated and deeper environments (Figure S7), consistent with their preference for low organic carbon flux rate (Altenbach *et al.*, 1999; Singh *et al.*, 2022; Singh *et al.*, 2012). Similarly, agglutinated species (*Crithionina pisum*, *Lagenammia*, *Recurvoides*, *Rhabdammina abyssorum*, *Trochammia squammata*, *Saccammia sphaerica*) occurring in similar type of niche (Figure S7a, b) because they can feed on suspended particles (Mojtahid *et al.*, 2010; Schönfeld, 2002). On the other hand, intermediate infaunal (II) taxa are less dependent on fresh organic input and can utilize refractory material (Goldstein & Corliss, 1994; Gooday *et al.*, 1992), allowing persistence under suboxic conditions (Fontanier *et al.*, 2006). Some II species recorded here (*e.g.* *Chillostomella ovoidea*, *C. oolina*, *Cassidulina*, *Globobulimina* sp., *Melonis* sp.) occur deeper in the sediments since at deepest sites, the oxygen penetration depth increased (Mojtahid *et al.*, 2010). Nevertheless, high oxygen levels permit both shallow and intermediate infaunal traits to extend into bathyal and abyssal zones (Gooday *et al.*, 1992).

## 9. CONCLUSION

This study demonstrates that benthic foraminiferal communities in the Southeastern Pacific (SEP) are not distributed along a continuous latitudinal gradient but are structured by discrete environmental thresholds that decouple taxonomic and functional diversity. Oxygen availability acts as the dominant ecological filter, defining the breadth of viable strategies across the region. Under depleted oxygen conditions, the communities are restricted by a narrow set of viable strategies, where high abundances mask a contraction of functional space. In contrast, sites with higher

species richness may exhibit functional expansion, although this relationship is not consistent across the region. These patterns indicate that functional redundancy becomes more pronounced under low-oxygen conditions, whereas areas characterized by greater environmental heterogeneity allow diversification of trait combinations. These dynamics highlight the importance of oxygen availability as a key factor that modulates not only community composition but also the balance between redundancy and differentiation in benthic ecosystem functioning.

The consistency between generalized additive models and trait-based patterns reinforces oxygen as the principal structuring force, while water mass properties and sedimentary proxies further shape both taxonomic and functional structure. These findings indicate that oxygen constrains which strategies can persist, whereas resource availability regulates their relative abundance. Although environmental drivers were assessed using a subset of stations with complete data, the consistency of these relationships with independent taxonomic, functional, and niche patterns supports their robustness across the broader region.

Overall, the SEP margin represents a mosaic of environmentally filtered assemblages rather than a simple latitudinal gradient. Given the current oceanic deoxygenation, the expansion of OMZ conditions may reduce functional space and constrain the range of ecological responses available to benthic communities.

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## APPENDIXES.

Appendix 1. Geochemistry sediment dataset from M77, SO156 and RR9702A cruise in the Southeastern Pacific. The data for the M77 cruise was obtained from Mollier-Vogel [et al.](#) (2012) and Sommer [et al.](#) (2019). Data for the RR9702A cruise was obtained from Prahl [et al.](#) (2006). Data for the SO156 cruise was obtained from Romero & Hebbeln (2003).

\*Note about total nitrogen and total carbon are the mean of the data from Mollier-Vogel [et al.](#) (2012) and Sommer [et al.](#) (2019)

ID	Cruise	Station	Long	Lat	Depth (m)	Total Organic Carbon [wt%]	Nitrogen Total [wt%]	CaCO <sub>3</sub> [wt%]	C/N ratio	Opal [wt%]	δ <sup>15</sup> N bulk [per mil]
Site_1_207	M77/2	772	-81.12	-1.95	207	1	0.12	9.63	8.71	2.98	4.62
Site_1_526	M77/2	767	-82.62	-1.75	526	2.92	0.33	9.84	10.09	2.97	4.81
Site_1_2092	M77/2	776	-81.20	-1.89	2092	1.70	0.22	41.59	8.36	8.23	4.74
Site_3_350	M77/2	744	-81.12	-3.75	350	2.99	0.36	6.86	9.42	3.12	4.78
Site_3_700	M77/2	757	-81.26	-3.85	700	3.31	0.40	6.34	9.34	3.27	4.4
Site_3_995	M77/2	753	-81.32	-3.95	995	3.27	0.36	-	9.26	7.02	4.27
Site_7_359	M77/2	716	-80.34	-8.00	359	9.70	1.25	11.72	-	-	-
Site_7_627	M77/2	723	-80.52	-7.87	627	4.91	0.58	18.32	8.91	3.64	4.5
Site_9_115	M77/2	694	-79.45	-9.05	115	1.57	0.16	25.66	11.11	2.61	5.27
Site_9_437	M77/2	692	-79.28	-9.28	437	8.45	0.77	15.59	10.43	3.53	4.53
Site_9_1105	M77/2	684	-79.90	-9.29	1105	5.27	0.61	19.56	8.34	4.55	4.45
Site_10_521	M77/1	553	-78.90	-10.44	521	4.16	0.41	-	10.2	1.21	6.83
Site_10_1004	M77/1	549	-78.52	-11.00	1004	-	-	-	-	-	-
Site_10_1923	M77/2	669	-78.77	-10.89	1923	2.98	0.36	6.83	8.26	3.55	5.04
Site_11_79	M77/1	540	-77.79	-11.00	79	3.26	0.5	-	6.53	12.88	4.82
Site_11_145	M77/1	470	-77.94	-11.00	145	7.48	0.91	-	8.21	17.62	6.02
Site_11_211	M77/2	676	-78.02	-11.08	211	-	0.77	-	-	9.78	6.11
Site_11_248	M77/1	583	-78.05	-11.11	248	-	-	-	-	-	-
Site_11_291	M77/1	582	-78.08	-11.16	291	10.25	1.37	-	7.4	9.64	6.09
Site_11_317	M77/1	473	-78.17	-11.00	317	12.32	1.4	-	7.35	9.87	6.68
Site_11_319	M77/1	449	-78.17	-11.00	319	12.2	1.46	-	7.28	9.12	6.28
Site_11_375	M77/1	482	-78.24	-11.00	375	13.98	1.71	-	8.43	5.72	6.35
Site_11_465	M77/1	456	-78.32	-11.00	465	6.87	0.7	-	9.74	2.43	7.27
Site_11_511	M77/1	516	-78.33	-11.00	511	9.49	1.2	-	7.88	6.93	6.23
Site_11_579	M77/1	487	-78.39	-11.00	579	4.75	0.82	-	5.98	2.93	6.18
Site_11_697	M77/1	459	-78.43	-11.00	697	6.55	0.73	-	8.92	2.44	5.01
Site_12_152	M77/1	619	-77.32	-12.30	152	8.85	1.2	-	7.59	17.39	7.23
Site_12_214	M77/2	635	-77.67	-12.09	214	12.75	1.32	-	-	9.9	6.67
Site_12_302	M77/1	616	-77.48	-12.38	302	-	-	-	-	-	-
Site_12_823	M77/1	622	-77.58	-12.55	823	-	-	-	7.81	4.94	5.82
Site_15_522	M77/1	421	-75.58	-15.19	522	6.74	0.81	-	8.28	-	7.36
Site_16_2575	GenesisIII RR-9702a	66MC	-77.10	-16.13	2575	-	-	-	-	-	-
Site_16_3228	GenesisIII RR-9702a	68MC	-76.38	-16.07	3228	-	-	-	-	-	-
Site_16_4124	GenesisIII RR-9702a	70MC	-76.01	-16.73	4124	-	-	-	-	-	-
Site_17_298	M77/1	403	-71.86	-17.43	298	1.7	0.22	-	9.51	2.22	9.17
Site_17_492	M77/1	406	-71.87	-17.47	492	2.37	0.3	-	8.92	2.73	8.61
Site_17_918	M77/1	410	-71.97	-17.64	918	2.76	0.33	-	9.12	3.85	8.19
Site_23_523	Sonne156	71	-70.6	23.9998	523	3.34	0.38	24.24	8.81	1.42	11.77

Site_23_1395	Sonne156	7114-1	-70.83	-24.00	1395	1.91	6.35	16.73	6.35	2.73	12.25
Site_23_3396	GenesisIII RR-9702a	50MC	-73.61	-23.61	3396	0.27	-	63.4	-	-	-
Site_25_1442	Sonne156	7121-1	-70.90	-26.00	1442	1.09	7.89	12.74	7.89	0.86	11.88
Site_26_673	Sonne156	7122-2	-70.85	-26.01	673	0.95	7.85	14.63	7.85	0.42	11.08
Site_26_1996	Sonne156	7116-1	-71.00	-26.00	1996	0.73	7.31	15.9	7.31	1.16	-
Site_27_557	Sonne156	7123-1	-71.05	-27.29	557	1.35	8.62	9.4	8.62	1.42	10.35
Site_28_473	Sonne156	7129-1	-71.33	-28.42	476	2.09	7.36	19.4	7.36	0.67	10.73
Site_28_1462	Sonne156	7127-1	-71.47	-28.38	1462	0.66	7.45	8.93	7.45	1.11	10.92
Site_28_1650	Sonne156	7131-1	-71.50	-28.38	1650	0.63	7.47	8.73	7.47	0.92	-
Site_28_2080	Sonne156	7130-1	-71.61	-28.42	2080	0.49	7.63	8.9	7.63	0.72	-
Site_29_635	Sonne156	7133-1	-71.64	-29.38	635	1.33	7.65	12.0	7.65	1.08	11.11
Site_29_1432	Sonne156	7135-1	-71.68	-29.67	1432	1.57	7.44	13.54	7.44	2.07	11.49
Site_29_3252	Sonne156	7132-1	-71.89	-29.47	3252	1.25	6.66	16.3	6.66	3.3	11.95
Site_30_1199	Sonne156	7137-2	-71.73	-30.17	1199	1.73	7.42	10.41	7.42	2.02	11.08
Site_30_2733	Sonne156	7138-1	-71.87	-30.13	2733	1.48	7.05	-	7.05	3.98	11.23
Site_30_3267	Sonne156	7139-1	-71.98	-30.20	3267	1.41	7.14	9.53	7.14	4.18	-
Site_31_183	Sonne156	7146-1	-71.63	-31.97	183	-	-	-	-	-	-
Site_31_301	Sonne156	7141-1	-71.75	-31.09	301	1.30	8.02	32.7	8.02	0.1	10.3
Site_31_326	Sonne156	7145-1	-71.70	-31.88	326	1.15	9.23	13.72	9.23	0.24	-
Site_31_398	Sonne156	7147-1	-71.67	-31.98	398	0.93	8.05	7.73	8.05	0.53	10.06
Site_31_481	Sonne156	7142-2	-71.75	-31.18	481	1.07	7.63	8.26	7.63	0.8	10.52
Site_31_1371	Sonne156	7143-1	-71.88	-31.11	1371	-	-	-	-	-	-
Site_31_2289	Sonne156	7148-1	-71.93	-31.97	2289	0.91	7.14	13.73	7.14	1.83	-
Site_31_3086	Sonne156	7149-1	-72.00	-31.49	3086	1.06	6.83	10.62	6.83	3.13	-
Site_32_1591	Sonne156	7150-1	-71.95	-32.28	1591	0.63	7.18	8.67	7.18	0.98	11.07
Site_32_3920	GenesisIII RR-9702a	48 MC5	-73.65	-32.59	3920	0.56	-	23.3	-	-	-
Site_33_420	Sonne156	7152-1	-72.11	-33.80	420	1.13	7.65	2.66	7.65	2.38	10.16
Site_33_863	Sonne156	7153-1	-72.16	-33.80	863	1.60	7.3	4.58	7.3	2.65	10.41
Site_33_1385	Sonne156	7154-2	-72.27	-33.80	1385	2.18	7.78	4.4	7.78	2.87	10.63
Site_33_3852	GenesisIII RR-9702a	46 MC4	-73.53	-33.28	3852	0.94	-	7.8	-	-	-
Site_34_1247	Sonne156	7156-1	-72.51	-34.58	1247	2.04	7.95	2.71	7.95	3.97	10.36
Site_35_506	Sonne156	7159-1	-73.23	-35.78	506	0.74	8.9	0.28	8.9	1.26	9.68
Site_35_1563	Sonne156	7158-1	-73.48	-35.78	1563	2.76	8.02	1.42	8.02	5.94	10.11
Site_35_2103	Sonne156	7157-1	-73.59	-35.78	2103	2.76	8	1.09	8	5.81	10.13
Site_36_366	Sonne156	7160-4	-73.07	-36.04	366	2.97	7.89	0.95	7.89	5.01	10
Site_36_538	Sonne156	7163-5	-73.91	-36.46	538	-	-	-	-	-	-
Site_36_1294	Sonne156	7166-3	-73.77	-36.47	1294	3.08	7.92	0.64	7.92	5.34	10.32
Site_36_2064	Sonne156	7167-4	-73.91	-36.45	2064	3.04	7.73	0.48	7.73	7.04	10.52
Site_37_124	Sonne156	7205-1	-73.73	-38.00	124	-	-	-	-	-	-
Site_37_1386	Sonne156	7171-2	-73.95	-37.40	1386	2.90	7.95	1.52	7.95	6.68	10.39
Site_38_281	Sonne156	7203-1	-73.96	-38.04	281	0.77	8.79	0.26	8.79	0.9	9.75
Site_38_700	Sonne156	7201-1	-74.06	-38.09	700	-	-	-	-	-	-
Site_38_2287	Sonne156	7198-1	-74.39	-38.17	2287	2.49	8.08	0.87	8.08	6.5	10.27
Site_39_112	Sonne156	7220-1	-73.90	-39.77	112	-	-	-	-	-	-
Site_39_174	Sonne156	7217-1	-73.93	-39.99	174	-	-	-	-	-	-
Site_39_498	Sonne156	7215-1	-74.00	-39.48	498	1.40	8.02	2.04	8.02	2.83	10.22
Site_39_638	Sonne156	7218-1	-73.88	-39.91	638	2.74	7.98	2.36	7.98	6.51	9.9
Site_39_772	Sonne156	7214-1	-74.17	-39.88	772	-	-	-	-	-	-
Site_39_890	Sonne156	7211-1	-74.27	-39.93	890	1.28	8.19	3.72	8.19	2.56	10.58
Site_39_1190	Sonne156	7213-1	-74.29	-39.73	1190	-	-	-	-	3.14	-
Site_39_1469	Sonne156	7212-1	-74.38	-39.70	1469	-	-	-	-	3.74	-
Site_39_1798	Sonne156	7208-1	-74.21	-39.01	1794	-	-	-	-	4.7	-
Site_39_2236	Sonne156	7207-1	-74.37	-39.07	2236	2.34	7.71	2.62	7.71	4.92	10.67
Site_40_165	Sonne156	7216-1	-73.94	-40.04	165	0.73	8.69	0.49	8.69	1.88	9.73
Site_40_816	Sonne156	7197-1	-73.94	-41.00	816	2.35	7.85	4.57	7.85	4.6	-

Site_40_3850	GenesisIII RR-9702a	27 MC5	-75.92	-40.48	3850	1.2	-	0.4	-	-	-
Site_41_308	Sonne156	7194-1	-74.43	-41.42	308	1.79	8.25	7.01	8.25	1.89	10.04
Site_41_521	Sonne156	7195-1	-74.41	-41.21	521	1.55	7.98	4.35	7.98	2.51	10.17
Site_42_160	Sonne156	7173-5	-74.56	-42.09	160	1.34	8.31	3.54	8.31	1.68	9.79
Site_42_206	Sonne156	7193-1	-74.71	-42.18	206	-	-	-	-	-	-
Site_42_299	Sonne156	7172-5	-74.79	-42.41	299	-	-	-	-	-	-
Site_42_301	Sonne156	7182-1	-74.84	-42.58	301	-	-	-	-	-	-
Site_42_909	Sonne156	7177-1	-74.84	-42.59	909	3.53	8.6	3.39	8.6	4.24	9.59
Site_42_1967	Sonne156	7181-1	-74.79	-42.41	1967	3.12	10.19	7.63	10.19	4.21	9.75
Site_42_2760	Sonne156	7175-4	-75.34	-42.57	2760	2.51	8.09	4.62	8.09	5.35	9.62
Site_43_1212	Sonne156	7179-1	-75.21	-42.45	1222	3.40	8.02	7.39	8.02	5.79	9.78
Site_42_1222	Sonne156	7174-2	-75.28	-43.37	1212	1.52	8.24	5.45	8.24	3.03	9.22
Site_43_3485	Sonne156	7180-1	-75.57	-43.37	3485	1.68	7.83	2.58	7.83	3.7	9.67
Site_43_3523	GenesisIII RR-9702a	12 MC4	-76.25	-43.42	3523	1.3	-	1.8	-	-	-
Site_44_476	Sonne156	7187-1	-75.17	-44.20	476	1.32	7.5	2.9	7.5	2.56	9.22
Site_44_868	Sonne156	7189-1	-75.59	-44.28	868	1.58	7.81	4.73	7.81	3.6	9.07
Site_44_1171	Sonne156	7186-1	-75.16	-44.15	1171	2.37	7.5	5.91	7.5	4.74	9.29
Site_44_1939	Sonne156	7191-1	-75.17	-44.20	1939	1.54	7.61	6.05	7.61	3.75	9.37
Site_44_3285	Sonne156	7190-1	-75.39	-44.28	3285	1.32	7.5	3.44	7.5	4.24	9.29
Site_46_3014	GenesisIII RR-9702a	8 MC8	-76.67	-46.35	3014	5.4	-	1	-	-	-
Site_50_3964	GenesisIII RR-9702a	1 MC5	-76.96	-50.65	3964	0.29	-	5.6	-	-	-

## Appendix 2. Dataset of environmental variables used for the GAM and OMI analyses.

\*: No data

ID	Longitude	Latitude	Depth	Temp	Salinity	Oxygen	Phosphate	Nitrate	Nitrite	Silicate	TOC	TN	CaCO <sub>3</sub>	C/N	Opal	δ15N
Site 1 207	-81.12	-1.95	207	13.23	34.93	27.78	1.84	22.63	0.07	17.73	1.01	0.12	9.63	8.71	2.98	4.62
Site 1 526	-81.20	-1.89	526	8.45	34.65	15.48	2.75	33.17	0.06	42.61	2.92	0.33	9.84	10.09	2.97	4.81
Site 1 2092	-82.62	-1.75	2092	2.20	34.66	100.83	2.81	37.45	0.16	134.96	1.70	0.22	41.59	8.36	8.23	4.74
Site 3 350	-81.12	-3.75	350	10.23	34.75	5.96	2.74	35.62	0.02	35.50	2.99	0.355	6.86	9.42	3.12	4.78
Site 3 700	-81.26	-3.85	700	6.24	34.58	37.98	3.12	44.12	0.04	64.23	3.31	0.40	6.34	9.34	3.27	4.4
Site 3 995	-81.32	-3.95	995	4.86	34.57	55.76	3.07	43.77	0.00	86.85	3.27	0.36	*	9.26	7.02	4.27
Site 7 359	-80.34	-8.00	359	10.26	34.75	2.36	2.77	32.27	0.44	32.28	9.70	1.25	11.72	*	*	*
Site 7 627	-80.52	-7.87	627	7.09	34.60	9.09	3.13	44.06	0.08	51.56	4.91	0.58	18.32	8.91	3.64	4.5
Site 9 115	-79.45	-9.05	115	14.44	35.02	4.61	2.47	24.31	0.17	21.46	1.57	0.16	25.66	11.11	2.61	5.27
Site 9 437	-79.28	-9.28	437	9.00	34.67	2.23	2.99	38.14	1.12	35.40	8.45	0.77	15.59	10.43	3.53	4.53
Site 9 1105	-79.90	-9.29	1105	4.32	34.56	56.26	3.25	45.63	0.02	68.41	5.27	0.61	19.56	8.34	4.55	4.45
Site 10 521	-78.90	-10.44	521	8.07	34.62	3.05	3.24	37.82	0.05	42.39	4.16	0.41	*	10.2	1.21	6.83
Site 10 1004	-78.52	-11.00	1004	4.50	34.55	49.19	3.37	49.23	0.07	69.69	*	*	*	*	*	*
Site 10 1923	-78.77	-10.89	1923	2.38	34.64	97.15	2.91	42.06	0.14	112.06	2.98	0.36	6.83	8.26	3.55	5.04
Site 11 79	-77.79	-11.00	79	13.64	34.96	2.30	2.73	31.74	0.40	22.95	3.26	0.5	*	6.53	12.88	4.82
Site 11 145	-77.94	-11.00	145	12.88	34.92	2.53	2.58	26.88	0.80	27.08	7.48	0.91	*	8.21	17.62	6.02
Site 11 211	-78.02	-11.08	211	12.10	34.87	2.14	2.49	25.09	1.12	24.90	*	0.77	*	*	9.78	6.11
Site 11 248	-78.05	-11.11	248	11.79	34.85	2.20	2.52	25.33	1.30	24.39	*	*	*	*	*	*
Site 11 291	-78.08	-11.16	291	11.31	34.82	2.31	2.61	26.11	2.04	24.73	10.25	1.37	*	7.4	9.64	6.09
Site 11 317	-78.17	-11.00	317	10.79	34.78	2.44	2.68	27.00	2.40	25.64	12.32	1.4	*	7.35	9.87	6.68
Site 11 319	-78.17	-11.00	319	10.79	34.78	2.44	2.68	27.00	2.40	25.64	12.2	1.46	*	7.28	9.12	6.28
Site 11 375	-78.24	-11.00	375	9.58	34.70	2.67	2.82	28.84	2.42	29.52	13.98	1.71	*	8.43	5.72	6.35
Site 11 465	-78.32	-11.00	465	8.43	34.64	3.67	3.01	31.07	1.01	36.82	6.87	0.7	*	9.74	2.43	7.27
Site 11 511	-78.33	-11.00	511	7.98	34.62	4.37	3.16	31.94	0.41	41.31	9.49	1.2	*	7.88	6.93	6.23
Site 11 579	-78.39	-11.00	579	7.24	34.58	6.34	3.41	36.28	0.32	47.82	4.75	0.82	*	5.98	2.93	6.18
Site 11 697	-78.43	-11.00	697	6.40	34.56	11.41	3.50	44.52	0.28	55.59	6.55	0.73	*	8.92	2.44	5.01
Site 12 152	-77.32	-12.30	152	13.17	34.94	2.44	2.61	16.75	4.47	26.40	8.85	1.2	*	7.59	17.39	7.23
Site 12 214	-77.67	-12.09	214	12.91	34.92	1.35	2.60	20.17	4.93	25.55	12.754	1.323	7.69	*	9.9	6.67
Site 12 302	-77.48	-12.38	302	11.35	34.82	3.15	2.58	23.17	5.76	26.36	*	*	*	*	*	*
Site 12 823	-77.58	-12.55	823	5.86	34.55	16.46	3.25	44.90	0.08	60.57	*	*	*	7.81	4.94	5.82
Site 15 522	-75.58	-15.19	522	7.57	34.57	14.04	3.02	39.58	0.12	40.70	6.74	0.81	*	8.28	*	7.36
Site 16 2575	-77.10	-16.13	2575	1.87	34.67	135.04	2.67	26.78	0.45	88.88	1.6	*	6.4	*	*	*
Site 16 3228	-76.38	-16.07	3228	1.78	34.68	144.88	2.79	32.61	0.18	107.71	1.6	*	0	*	*	*
Site 16 4124	-76.01	-16.73	4124	1.78	34.70	148.53	2.79	36.93	0.05	123.53	0.67	*	0	*	*	*
Site 17 298	-71.86	-17.43	298	9.92	34.70	2.71	2.89	27.03	0.61	25.88	1.7	0.22	*	9.51	2.22	9.17
Site 17 492	-71.87	-17.47	492	7.25	34.54	23.81	3.20	43.65	0.20	29.79	2.37	0.3	*	8.92	2.73	8.61
Site 17 918	-71.97	-17.64	918	4.48	34.52	59.17	3.30	47.52	0.01	56.39	2.76	0.33	*	9.12	3.85	8.19
Site 23 523	-70.60	-24.00	523	7.37	34.51	29.57	2.30	31.94	0.04	25.45	3.34	0.38	24.24	8.81	1.42	11.77
Site 23 3396	-73.61	-23.61	3396	1.75	34.68	134.07	2.46	26.62	0.08	69.95	0.27	*	63.4	*	*	*
Site 25 1442	-70.90	-26.00	1442	2.96	34.57	98.99	2.83	39.40	0.07	79.28	1.09	7.89	12.74	7.89	0.86	11.88
Site 26 673	-70.85	-26.01	673	5.94	34.46	94.95	2.57	42.39	0.03	28.95	0.95	7.85	14.63	7.85	0.42	11.08
Site 26 1996	-71.00	-26.00	1996	2.24	34.63	113.61	2.62	34.26	0.22	89.63	0.73	7.31	15.9	7.31	1.16	*
Site 27 557	-71.05	-27.29	557	6.97	34.44	69.44	2.67	43.97	0.03	24.85	1.35	8.62	9.4	8.62	1.42	10.35
Site 28 473	-71.33	-28.42	476	7.78	34.49	5.31	2.74	42.30	0.06	23.07	2.09	7.36	19.4	7.36	0.67	10.73
Site 28 1462	-71.47	-28.38	1462	2.95	34.57	103.26	2.81	39.89	0.04	88.61	0.66	7.45	8.93	7.45	1.11	10.92
Site 28 1650	-71.50	-28.38	1650	2.63	34.60	109.90	2.74	38.35	0.09	95.85	0.63	7.47	8.73	7.47	0.92	*
Site 28 2080	-71.61	-28.42	2080	2.16	34.64	124.06	2.59	34.91	0.14	102.38	0.49	7.63	8.9	7.63	0.72	*
Site 29 635	-71.64	-29.38	635	5.96	34.43	76.46	2.52	45.07	0.00	26.31	1.33	7.65	12.89	7.65	1.08	11.11
Site 29 1432	-71.68	-29.67	1432	3.00	34.57	104.97	2.81	40.18	0.01	91.28	1.57	7.44	13.54	7.44	2.07	11.49
Site 29 3252	-71.89	-29.47	3252	1.75	34.68	144.27	2.45	32.40	0.07	105.87	1.25	6.66	16.3	6.66	3.3	11.95
Site 30 1199	-71.73	-30.17	1199	3.52	34.53	96.98	2.86	41.71	0.03	78.35	1.73	7.42	10.41	7.42	2.02	11.08
Site 30 2733	-71.87	-30.13	2733	1.83	34.67	144.22	2.47	33.53	0.06	111.30	1.48	7.05	*	7.05	3.98	11.23
Site 30 3267	-71.98	-30.20	3267	1.75	34.68	146.16	2.46	33.34	0.04	112.16	1.41	7.14	9.53	7.14	4.18	*
Site 31 183	-71.63	-31.97	183	10.84	34.60	13.87	2.66	24.61	0.08	19.27	*	*	*	*	*	*
Site 31 301	-71.75	-31.09	301	9.52	34.62	9.02	2.68	28.52	0.15	23.24	1.30	8.02	32.7	8.02	0.1	10.3
Site 31 326	-71.70	-31.88	326	9.15	34.58	16.23	2.69	32.27	0.04	23.67	1.15	9.23	13.72	9.23	0.24	*
Site 31 398	-71.67	-31.98	398	8.14	34.52	33.28	2.60	35.07	0.03	22.81	0.93	8.05	7.73	8.05	0.53	10.06
Site 31 481	-71.75	-31.18	481	7.03	34.47	60.19	2.40	38.03	0.03	20.74	1.07	7.63	8.26	7.63	0.8	10.52
Site 31 1371	-71.88	-31.11	1371	3.11	34.56	105.86	2.82	40.16	0.02	90.89	*	*	*	*	*	*
Site 31 2289	-71.93	-31.97	2289	2.00	34.65	140.96	2.55	36.01	0.04	120.16	0.91	7.14	13.73	7.14	1.83	*
Site 31 3086	-72.00	-31.49	3086	1.79	34.68	148.49	2.47	34.79	0.03	121.42	1.06	6.83	10.62	6.83	3.13	*
Site 32 1591	-71.95	-32.28	1591	2.69	34.59	116.73	2.72	38.47	0.00	102.98	0.63	7.18	8.67	7.18	0.98	11.07
Site 32 3920	-73.65	-32.59	3920	1.72	34.69	160.07	2.43	34.83	0.01	123.86	0.56	*	23.3	*	*	*
Site 33 420	-72.11	-33.80	420	7.65	34.45	55.70	2.88	34.29	0.01	22.88	1.13	7.65	2.66	7.65	2.38	10.16
Site 33 863	-72.16	-33.80	863	4.41	34.37	136.52	2.52	33.87	0.02	41.92	1.60	7.3	4.58	7.3	2.65	10.41
Site 33 1385	-72.27	-33.80	1385	3.04	34.55	112.76	2.74	37.82	0.03	90.73	2.18	7.78	4.4	7.78	2.87	10.63
Site 33 3852	-73.53	-33.28	3852	1.73	34.69	160.77	2.43	34.99	0.03	125.42	0.94	*	7.8	*	*	*
Site 34 1247	-72.51	-34.58	1247	3.31	34.51	111.58	2.70	36.15	0.05	78.55	2.04	7.95	2.71	7.95	3.97	10.36
Site 35 506	-73.23	-35.78	506	6.20	34.33	158.69	2.11	29.89	0.04	19.26	0.74	8.9	0.28	8.9	1.26	9.68
Site 35 1563	-73.48	-35.78	1563	2.75	34.58	118.15	2.68	36.74	0.08	97.01	2.76	8.02	1.42	8.02	5.94	10.11
Site 35 2103	-73.59	-35.78	2103	2.21	34.64	130.14	2.61	37.32	0.04	112.19	2.76	8	1.09	8	5.81	10.13

Site 36 366	-73.07	-36.04	366	8.01	34.46	51.19	2.57	35.00	0.04	24.27	2.97	7.89	0.95	7.89	5.01	10
Site 36 538	-73.91	-36.46	538	5.74	34.30	194.43	1.95	29.22	0.04	18.20	*	0.263	*	*	*	*
Site 36 1294	-73.77	-36.47	1294	3.19	34.51	117.67	2.60	33.84	0.12	76.22	3.08	7.92	0.64	7.92	5.34	10.32
Site 36 2064	-73.91	-36.45	2064	2.25	34.64	127.67	2.62	37.36	0.08	111.00	3.04	7.73	0.48	7.73	7.04	10.52
Site 37 124	-73.73	-38.00	124	10.26	34.23	87.64	1.84	25.48	0.04	19.90	*	*	*	*	*	*
Site 37 1386	-73.95	-37.40	1386	3.00	34.53	119.18	2.61	34.24	0.15	81.79	2.90	7.95	1.52	7.95	6.68	10.39
Site 38 281	-73.96	-38.04	281	8.81	34.50	46.86	2.35	31.24	0.04	23.31	0.77	8.79	0.26	8.79	0.9	9.75
Site 38 700	-74.06	-38.09	700	4.77	34.27	198.23	1.86	24.97	0.06	20.13	*	*	*	*	*	*
Site 38 2287	-74.39	-38.17	2287	2.13	34.66	128.29	2.60	37.22	0.19	116.05	2.49	8.08	0.87	8.08	6.5	10.27
Site 39 112	-73.90	-39.77	112	10.15	34.05	139.50	1.19	15.42	0.05	9.76	0.586	0.064	0.1	9.18	1.49	9.59
Site 39 174	-73.93	-39.99	174	9.41	34.34	68.81	1.82	24.26	0.02	14.10	*	*	*	*	*	*
Site 39 498	-74.00	-39.48	498	5.95	34.28	183.64	1.61	23.34	0.03	13.03	1.40	8.02	2.04	8.02	2.83	10.22
Site 39 772	-74.17	-39.88	772	4.49	34.28	175.81	1.78	21.91	0.12	20.41	*	*	*	*	*	*
Site 39 890	-74.27	-39.93	890	4.08	34.33	153.53	1.93	22.36	0.15	28.47	1.28	8.19	3.72	8.19	2.56	10.58
Site 39 1190	-74.29	-39.73	1190	3.30	34.46	132.39	2.39	29.58	0.20	59.53	*	*	*	*	3.14	*
Site 39 1469	-74.38	-39.70	1469	2.85	34.55	120.23	2.62	34.70	0.24	84.95	*	*	*	*	3.74	*
Site 39 1798	-74.21	-39.01	1794	2.48	34.61	118.41	2.67	36.99	0.21	103.12	*	*	*	*	4.7	*
Site 40 165	-73.94	-40.04	165	9.49	34.31	76.06	1.75	23.54	0.02	13.57	0.73	8.69	0.49	8.69	1.88	9.73
Site 40 816	-73.94	-41.00	816	4.32	34.29	160.56	1.79	20.96	0.17	21.70	2.35	7.85	4.57	7.85	4.6	*
Site 40 3850	-75.92	-40.48	3850	2.06	34.72	162.49	2.39	33.07	0.65	116.84	1.20	*	0.4	*	*	*
Site 41 308	-74.43	-41.42	308	7.85	34.37	83.97	1.85	18.47	0.08	11.68	1.79	8.25	7.01	8.25	1.89	10.04
Site 41 521	-74.41	-41.21	521	5.78	34.25	183.37	1.56	18.75	0.08	10.62	1.55	7.98	4.35	7.98	2.51	10.17
Site 42 160	-74.56	-42.09	160	9.19	34.26	105.67	1.32	17.81	0.14	8.68	1.34	8.31	3.54	8.31	1.68	9.79
Site 42 206	-74.71	-42.18	206	8.69	34.36	77.21	1.71	21.05	0.13	10.99	*	*	*	*	*	*
Site 42 299	-74.79	-42.41	299	7.75	34.35	91.18	1.80	17.90	0.17	10.52	1.56	0.175	4.72	8.89	1.56	10
Site 42 301	-74.84	-42.58	301	7.72	34.35	92.06	1.80	18.01	0.17	10.40	*	*	*	*	*	*
Site 42 909	-74.84	-42.59	909	4.03	34.32	141.52	1.93	21.56	0.23	26.72	3.53	8.6	3.39	8.6	4.24	9.59
Site 42 1222	-75.28	-43.37	1212	3.17	34.46	129.37	2.49	32.90	0.27	62.25	1.52	8.24	5.45	8.24	3.03	9.22
Site 42 1967	-74.79	-42.41	1967	2.30	34.63	116.19	2.66	37.27	0.40	108.36	3.12	10.19	7.63	10.19	4.21	9.75
Site 42 2760	-75.34	-42.57	2760	2.05	34.70	127.26	2.50	35.46	0.59	119.66	2.51	8.09	4.62	8.09	5.35	9.62
Site 43 1212	-75.21	-42.45	1222	3.17	34.47	129.69	2.46	31.83	0.26	62.57	3.40	8.02	7.39	8.02	5.79	9.78
Site 43 3523	-76.25	-43.42	3523	2.14	34.73	155.75	2.40	32.57	0.85	113.00	1.30	*	1.8	*	*	*
Site 44 476	-75.17	-44.20	476	5.87	34.24	164.59	1.61	18.37	0.15	8.70	1.32	7.5	2.9	7.5	2.56	9.22
Site 44 868	-75.59	-44.28	868	4.14	34.29	145.95	2.07	24.64	0.17	25.94	1.58	7.81	4.73	7.81	3.6	9.07
Site 44 1171	-75.16	-44.15	1171	3.23	34.44	130.39	2.48	33.28	0.25	58.69	2.37	7.5	5.91	7.5	4.74	9.29
Site 44 1939	-75.17	-44.20	1939	2.28	34.63	116.04	2.66	37.27	0.43	107.87	1.54	7.61	6.05	7.61	3.75	9.37
Site 44 3285	-75.39	-44.28	3285	2.13	34.73	142.26	2.41	32.94	0.84	113.57	1.32	7.5	3.44	7.5	4.24	9.29
Site 46 3014	-76.67	-46.35	3014	2.13	34.73	120.85	2.42	32.90	0.85	111.31	1.00	*	5.4	*	*	*
Site 50 3964	-76.96	-50.65	3964	2.64	34.77	79.83	2.28	26.75	1.39	84.30	1.25	*	0.29	*	*	*

### Appendix 3. List of living benthic foraminifera species present in the Southeastern Pacific Ocean

Class	Order	Family	Genus	Specie	Authority
Monothalamea	Astrorhizida	Astrorhizidae	Astrorhiza	<i>Astrorhiza arenaria</i>	Carpenter in Norman, 1877
		Astrorhizidae	Astrorhiza	<i>Astrorhiza granulosa</i>	(Brady, 1879)
		Astrorhizidae	Astrorhizoides	<i>Astrorhizoides cornutus</i>	(Brady, 1879)
		Crithioninidae	Crithionina	<i>Crithionina pisum</i>	Goës, 1896
		Dendrophryidae	Psammatodendron	<i>Psammatodendron arborescens</i>	Norman in Brady, 1881
		Hyperamminidae	Hyperammina	<i>Hyperammina</i>	Brady, 1878
		Hyperamminidae	Hyperammina	<i>Hyperammina echinata</i>	Saidova, 1970
		Hyperamminidae	Hyperammina	<i>Hyperammina elongata</i>	Brady, 1878
		Hyperamminidae	Hyperammina	<i>Hyperammina friabilis</i>	Brady, 1884
		Hyperamminidae	Saccorhiza	<i>Saccorhiza ramosa</i>	(Brady, 1879)
		Psammosphaeridae	Psammosphaera	<i>Psammosphaera fusca</i>	Schulze, 1875
Rhabdamminidae	Bathysiphon	<i>Bathysiphon capillare</i>	Folin, 1886		

		Rhabdamminidae	Bathysiphon	<i>Bathysiphon fusca</i>	Cushman, 1927
		Rhabdamminidae	Bathysiphon	<i>Bathysiphon</i>	Sars, 1872
		Rhabdamminidae	Marsipella	<i>Marsipella elongata</i>	Norman, 1878
		Rhabdamminidae	Psammosiphonella	<i>Psammosiphonella discreta</i>	(Brady, 1881)
		Rhabdamminidae	Rhabdammina	<i>Rhabdammina abyssorum</i>	Sars in Carpenter, 1869
		Rhabdamminidae	Rhabdammina	<i>Rhabdammina pacifica</i>	Shchedrina, 1952
		Rhabdamminidae	Rhabdammina	<i>Rhabdammina sp.</i>	Sars in Carpenter, 1869
		Rhabdamminidae	Rhabdamminella	<i>Rhabdamminella cylindrica</i>	(Brady, 1882)
		Rhizamminidae	Rhizammina	<i>Rhizammina algaeformis</i>	Brady, 1879
		Rhizamminidae	Rhizammina	<i>Rhizammina</i>	Brady, 1879
		Saccamminidae	Astrammmina	<i>Astrammmina rara</i>	Rhumbler, 1931
		Saccamminidae	Lagenammina	<i>Lagenammina ampullacea</i>	(Brady, 1881)
		Saccamminidae	Lagenammina	<i>Lagenammina difflugiformis</i>	(Brady, 1879)
		Saccamminidae	Saccammina	<i>Saccammina sphaerica</i>	Brady, 1871
		Saccamminidae	Technitella	<i>Technitella</i>	Norman, 1878
		Saccamminidae	Technitella	<i>Technitella melo</i>	Norman, 1878
		Saccamminidae	Thurammmina	<i>Thurammmina albicans</i>	Brady, 1879
Globotalamea	Lituolida	Adercotrymidae	Adercotryma	<i>Adercotryma glomeratum</i>	(Brady, 1878)
		Ammomassiliniidae	Ammomassilina	<i>Ammomassilina alveoliniformis</i>	(Millett, 1898)
		Ammosphaeroidinidae	Ammosphaeroidina	<i>Ammosphaeroidina grandis</i>	Cushman, 1910
		Ammosphaeroidinidae	Cribrostomoides	<i>Cribrostomoides evolutus</i>	(Natland, 1938)
		Ammosphaeroidinidae	Cribrostomoides	<i>Cribrostomoides jeffreysii</i>	(Williamson, 1858)
		Ammosphaeroidinidae	Cribrostomoides	<i>Cribrostomoides nitidus</i>	(Goës, 1896)
		Ammosphaeroidinidae	Cribrostomoides	<i>Cribrostomoides sphaerilocula</i>	(Cushman, 1910)
		Ammosphaeroidinidae	Cribrostomoides	<i>Cribrostomoides subglobosus</i>	(Cushman, 1910)
		Ammosphaeroidinidae	Recurvoidella	<i>Recurvoidella bradyi</i>	(Robertson, 1891)
		Ammosphaeroidinidae	Recurvoides	<i>Recurvoides contortus</i>	Earland, 1934
		Ammosphaeroidinidae	Recurvoides	<i>Recurvoides turbinatus</i>	(Brady, 1881)
		Discamminidae	Glaphyrammina	<i>Glaphyrammina americana</i>	(Cushman, 1910)
		Haplophragmoididae	Buzasina	<i>Buzasina ringens</i>	(Brady, 1879)
		Haplophragmoididae	Buzasina	<i>Buzasina sp.</i>	Loeblich & Tappan, 1985
		Haplophragmoididae	Evolutinella	<i>Evolutinella rotulata</i>	(Brady, 1881)

	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides bulloides</i>	Beissel, 1886
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides mexicanus</i>	Kornfeld, 1931
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides planissima</i>	Cushman, 1927
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides pusillus</i>	Collins, 1974
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides sp.</i>	Cushman, 1910
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides subsphaeroides</i>	Subbotina, 1950
	Haplophragmoididae	Haplophragmoides	<i>Haplophragmoides wilberti</i>	Andersen, 1953
	Haplophragmoididae	Labrospira	<i>Labrospira crassimargo</i>	(Norman, 1892)
	Haplophragmoididae	Labrospira	<i>Labrospira robusta</i>	Cushman & McCulloch, 1948
	Haplophragmoididae	Veleroninoides	<i>Veleroninoides scitulus</i>	(Brady, 1881)
	Lituolidae	Ammobaculites	<i>Ammobaculites agglutinans</i>	(d'Orbigny, 1846)
	Lituolidae	Ammomarginulina	<i>Ammomarginulina catenulata</i>	(Cushman & McCulloch, 1939)
	Lituolidae	Eratidus	<i>Eratidus foliaceus</i>	(Brady, 1881)
	Prolixoplectidae	Eggerelloides	<i>Eggerelloides advena</i>	(Cushman, 1922)
	Prolixoplectidae	Eggerelloides	<i>Eggerelloides scabrum</i>	(Williamson, 1858)
	Sigmoilopsidae	Spirosigmoilinella	<i>Spirosigmoilinella</i>	Matsunaga, 1955
	Spiroplectamminidae	Spiroplectammina	<i>Spiroplectammina biformis</i>	(Parker & Jones, 1865)
	Trochamminidae	Ammoglobigerina	<i>Ammoglobigerina globigeriniformis</i>	(Parker & Jones, 1865)
	Trochamminidae	Ammoglobigerina	<i>Ammoglobigerina globulosa</i>	(Cushman, 1920)
	Trochamminidae	Paratrochammina	<i>Paratrochammina challengerii</i>	Brönnimann & Whittaker, 1988
	Trochamminidae	Paratrochammina	<i>Paratrochammina globorotaliformis</i>	(Zheng, 1988)
	Trochamminidae	Polystomammina	<i>Polystomammina nitida</i>	(Brady, 1881)
	Trochamminidae	Portatrochammina	<i>Portatrochammina karica</i>	(Shchedrina, 1946)
	Trochamminidae	Portatrochammina	<i>Portatrochammina murrayi</i>	Brönnimann & Zaninetti, 1984
	Trochamminidae	Portatrochammina	<i>Portatrochammina pacifica</i>	(Cushman, 1925)
	Trochamminidae	Trochammina	<i>Trochammina inflata</i>	(Montagu, 1808)
	Trochamminidae	Trochammina	<i>Trochammina intermedia</i>	Rhumbler, 1938
	Trochamminidae	Trochammina	<i>Trochammina nana</i>	(Brady, 1881)
	Trochamminidae	Trochammina	<i>Trochammina</i>	Parker & Jones, 1859

Globothalamea		Trochamminidae	Trochammina	<i>Trochammina squamata</i>	Jones & Parker, 1860
		Trochamminidae	Trochammina	<i>Trochammina triloba</i>	van Voorthuysen, 1969
	Loftusiida	Cyclamminidae	Alveolophragmium	<i>Alveolophragmium orbiculatum</i>	Shchedrina, 1936
		Cyclamminidae	Cyclammina	<i>Cyclammina cancellata</i>	Brady, 1879
		Cyclamminidae	Cyclammina	<i>Cyclammina trullissata</i>	(Brady, 1879)
		Globotextulariidae	Liebusella	<i>Liebusella goesi</i>	Höglund, 1947
		Globotextulariidae	Rhumlerella	<i>Rhumlerella humboldti</i>	(Todd & Brönniman, 1957)
		Globotextulariidae	Verneuilinulla	<i>Verneuilinulla propinqua</i>	(Brady, 1884)
	Robertinida	Epistominidae	Hoeglundina	<i>Hoeglundina elegans</i>	(d'Orbigny, 1826)
		Robertinidae	Robertina	<i>Robertina sp.</i>	d'Orbigny, 1846
		Robertinidae	Robertinoides	<i>Robertinoides bradyi</i>	(Cushman & Parker, 1936)
		Robertinidae	Robertinoides	<i>Robertinoides oceanicum</i>	(Cushman & Parker, 1947)
	Rotaliida	Alabaminidae	Oridorsalis	<i>Oridorsalis pauciapertura</i>	Belford, 1966
		Alabaminidae	Oridorsalis	<i>Oridorsalis tenerus subsp. profundus</i>	Saidova, 1975
		Alabaminidae	Oridorsalis	<i>Oridorsalis umbonatus</i>	(Reuss, 1851)
		Anomalinidae	Anomalinoides	<i>Anomalinoides</i>	Brotzen, 1942
		Astrononionidae	Astrononion	<i>Astrononion echolsi</i>	Kennett, 1967
		Astrononionidae	Astrononion	<i>Astrononion stellatum</i>	(Terquem, 1882)
		Bagginidae	Rotalina	<i>Rotalina pleurostomata</i>	Schlumberger, 1883
		Bolivinitidae	Bolivina	<i>Bolivina (Loxostomum) boltovskoyi</i>	Smith, 1963
		Bolivinitidae	Bolivina	<i>Bolivina (Loxostomum) salvadorensis</i>	Smith, 1963
		Bolivinitidae	Bolivina	<i>Bolivina alata</i>	(Seguenza, 1862)
		Bolivinitidae	Bolivina	<i>Bolivina costata</i>	d'Orbigny, 1839
		Bolivinitidae	Bolivina	<i>Bolivina humilis</i>	Cushman & McCulloch, 1942
		Bolivinitidae	Bolivina	<i>Bolivina interjuncta</i>	Cushman, 1926
		Bolivinitidae	Bolivina	<i>Bolivina pacifica</i>	Cushman & McCulloch, 1942
		Bolivinitidae	Bolivina	<i>Bolivina plicata</i>	d'Orbigny, 1839
Bolivinitidae	Bolivina	<i>Bolivina seminuda</i>	Cushman, 1911		
Bolivinitidae	Bolivina	<i>Bolivina spissa</i>	Cushman, 1926		

		Bolivinitidae	Bolivina	<i>Bolivina subadvena</i>	Cushman, 1926		
		Bolivinitidae	Bolivina	<i>Bolivina subadvena subsp. acuminata</i>	Natland, 1946		
		Bolivinitidae	Bolivina	<i>Bolivina subaenariensis</i>	Cushman, 1922		
		Bolivinitidae	Bolivina	<i>Bolivina variabilis</i>	(Williamson, 1858)		
		Bolivinitidae	Bolivinita	<i>Bolivinita minuta</i>	(Natland, 1938)		
		Bolivinitidae	Fursenkoina	<i>Fursenkoina bradyi</i>	(Cushman, 1922)		
		Bolivinitidae	Fursenkoina	<i>Fursenkoina schreibersiana</i>	(Czjžek, 1848)		
		Bolivinitidae	Fursenkoina	<i>Fursenkoina texturata</i>	(Brady, 1884)		
		Bolivinitidae	Loxostomina	<i>Loxostomina limbata</i>	(Brady, 1881)		
		Bolivinitidae	Lugdunum	<i>Lugdunum hantkenianum</i>	(Brady, 1881)		
		Bolivinitidae	Pseudobrivalina	<i>Pseudobrivalina lobata</i>	(Brady, 1881)		
		Bolivinitidae	Rutherfordoides	<i>Rutherfordoides rotundatus</i>	(Parr, 1950)		
		Bolivinitidae	Suggrunda	<i>Suggrunda eckisi</i>	Natland, 1950		
		Bolivinitidae	Suggrunda	<i>Suggrunda kleinpelli</i>	Bramlette, 1951		
		Bolivinitidae	Suggrunda	<i>Suggrunda porosa</i>	Hoffmeister & Berry, 1937		
		Buliminellidae	Buliminella	<i>Buliminella elegantissima var. limbosa</i>	Cushman & McCulloch, 1948		
		Buliminellidae	Buliminella	<i>Buliminella elegantissima var. tenuis</i>	Cushman & McCulloch, 1948		
		Globothalamea	Rotaliida	Buliminidae	Bulimina	<i>Bulimina aculeata</i>	d'Orbigny, 1826
				Buliminidae	Bulimina	<i>Bulimina denudata</i>	Cushman & Parker, 1938
				Buliminidae	Bulimina	<i>Bulimina elongata</i>	d'Orbigny, 1846
Buliminidae	Bulimina			<i>Bulimina marginata</i>	d'Orbigny, 1826		
Buliminidae	Bulimina			<i>Bulimina pagoda</i>	Cushman, 1927		
Buliminidae	Bulimina			<i>Bulimina striata</i>	d'Orbigny in Guérin-Méneville, 1832		
Buliminidae	Bulimina			<i>Bulimina</i>	d'Orbigny, 1826		
Buliminidae	Bulimina			<i>Bulimina truncana</i>	Gümbel, 1870		
Buliminidae	Praeglobbulimina			<i>Praeglobbulimina primitiva</i>	(Todd, 1952)		
Buliminidae	Praeglobbulimina			<i>Praeglobbulimina spinescens</i>	(Brady, 1884)		
Buliminidae	Protoglobbulimina			<i>Protoglobbulimina pupoides</i>	(d'Orbigny, 1846)		
Cancrisidae	Cancris			<i>Cancris auricula</i>	(Fichtel & Moll, 1798)		
Cancrisidae	Cancris			<i>Cancris carmenensis</i>	Natland, 1950		

		Cancrisidae	Cancris	<i>Cancris oblongus</i>	(d'Orbigny, 1839)
		Cancrisidae	Cancris	<i>Cancris</i>	Montfort, 1808
		Cancrisidae	Gyroidinoides	<i>Gyroidinoides</i>	Brotzen, 1942
		Cancrisidae	Valvulineria	<i>Valvulineria araucana</i>	(d'Orbigny, 1839)
		Cancrisidae	Valvulineria	<i>Valvulineria bradyana</i>	(Fornasini, 1900)
		Cancrisidae	Valvulineria	<i>Valvulineria glabra</i>	Cushman, 1927
		Cancrisidae	Valvulineria	<i>Valvulineria inflata</i>	(d'Orbigny, 1839)
		Cancrisidae	Valvulineria	<i>Valvulineria inaequalis</i>	(d'Orbigny, 1839)
		Cassidulinidae	Cassidulina	<i>Cassidulina auka</i>	Boltovskoy & Theyer, 1970
		Cassidulinidae	Cassidulina	<i>Cassidulina carinata</i>	Silvestri, 1896
		Cassidulinidae	Cassidulina	<i>Cassidulina laevigata</i>	d'Orbigny, 1826
		Cassidulinidae	Cassidulina	<i>Cassidulina</i>	d'Orbigny, 1826
		Cassidulinidae	Cassidulinoides	<i>Cassidulinoides parkeriana</i>	(Brady, 1881)
		Cassidulinidae	Cassidulinoides	<i>Cassidulinoides</i>	Cushman, 1927
		Cassidulinidae	Ehrenbergina	<i>Ehrenbergina compressa</i>	Cushman, 1927
		Cassidulinidae	Ehrenbergina	<i>Ehrenbergina trigona</i>	Goës, 1896
		Cassidulinidae	Globocassidulina	<i>Globocassidulina crassa</i>	(d'Orbigny, 1839)
		Cassidulinidae	Globocassidulina	<i>Globocassidulina minima</i>	(Saidova, 1975)
		Cassidulinidae	Globocassidulina	<i>Globocassidulina</i>	Voloshinova, 1960
		Cassidulinidae	Globocassidulina	<i>Globocassidulina subglobosa</i>	(Brady, 1881)
		Cassidulinidae	Paracassidulina	<i>Paracassidulina minuta</i>	(Cushman, 1933)
		Cassidulinidae	Takayanagia	<i>Takayanagia delicata</i>	(Cushman, 1927)
		Chilostomellidae	Chilostomella	<i>Chilostomella oolina</i>	Schwager, 1878
		Chilostomellidae	Chilostomella	<i>Chilostomella ovoidea</i>	Reuss, 1850
Chilostomellidae	Chilostomella	<i>Chilostomella</i>	Reuss in Czjžek, 1849		
Globothalamea	Rotaliida	Cibicididae	Cibicides	<i>Cibicides pachyderma</i>	(Rzehak, 1886)
		Cibicididae	Cibicidoides	<i>Cibicidoides cicatricosus</i>	(Schwager, 1866)
		Cibicididae	Cibicidoides	<i>Cibicidoides dispars</i>	(d'Orbigny, 1839)
		Cibicididae	Cibicidoides	<i>Cibicidoides mckannai</i>	(Galloway & Wissler, 1927)
		Cibicididae	Cibicidoides	<i>Cibicidoides mundulus</i>	(Brady, Parker & Jones, 1888)
		Cibicididae	Cibicidoides	<i>Cibicidoides robertsonianus</i>	(Brady, 1881)
		Cibicididae	Cibicidoides	<i>Cibicidoides</i>	Thalman, 1939

	Cibicididae	Cibicidoidea	<i>Cibicidoidea variabilis</i>	(d'Orbigny, 1826)
	Cibicididae	Heterolepa	<i>Heterolepa bradyi</i>	(Trauth, 1918)
	Cibicididae	Heterolepa	<i>Heterolepa broeckhiana</i>	(Karrer, 1878)
	Cibicididae	Heterolepa	<i>Heterolepa</i>	Franzenau, 1884
	Cibicididae	Lobatula	<i>Lobatula lobatula</i>	(Walker & Jacob, 1798)
	Cibicididae	Lobatula	<i>Lobatula ungeriana</i>	(d'Orbigny, 1846)
	Cibicididae	Lobatula	<i>Lobatula wuellerstorfi</i>	(Schwager, 1866)
	Discorbidae	Discorbis	<i>Discorbis peruvianus</i>	(d'Orbigny, 1839)
	Discorbidae	Discorbis	<i>Discorbis</i>	Lamarck, 1804
	Discorbidae	Discorbis	<i>Discorbis vilardeboanus</i>	(d'Orbigny, 1839)
	Discorbinellidae	Discorbinella	<i>Discorbinella bertheloti</i>	(d'Orbigny, 1839)
	Discorbinellidae	Hanzawaia	<i>Hanzawaia boueana</i>	(d'Orbigny, 1846)
	Discorbinellidae	Hanzawaia	<i>Hanzawaia mexicana</i>	Lankford, 1973
	Discorbinellidae	Hanzawaia	<i>Hanzawaia prona</i>	Poag, 1966
	Epistomariidae	Nuttallides	<i>Nuttallides umbonifer</i>	(Cushman, 1933)
	Eponidae	Eponides	<i>Eponides</i>	Montfort, 1808
	Gavelinellidae	Gyroidina	<i>Gyroidina altiformis</i>	Stewart & Stewart, 1930
	Gavelinellidae	Gyroidina	<i>Gyroidina gemma</i>	Bandy, 1953
	Gavelinellidae	Gyroidina	<i>Gyroidina lamarckiana</i>	(d'Orbigny, 1839)
	Gavelinellidae	Gyroidina	<i>Gyroidina polia</i>	(Phleger & Parker, 1951)
	Gavelinellidae	Gyroidina	<i>Gyroidina orbicularis</i>	d'Orbigny in Parker, Jones & Brady, 1865
	Gavelinellidae	Gyroidina	<i>Gyroidina rothwelli</i>	Natland, 1950
	Gavelinellidae	Gyroidina	<i>Gyroidina multilocula</i>	Coryell & Mossman, 1942
	Gavelinellidae	Gyroidina	<i>Gyroidina</i>	d'Orbigny, 1826
	Gavelinellidae	Hansenisca	<i>Hansenisca soldanii</i>	(d'Orbigny, 1826)
Rotaliida	Globobuliminidae	Globobulimina	<i>Globobulimina affinis</i>	(d'Orbigny, 1839)
	Globobuliminidae	Globobulimina	<i>Globobulimina auriculata</i>	(Bailey, 1851)
	Globobuliminidae	Globobulimina	<i>Globobulimina hoeglundi</i>	Uchio, 1960
	Globobuliminidae	Globobulimina	<i>Globobulimina notovata</i>	(Chapman, 1941)
	Globobuliminidae	Globobulimina	<i>Globobulimina pacifica</i>	Cushman, 1927
	Globobuliminidae	Globobulimina	<i>Globobulimina</i>	Cushman, 1927
	Globobuliminidae	Globobulimina	<i>Globobulimina spinifera</i>	Uchio, 1960

		Globobuliminidae	Globobulimina	<i>Globobulimina subaffinis</i>	(Cushman, 1921)
		Globobuliminidae	Globobulimina	<i>Globobulimina turgida</i>	(Bailey, 1851)
		Melonidae	Melonis	<i>Melonis affinis</i>	(Reuss, 1851)
		Melonidae	Melonis	<i>Melonis pompilioides</i>	(Fichtel & Moll, 1798)
		Nonionidae	Nonion	<i>Nonion commune</i>	(d'Orbigny, 1846)
		Nonionidae	Nonion	<i>Nonion pacificum</i>	(Cushman, 1924)
		Nonionidae	Nonionella	<i>Nonionella astricta</i>	McCulloch, 1965
		Nonionidae	Nonionella	<i>Nonionella auris</i>	(d'Orbigny, 1839)
		Nonionidae	Nonionella	<i>Nonionella</i>	Cushman, 1926
		Nonionidae	Nonionella	<i>Nonionella stella</i>	Cushman & Moyer, 1930
		Nonionidae	Nonionoides	<i>Nonionoides grateloupii</i>	(d'Orbigny, 1839)
		Nonionidae	Nonionoides	<i>Nonionoides turgidus</i>	(Williamson, 1858)
		Notorotaliidae	Buccella	<i>Buccella peruviana</i>	(d'Orbigny, 1839)
		Planulinidae	Planulina	<i>Planulina ecuadorana</i>	Cushman & Stevenson, 1948
		Pseudoparrellidae	Epistominella	<i>Epistominella afueraensis</i>	McCulloch, 1977
		Pseudoparrellidae	Epistominella	<i>Epistominella exigua</i>	(Brady, 1884)
		Pseudoparrellidae	Epistominella	<i>Epistominella obesa</i>	Bandy & Arnal, 1957
		Pseudoparrellidae	Epistominella	<i>Epistominella pulchella</i>	Husezima & Maruhasi, 1944
		Pseudoparrellidae	Epistominella	<i>Epistominella</i>	Husezima & Maruhasi, 1944
		Pseudoparrellidae	Pseudoparrella	<i>Pseudoparrella pacifica</i>	(Cushman, 1927)
Pseudoparrellidae	Pseudoparrella	<i>Pseudoparrella subperuviana</i>	(Cushman, 1926)		
Globothalamea	Rotaliida	Pulleniidae	Pullenia	<i>Pullenia bulloides</i>	(d'Orbigny, 1846)
		Pulleniidae	Pullenia	<i>Pullenia elegans</i>	Cushman & Todd, 1943
		Pulleniidae	Pullenia	<i>Pullenia quinqueloba</i>	(Reuss, 1851)
		Pulleniidae	Pullenia	<i>Pullenia</i>	Parker & Jones in Carpenter <b>et al.</b> , 1862
		Pulleniidae	Pullenia	<i>Pullenia subcarinata</i>	(d'Orbigny, 1839)
		Pulleniidae	Pulleniella	<i>Pulleniella asymmetrica</i>	Ujiié, 1990
		Rosalinidae	Rosalina	<i>Rosalina globularis</i>	d'Orbigny, 1826
		Sphaeroidinidae	Eusphaeroidina	<i>Eusphaeroidina</i>	Ujiié, 1990

		Sphaeroidinidae	Sphaeroidina	<i>Sphaeroidina bulloides</i>	d'Orbigny in Deshayes, 1832
		Sphaeroidinidae	Sphaeroidina	<i>Sphaeroidina</i>	d'Orbigny, 1826
		Stainforthiidae	Cassidelina	<i>Cassidelina spinescens</i>	(Cushman, 1911)
		Stainforthiidae	Stainforthia	<i>Stainforthia fusiformis</i>	(Williamson, 1858)
		Turrilinidae	Ebuliminella	<i>Ebuliminella basispinata</i>	(Stewart & Stewart, 1930)
		Turrilinidae	Ebuliminella	<i>Ebuliminella curta</i>	(Cushman, 1925)
		Turrilinidae	Ebuliminella	<i>Ebuliminella exilis</i>	(Brady, 1884)
		Uvigerinidae	Siphouvigerina	<i>Siphouvigerina hispida</i>	(Schwager, 1866)
		Uvigerinidae	Siphouvigerina	<i>Siphouvigerina proboscidea</i>	(Schwager, 1866)
		Uvigerinidae	Trifarina	<i>Trifarina angulosa</i>	(Williamson, 1858)
		Uvigerinidae	Uvigerina	<i>Uvigerina auberiana</i>	d'Orbigny, 1839
		Uvigerinidae	Uvigerina	<i>Uvigerina canariensis</i>	d'Orbigny, 1839
		Uvigerinidae	Uvigerina	<i>Uvigerina dirupta</i>	Todd, 1948
		Uvigerinidae	Uvigerina	<i>Uvigerina peregrina</i>	Cushman, 1923
		Uvigerinidae	Uvigerina	<i>Uvigerina semiornata</i>	d'Orbigny, 1846
		Uvigerinidae	Uvigerina	<i>Uvigerina senticosa</i>	Cushman, 1927
		Uvigerinidae	Uvigerina	<i>Uvigerina shiwoensis</i>	Asano, 1958
		Uvigerinidae	Uvigerina	<i>Uvigerina</i>	d'Orbigny, 1826
		Uvigerinidae	Uvigerina	<i>Uvigerina striata</i>	d'Orbigny, 1839
Globothalamea	Textulariida	Eggerellidae	Dorothia	<i>Dorothia bradyana</i>	Cushman, 1936
		Eggerellidae	Dorothia	<i>Dorothia pseudoturris</i>	(Cushman, 1922)
		Eggerellidae	Dorothia	<i>Dorothia scabra</i>	(Brady, 1884)
		Eggerellidae	Eggerella	<i>Eggerella bradyi</i>	(Cushman, 1911)
		Eggerellidae	Eggerella	<i>Eggerella</i>	Cushman, 1933
		Eggerellidae	Karrieriella	<i>Karrieriella baccata</i>	(Schwager, 1866)
		Eggerellidae	Karrieriella	<i>Karrieriella bradyi</i>	(Cushman, 1911)
		Eggerellidae	Martinottiella	<i>Martinottiella communis</i>	(d'Orbigny, 1846)
		Eggerellidae	Martinottiella	<i>Martinottiella nodulosa</i>	Cushman, 1933
		Textulariidae	Sahulia	<i>Sahulia conica</i>	(d'Orbigny, 1839)
		Textulariidae	Textularia	<i>Textularia earlandi</i>	Parker, 1952
		Textulariidae	Textularia	<i>Textularia gramen</i>	d'Orbigny, 1846
		Textulariidae	Textularia	<i>Textularia lythostrota</i>	(Schwager, 1866)

		Textulariidae	Textularia	<i>Textularia porrecta</i>	Brady, 1884
		Valvulinidae	Cribrogoesella	<i>Cribrogoesella pacifica</i>	Cushman & McCulloch, 1939
Nodosariata	Lagenidae	Lagenidae	Hyalinonetrion	Hyalinonetrion clavatum	(d'Orbigny, 1846)
		Lagenidae	Hyalinonetrion	Hyalinonetrion gracillimum	(Seguenza, 1862)
		Lagenidae	Lagena	Lagena aspera	Reuss, 1862
		Lagenidae	Lagena	Lagena hispidula	Cushman, 1913
		Lagenidae	Lagena	Lagena parvulipora	Bandy, 1949
		Lagenidae	Lagena	Lagena striata	(d'Orbigny, 1839)
		Lagenidae	Lagena	Lagena substriata	Williamson, 1848
		Lagenidae	Lagena	Lagena sulcata	(Walker & Jacob, 1798)
		Lagenidae	Procerolagena	Procerolagena distoma	(Parker & Jones, 1864)
		Lagenidae	Procerolagena	Procerolagena gracilis	(Williamson, 1848)
		Lagenidae	Reussoolina	Reussoolina apiculata	(Reuss, 1851)
		Lagenidae	Reussoolina	Reussoolina laevis	(Montagu, 1803)
		Nodosariidae	Dentalina	Dentalina	Risso, 1826
	Nodosariidae	Nodosariidae	Laevidentalina	Laevidentalina baggi	(Galloway & Wissler, 1927)
		Nodosariidae	Laevidentalina	Laevidentalina communis	(d'Orbigny, 1826)
		Nodosariidae	Laevidentalina	Laevidentalina elegans	(d'Orbigny, 1846)
		Nodosariidae	Laevidentalina	Laevidentalina filiformis	(d'Orbigny, 1826)
Nodosariidae		Laevidentalina	Laevidentalina subsoluta	(Cushman, 1923)	
Nodosariata	Polymorphinida	Ellipsolagenidae	Fissurina	<i>Fissurina annectens</i>	(Burrows & Holland, 1895)
		Ellipsolagenidae	Fissurina	<i>Fissurina circularis</i>	Todd, 1954
		Ellipsolagenidae	Fissurina	<i>Fissurina cucullata</i>	Silvestri, 1902
		Ellipsolagenidae	Fissurina	<i>Fissurina cucurbitasema</i>	Loeblich & Tappan, 1953
		Ellipsolagenidae	Fissurina	<i>Fissurina gravata</i>	McCulloch, 1977
		Ellipsolagenidae	Fissurina	<i>Fissurina granifera</i>	(Buchner, 1940)
		Ellipsolagenidae	Fissurina	<i>Fissurina laevigata</i>	Reuss, 1850
		Ellipsolagenidae	Fissurina	<i>Fissurina marginata</i>	(Montagu, 1803)
		Ellipsolagenidae	Fissurina	<i>Fissurina orbignyana</i>	Seguenza, 1862
		Ellipsolagenidae	Fissurina	<i>Fissurina (Fissurina) ovata</i>	Seguenza, 1862
		Ellipsolagenidae	Fissurina	<i>Fissurina piriformis</i>	(Buchner, 1940)
		Ellipsolagenidae	Fissurina	<i>Fissurina</i>	Reuss, 1850
		Ellipsolagenidae	Fissurina	<i>Fissurina marginata</i>	(Montagu, 1803)

		Ellipsolagenidae	Fissurina	<i>Fissurina susanae</i>	Figuroa, Marchant & Ramirez in Figuroa <b>et al.</b> , 2006
		Ellipsolagenidae	Galwayella	<i>Galwayella trigonoornata</i>	(Brady, 1881)
		Ellipsolagenidae	Oolina	<i>Oolina globosa</i>	(Montagu, 1803)
		Ellipsolagenidae	Palliolatella	<i>Palliolatella bradii</i>	(Silvestri, 1902)
		Ellipsolagenidae	Parafissurina	<i>Parafissurina lateralis</i>	(Cushman, 1913)
		Ellipsolagenidae	Parafissurina	<i>Parafissurina ovata</i>	(Wiesner, 1931)
		Ellipsolagenidae	Parafissurina	<i>Parafissurina</i>	Parr, 1947
		Ellipsolagenidae	Parafissurina	<i>Parafissurina ventricosa</i>	(Silvestri, 1904)
		Ellipsolagenidae	Vasicostella	<i>Vasicostella inflatiperforata</i>	(McCulloch, 1977)
		Glandulinidae	Glandulina	<i>Glandulina laevigata</i>	(d'Orbigny, 1826)
		Glandulinidae	Glandulina	<i>Glandulina ovula</i>	d'Orbigny, 1846
		Glandulinidae	Glandulina	<i>Glandulina symmetrica</i>	(McCulloch, 1977)
		Pleurostomellidae	Obesopleurostomella	<i>Obesopleurostomella brevis</i>	(Schwager, 1866)
		Polymorphinidae	Pyulina	<i>Pyulina</i>	d'Orbigny, 1839
		Polymorphinidae	Pyulina	<i>Pyulina angusta</i>	(Egger, 1857)
		Vaginulinidae	Astacolus	<i>Astacolus insolitus</i>	(Schwager, 1866)
Nodosariata	Vaginulinida	Vaginulinidae	Lenticulina	<i>Lenticulina convergens</i>	(Bornemann, 1855)
		Vaginulinidae	Lenticulina	<i>Lenticulina gibba</i>	(d'Orbigny, 1839)
		Vaginulinidae	Lenticulina	<i>Lenticulina inornata</i>	(d'Orbigny, 1846)
		Vaginulinidae	Lenticulina	<i>Lenticulina pliocaena</i>	(Silvestri, 1898)
		Vaginulinidae	Lenticulina	<i>Lenticulina rotulata</i>	(Lamarck, 1804)
		Vaginulinidae	Lenticulina	<i>Lenticulina</i>	Lamarck, 1804
		Vaginulinidae	Lenticulina	<i>Lenticulina thalmani</i>	(Hessland, 1943)
		Vaginulinidae	Saracenaria	<i>Saracenaria stolidota</i>	Loeblich & Tappan, 1994
		Vaginulinidae	Vaginulina	<i>Vaginulina americana</i>	Cushman, 1923
		Ammolagenidae	Ammolagena	<i>Ammolagena clavata</i>	(Jones & Parker, 1860)
		Hormosinellidae	Hormosinella	<i>Hormosinella distans</i>	(Brady, 1881)
		Hormosinellidae	Hormosinelloides	<i>Hormosinelloides guttifer</i>	(Brady, 1881)
		Hormosinidae	Hormosina	<i>Hormosina globulifera</i>	Brady, 1879
		Hormosinidae	Nodosinum	<i>Nodosinum gaussicum</i>	(Rhumbler, 1913)

		Hormosinidae	Pseudonodosinella	<i>Pseudonodosinella nodulosa</i>	(Brady, 1879)
		Reophacidae	Nodulina	<i>Nodulina dentaliniformis</i>	(Brady, 1881)
		Reophacidae	Reophax	<i>Reophax apiculatus</i>	Zheng, 1988
		Reophacidae	Reophax	<i>Reophax bradyi</i>	Brönnimann & Whittaker, 1980
		Reophacidae	Reophax	<i>Reophax duplex</i>	Grzybowski, 1896
		Reophacidae	Reophax	<i>Reophax fusiformis</i>	(Williamson, 1858)
		Reophacidae	Reophax	<i>Reophax globosus</i>	Sliter, 1968
		Reophacidae	Reophax	<i>Reophax insectus</i>	Goës, 1896
		Reophacidae	Reophax	<i>Reophax pilulifer</i>	Brady, 1884
		Reophacidae	Reophax	<i>Reophax pisiformis</i>	Zheng, 1988
		Reophacidae	Reophax	<i>Reophax regularis</i>	Höglund, 1947
		Reophacidae	Reophax	<i>Reophax scorpiurus</i>	Montfort, 1808
		Reophacidae	Reophax	<i>Reophax subdentaliniformis</i>	Parr, 1950
		Reophacidae	Reophax	<i>Reophax subfusiformis</i>	Earland Em. Höglund, 1947
		Cornuspiridae	Cornuspira	<i>Cornuspira involvens</i>	(Reuss, 1850)
Tubothalamea	Miliolida	Hauerinidae	Miliolinella	<i>Miliolinella subrotunda</i>	(Montagu, 1803)
		Hauerinidae	Nummoloculina	<i>Nummoloculina</i>	Steinmann, 1881
		Hauerinidae	Pyrgo	<i>Pyrgo depressa</i>	(d'Orbigny, 1826)
		Hauerinidae	Pyrgo	<i>Pyrgo elongata</i>	(d'Orbigny, 1826)
		Hauerinidae	Pyrgo	<i>Pyrgo murrhina</i>	(Schwager, 1866)
		Hauerinidae	Pyrgo	<i>Pyrgo sarsi</i>	(Schlumberge, 1891)
		Hauerinidae	Pyrgo	<i>Pyrgo</i>	Defrance, 1824
		Hauerinidae	Pyrgoella	<i>Pyrgoella irregularis</i>	(d'Orbigny, 1839)
		Hauerinidae	Pyrgoella	<i>Pyrgoella sphaera</i>	(d'Orbigny, 1839)
		Hauerinidae	Quinqueloculina	<i>Quinqueloculina seminulum</i>	(Linnaeus, 1758)
		Hauerinidae	Quinqueloculina	<i>Quinqueloculina</i>	d'Orbigny, 1826
		Hauerinidae	Quinqueloculina	<i>Quinqueloculina weaveri</i>	Rau, 1948
		Hauerinidae	Sigmoilina	<i>Sigmoilina</i>	Schlumberger, 1887
		Hauerinidae	Spirosigmoilina	<i>Spirosigmoilina tenuis</i>	(Czjžek, 1848)
		Hauerinidae	Triloculina	<i>Triloculina tricarinata</i>	d'Orbigny in Deshayes, 1832
		Hauerinidae	Triloculina	<i>Triloculina trigonula</i>	(Lamarck, 1804)
		Miliolidae	Planispirinoides	<i>Planispirinoides bucculentus</i>	(Brady, 1884)

		Nubeculariidae	Nubeculina	<i>Nubeculina</i>	Cushman, 1924
		Spiroloculinidae	Spiroloculina	<i>Spiroloculina robusta</i>	Brady, 1884
		Spiroloculinidae	Spiroloculina	<i>Spiroloculina</i>	d'Orbigny, 1826
		Ammodiscidae	Arenoturrspirillina	<i>Arenoturrspirillina catinus</i>	(Höglund, 1947)
	Spirillinida	Ammodiscidae	Ammodiscus	<i>Ammodiscus incertus</i>	(d'Orbigny, 1839)
		Ammodiscidae	Ammodiscus	<i>Ammodiscus planorbis</i>	Höglund, 1947
		Ammodiscidae	Ammodiscus	<i>Ammodiscus tenuis</i>	(Brady, 1881)
		Ammodiscidae	Glomospira	<i>Glomospira gordialis</i>	(Jones & Parker, 1860)
		Ammodiscidae	Tolypammina	<i>Tolypammina vagans</i>	(Brady, 1879)
		Spirillinidae	Spirillina	<i>Spirillina</i>	Ehrenberg, 1843

### Appendix 3. Taxonomic and Functional Indices of benthic foraminifera from the Southeastern Pacific

ID	Abundance (ind/50cm <sup>3</sup> )	Taxa	Simpson	Shannon	Pielou	Fisher_α	FRic	FDiv	FDis	FEve	RaoQ
Site_1_207	2222.22	26	0.85	2.29	0.70	5.54	16.94	0.68	3.78	0.43	15.33
Site_1_526	850.21	35	0.92	2.86	0.81	10.45	13.65	0.66	4.29	0.53	18.93
Site_1_2092	197.55	40	0.95	3.31	0.90	20.47	97.72	0.79	4.79	0.64	24.39
Site_3_350	380.39	11	0.67	1.57	0.66	2.53	1.78	0.70	3.02	0.55	10.28
Site_3_700	554.35	13	0.44	1.01	0.39	2.90	4.47	0.86	2.97	0.64	12.66
Site_3_995	195.04	40	0.94	3.22	0.87	16.21	95.80	0.72	4.35	0.62	20.22
Site_7_359	5376.96	10	0.18	0.48	0.21	1.54	2.77	0.96	0.59	0.53	1.94
Site_7_627	437.36	19	0.71	1.72	0.58	5.22	14.27	0.93	2.95	0.50	12.48
Site_9_115	72709.28	13	0.39	0.95	0.37	2.39	1.13	0.91	1.87	0.53	7.07
Site_9_437	19184.61	17	0.70	1.68	0.59	2.85	4.74	0.84	2.01	0.54	7.03
Site_9_1105	193.90	19	0.75	2.10	0.71	8.93	43.76	0.88	4.41	0.60	21.51
Site_10_521	3523.61	24	0.89	2.47	0.78	5.49	17.36	0.75	3.70	0.46	15.53
Site_10_1004	1080.51	36	0.88	2.81	0.79	13.66	79.77	0.79	4.81	0.60	24.64
Site_10_1923	615.41	33	0.91	2.92	0.83	13.28	74.70	0.75	4.98	0.60	25.58
Site_11_79	34709.82	12	0.57	1.12	0.45	1.83	9.05	0.43	3.18	0.22	10.33
Site_11_145	486.19	6	0.39	0.77	0.43	1.20	0.16	0.87	1.91	0.44	5.69
Site_11_211	275.82	7	0.25	0.61	0.31	1.49	0.25	0.96	1.15	0.29	3.72
Site_11_248	105558.44	11	0.18	0.48	0.20	1.66	1.72	0.93	0.79	0.26	2.33
Site_11_291	30088.31	12	0.57	1.30	0.52	2.39	4.78	0.89	2.01	0.58	6.21
Site_11_317	52324.32	10	0.51	1.06	0.46	1.78	1.50	0.91	1.40	0.50	3.84
Site_11_319	25854.59	17	0.63	1.56	0.55	3.13	5.54	0.93	2.05	0.32	7.26
Site_11_375	29295.62	17	0.71	1.72	0.61	3.17	4.75	0.83	2.45	0.30	7.53
Site_11_465	15370.69	35	0.75	2.05	0.58	7.00	26.13	0.90	2.80	0.49	11.01

Site 11 511	3943.61	22	0.80	2.09	0.68	5.65	15.62	0.76	3.57	0.48	13.53
Site 11 579	2420.38	19	0.74	1.91	0.65	5.26	7.32	0.66	4.46	0.56	20.38
Site 11 697	3146.69	34	0.88	2.45	0.70	7.02	66.48	0.70	4.78	0.43	23.18
Site 12 152	38.67	2	0.05	0.12	0.17	0.45	0	0	0.21	0	0.43
Site 12 214	14760.30	14	0.27	0.76	0.29	2.51	5.60	0.95	1.25	0.61	4.40
Site 12 302	41601.51	15	0.65	1.41	0.52	2.30	5.03	0.87	1.70	0.29	5.23
Site 12 823	1527.21	36	0.91	2.79	0.78	9.06	89.27	0.69	4.72	0.50	22.56
Site 15 522	470.93	10	0.70	1.55	0.67	2.36	4.95	0.81	3.84	0.33	16.38
Site 16 2575	113.85	23	0.80	2.21	0.71	8.45	25.78	0.74	2.58	0.50	8.78
Site 16 3228	39.41	20	0.89	2.59	0.87	10.23	33.24	0.81	4.78	0.63	24.11
Site 16 4124	6.59	6	0.76	1.61	0.90	6.33	7.57	0.84	5.23	0.69	29.07
Site 17 298	852.22	15	0.83	2.11	0.78	3.20	2.83	0.77	3.00	0.57	11.03
Site 17 492	1076.92	18	0.86	2.35	0.81	7.03	14.73	0.71	4.66	0.64	21.87
Site 17 918	162.16	9	0.86	2.09	0.95	16.36	20.69	0.71	3.96	0.76	18.11
Site 23 523	120.38	30	0.92	2.88	0.85	10.05	28.92	0.80	4.10	0.58	17.37
Site 23 3396	24.35	15	0.88	2.40	0.89	8.35	10.32	0.58	3.66	0.63	14.70
Site 25 1442	31.21	25	0.93	2.98	0.92	20.44	14.95	0.83	4.68	0.65	23.09
Site 26 673	2.55	13	0.90	2.41	0.94	10.35	9.71	0.83	5.20	0.80	28.36
Site 26 1996	16.56	4	0.75	1.39	1.00	0	0.05	0.76	3.21	0.96	10.52
Site 27 557	40.76	17	0.78	2.03	0.72	7.57	45.34	0.74	3.72	0.65	15.64
Site 28 473	87.90	26	0.89	2.60	0.80	9.47	28.30	0.84	4.46	0.65	21.02
Site 28 1462	59.24	24	0.89	2.59	0.82	10.48	82.83	0.84	5.00	0.57	26.06
Site 28 1650	42.04	18	0.90	2.57	0.89	8.15	13.69	0.88	4.68	0.63	22.54
Site 28 2080	15.92	11	0.84	2.10	0.87	7.50	13.07	0.92	4.85	0.80	23.85
Site 29 635	56.05	25	0.92	2.79	0.87	11.65	67.06	0.71	3.93	0.60	17.04
Site 29 1432	246.50	54	0.93	3.16	0.79	17.06	101.46	0.80	4.60	0.50	22.31
Site 29 3252	58.60	26	0.91	2.76	0.85	12.07	90.27	0.77	4.95	0.60	25.84
Site 30 1199	84.08	29	0.73	2.17	0.65	11.48	83.54	0.81	3.95	0.69	18.53
Site 30 2733	7.01	4	0.55	1.03	0.75	2.26	0.61	0.88	3.63	0.71	14.82
Site 30 3267	43.95	14	0.76	1.81	0.69	5.30	7.53	0.83	4.19	0.63	18.63
Site 31 183	5580.25	21	0.60	1.14	0.38	2.58	7.24	0.79	3.04	0.24	10.17
Site 31 301	86.62	15	0.84	2.16	0.80	4.31	23.01	0.71	4.02	0.51	17.62
Site 31 326	136.31	13	0.69	1.57	0.61	3.05	5.07	0.82	3.75	0.52	15.21
Site 31 398	1010.19	51	0.91	2.84	0.72	10.07	38.60	0.76	4.20	0.38	18.61
Site 31 481	187.26	45	0.96	3.37	0.89	14.82	80.69	0.76	4.28	0.54	19.78
Site 31 1371	92.99	34	0.92	2.90	0.82	13.93	85.25	0.84	4.49	0.56	21.00
Site 31 2289	37.58	26	0.93	2.98	0.92	17.77	81.48	0.69	5.36	0.63	29.68
Site 31 3086	56.69	24	0.90	2.68	0.84	10.79	21.61	0.80	4.90	0.58	25.35
Site 32 1591	262.42	48	0.91	2.96	0.76	14.08	94.49	0.91	4.28	0.50	19.74

Site 32 3920	77.18	21	0.87	2.37	0.78	6.13	16.24	0.62	3.42	0.60	12.76
Site 33 420	60.51	28	0.67	1.72	0.52	6.66	26.67	0.70	4.11	0.58	18.06
Site 33 863	54.95	37	0.80	2.48	0.69	10.97	28.32	0.83	4.03	0.53	17.72
Site 33 1385	280.25	22	0.88	2.52	0.81	8.98	12.32	0.85	4.38	0.63	19.86
Site 33 3852	196.82	18	0.86	2.33	0.81	7.17	23.66	0.86	3.53	0.70	15.12
Site 34 1247	40.13	14	0.86	2.20	0.83	5.58	3.33	0.60	3.87	0.62	16.81
Site 35 506	150.96	20	0.72	1.88	0.63	5.21	9.91	0.78	4.30	0.53	19.40
Site 35 1563	109.55	20	0.64	1.64	0.55	5.86	12.09	0.85	3.96	0.61	17.46
Site 35 2103	24.20	15	0.89	2.43	0.90	9.15	25.25	0.69	4.65	0.70	24.12
Site 36 366	38.22	8	0.61	1.28	0.62	2.48	0.96	0.73	3.74	0.38	15.42
Site 36 538	175.80	28	0.77	2.01	0.60	7.79	21.26	0.81	4.49	0.62	20.65
Site 36 1294	8.28	6	0.77	1.61	0.90	4.32	11.08	0.83	4.97	0.92	24.90
Site 36 2064	8.28	8	0.83	1.93	0.93	8.85	1.57	0.67	3.74	0.61	14.40
Site 37 124	816.56	14	0.73	1.63	0.62	2.20	6.74	0.82	3.54	0.33	13.19
Site 37 1386	15.92	12	0.80	2.06	0.83	9.06	5.77	0.89	4.92	0.80	25.04
Site 38 281	9.55	4	0.73	1.34	0.97	1.78	0.86	0.84	3.77	0.91	14.34
Site 38 700	45.86	23	0.92	2.85	0.91	11.68	14.99	0.78	3.90	0.59	17.09
Site 38 2287	11.46	9	0.83	1.98	0.90	7.16	4.64	0.81	5.81	0.84	34.90
Site 39 112	5.10	4	0.72	1.32	0.95	3.18	1.13	0.75	4.05	0.78	17.38
Site 39 174	268.79	17	0.74	1.72	0.61	3.55	17.00	0.62	4.51	0.54	21.01
Site 39 498	5.10	3	0.59	0.97	0.89	1.74	0.00	0.00	3.04	0.00	9.83
Site 39 772	136.31	39	0.90	2.84	0.77	13.97	19.69	0.68	4.35	0.58	19.53
Site 39 890	40.76	20	0.88	2.46	0.82	9.99	16.06	0.74	3.80	0.52	16.79
Site 39 1190	134.39	41	0.90	2.84	0.76	15.18	20.44	0.85	4.81	0.55	23.58
Site 39 1469	28.03	10	0.73	1.70	0.74	4.04	6.60	0.72	4.06	0.60	17.98
Site 39 1798	3.82	4	0.67	1.24	0.90	5.24	2.13	0.83	4.42	0.89	19.88
Site 40 165	217.20	18	0.70	1.68	0.58	4.05	55.56	0.69	3.91	0.44	16.87
Site 40 816	19.75	17	0.92	2.70	0.95	15.43	10.23	0.85	3.98	0.80	16.88
Site 40 3850	27.09	16	0.84	2.30	0.83	12.73	18.04	0.75	3.67	0.66	16.73
Site 41 308	424.20	44	0.72	2.15	0.57	10.58	91.47	0.74	3.90	0.51	17.96
Site 41 521	38.22	13	0.58	1.50	0.58	5.11	8.69	0.66	3.83	0.66	18.63
Site 42 160	130.57	21	0.85	2.33	0.77	5.86	24.43	0.84	4.11	0.68	18.02
Site 42 206	63.69	16	0.81	2.12	0.77	5.38	23.24	0.74	4.09	0.66	18.57
Site 42 299	421.66	49	0.90	2.78	0.71	12.22	37.26	0.86	4.43	0.54	20.56
Site 42 301	128.03	30	0.88	2.60	0.77	9.77	16.97	0.83	4.69	0.58	22.37
Site 42 909	14.65	12	0.90	2.38	0.96	10.12	12.21	0.71	4.29	0.60	19.87
Site 42 1222	31.85	17	0.88	2.47	0.87	9.07	9.93	0.61	4.46	0.61	20.71
Site 42 1967	32.48	12	0.72	1.69	0.68	4.95	18.70	0.83	4.40	0.68	19.83
Site 42 2760	8.92	6	0.71	1.48	0.82	3.98	0.08	0.69	3.88	0.64	15.29

Site 43 1212	68.15	27	0.85	2.49	0.76	11.62	16.27	0.69	4.17	0.55	18.65
Site 43 3523	89.59	26	0.92	2.80	0.86	11.06	29.23	0.79	4.52	0.55	22.06
Site 44 476	32.48	15	0.87	2.31	0.85	7.16	15.90	0.84	4.13	0.70	17.69
Site 44 868	4.46	6	0.82	1.75	0.98	19.95	1.76	0.64	3.80	0.86	16.32
Site 44 1171	33.76	14	0.87	2.29	0.87	6.21	6.78	0.76	4.05	0.66	18.28
Site 44 1939	16.56	9	0.79	1.87	0.85	4.88	5.67	0.85	4.97	0.70	24.93
Site 44 3285	19.75	8	0.73	1.58	0.76	3.49	6.20	0.82	5.25	0.65	27.62
Site 46 3014	77.83	20	0.88	2.40	0.80	6.77	24.78	0.81	3.11	0.59	11.83
Site 50 3964	64.99	21	0.92	2.72	0.89	6.99	15.99	0.69	4.48	0.46	21.29

Appendix 4. Species of benthic foraminifera with abundance of >100 ind/50cm<sup>3</sup> in all marine surface sediment stations of the Southeastern Pacific.

Species	Abundance (ind/cm <sup>3</sup> )	Species	Abundance (ind/cm <sup>3</sup> )	Species	Abundance (ind/cm <sup>3</sup> )
<i>Bolivina seminuda</i>	297513.1148	<i>Eggerella bradyi</i>	890.4306983	<i>Eratidus foliaceus</i>	196.4395842
<i>Bolivina costata</i>	49807.72173	<i>Globocassidulina crassa</i>	855.4442349	<i>Eggerelloides scabrum</i>	187.7608415
<i>Bolivina plicata</i>	24447.11439	<i>Takayanagia delicata</i>	852.1055755	<i>Uvigerina auferiana</i>	181.0884174
<i>Bolivina interjuncta</i>	13846.08513	<i>Hoeglundina elegans</i>	826.6384681	<i>Cyclammina cancellata</i>	179.9099485
<i>Nonionella stella</i>	12912.80563	<i>Epistominella exigua</i>	757.8257267	<i>Melonis affinis</i>	176.7979682
<i>Valvulineria glabra</i>	12276.00427	<i>Pullenia elegans</i>	719.8391069	<i>Ammobaculites agglutinans</i>	176.5745616
<i>Uvigerina peregrina</i>	7459.73682	<i>Uvigerina striata</i>	712.3500255	<i>Glaphyrammina americana</i>	168.5770594
<i>Cancris carmenensis</i>	7152.499951	<i>Reophax scorpiurus</i>	690.4081316	<i>Pullenia subcarinata</i>	166.0457993
<i>Stainforthia fusiformis</i>	6447.063767	<i>Pseudoparrella subperuviana</i>	665.099804	<i>Recurvoides contortus</i>	156.6548545
<i>Epistominella obesa</i>	6123.406176	<i>Lobatula wuellerstorfi</i>	640.7627639	<i>Reophax bradyi</i>	154.7770701
<i>Buliminella elegantissima tenuis</i>	5479.454832	<i>Hanzawaia prona</i>	527.8350515	<i>Ammodiscus incertus</i>	150.791102
<i>Cassidulina auka</i>	4711.16336	<i>Cassidulina carinata</i>	520.413514	<i>Adercotryma glomeratum</i>	150.1245927
<i>Valvulineria inflata</i>	2617.197452	<i>Ebuliminella curta</i>	518.1479153	<i>Nodulina dentaliniformis</i>	147.595177
<i>Buliminella elegantissima limbosa</i>	2052.548198	<i>Suggrunda porosa</i>	438.4322437	<i>Oridorsalis umbonatus</i>	141.6770055
<i>Gyroidina soldanii var multilocula</i>	1924.478004	<i>Nonion commune</i>	405.0438932	<i>Hanzawaia mexicana</i>	141.6409699
<i>Bolivina spissa</i>	1911.722829	<i>Pseudoparrella pacifica</i>	387.9809104	<i>Chilostomella oolina</i>	137.2697182
<i>Protoglobobulimina pupoides</i>	1644.692942	<i>Ebuliminella basispinata</i>	324.3799292	<i>Cassidulina laevigata</i>	128.0254777

<i>Bolivina alata</i>	1582.880978	<i>Pseudobrizalina lobata</i>	285.1161164	<i>Lagenammina difflugiformis</i>	122.4833748
<i>Bolivinina minuta</i>	1514.238575	<i>Reophax apiculatus</i>	277.7199072	<i>Globobulimina affinis</i>	114.6496815
<i>Trifarina angulosa</i>	1267.635619	<i>Nonionoides grateloupii</i>	273.468043	<i>Glomospira gordialis</i>	111.8838404
<i>Fursenkoina texturata</i>	1164.04971	<i>Cancris oblongus</i>	206.4912026	<i>Globocassidulina subglobosa</i>	107.3386082
<i>Hansenisca soldanii</i>	974.315306	<i>Bolivina subadvena</i> subsp. <i>Acuminata</i>	204.3981278	<i>Ammoglobigerina globigeriniformis</i>	104.2367674

Table S5. Matrix of functional traits of benthic foraminifera in the Southeastern Pacific. SI: Shallow infaunal. II: Intermedia infaunal

Specie	Type of wall	Morphology	Microhabitat	Oxygen condition	Trophic behavior
<i>Adercotryma glomeratum</i>	Agglutinated	Globose, subglobose, ovoid or streptospiral	Epifaunal	Oxic	Herbivore / Phytodetrivore
<i>Alveolophragmium orbiculatum</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Ammobaculites agglutinans</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Ammodiscus catinus</i>	Agglutinated	Planispiral	Epifaunal	Hypoxic - Anoxic	Detritivore
<i>Ammodiscus incertus</i>	Agglutinated	Planispiral	Epifaunal	Hypoxic - Anoxic	Detritivore
<i>Ammodiscus planorbis</i>	Agglutinated	Planispiral	Epifaunal	Hypoxic - Anoxic	Detritivore
<i>Ammodiscus tenuis</i>	Agglutinated	Planispiral	Epifaunal	Hypoxic - Anoxic	Detritivore
<i>Ammoglobigerina globigeriniformis</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ammoglobigerina globulosa</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ammolagena clavata</i>	Agglutinated	Irregular	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ammomarginulina catenulata</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Ammomassilina alveoliniformis</i>	Porcellaneous	Miliolide	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ammosphaeroidina grandis</i>	Agglutinated	Globose, subglobose, ovoid or streptospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Anomalinoidea sp</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Astacolus insolitus</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore

<i>Astrammina rara</i>	Agglutinated	Irregular	Intermedia infaunal	Oxic	Carnivore Bacterivore
<i>Astrononion echolsi</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Phytodetrivore
<i>Astrononion stellatum</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Astrorhiza arenaria</i>	Agglutinated	Irregular	Intermedia infaunal	Oxic	Bacterivore
<i>Astrorhiza granulosa</i>	Agglutinated	Irregular	Intermedia infaunal	Oxic	Bacterivore
<i>Astrorhizoides cornutus</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Bathysiphon capillare</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Phytodetrivore
<i>Bathysiphon fusca</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Phytodetrivore
<i>Bathysiphon sp.</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Phytodetrivore
<i>Bolivina (Loxostomum) boltovskoyi</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina (Loxostomum) salvadorensis</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina alata</i>	Hyaline	Biserial	Shallow Infaunal	Hypoxic - Anoxic	Phytodetrivore
<i>Bolivina alata</i>	Hyaline	Biserial	Shallow Infaunal	Hypoxic - Anoxic	Phytodetrivore
<i>Bolivina costata</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina interjuncta</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina pacifica</i>	Hyaline	Biserial	Epifaunal	Hypoxic	Phytodetrivore
<i>Bolivina plicata</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina seminuda</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina spissa</i>	Hyaline	Biserial	Epifaunal	Oxic - Anoxic	Phytodetrivore
<i>Bolivina subadvena</i>	Hyaline	Biserial	Epifaunal	Hypoxic	Phytodetrivore
<i>Bolivina subadvena subsp. Acuminata</i>	Hyaline	Biserial	Epifaunal	Hypoxic	Phytodetrivore
<i>Bolivina subaenariensis</i>	Hyaline	Biserial	Shallow Infaunal	Hypoxic	Phytodetrivore
<i>Bolivina variabilis</i>	Hyaline	Biserial	Epifaunal	Oxic - Hypoxic	Phytodetrivore
<i>Bolivinita minuta</i>	Hyaline	Biserial	Epifaunal	Hypoxic - Anoxic	Phytodetrivore
<i>Buccella peruviana</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Bulimina aculeata</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Bulimina aculeata</i>	Hyaline	Triserial	SI/II	Oxic - Hypoxic	Detritivore

<i>Bulimina denudata</i>	Hyaline	Triserial	SI/II	Oxic - Hypoxic	Detritivore
<i>Bulimina elongata</i>	Hyaline	Triserial	SI/II	Oxic - Hypoxic	Detritivore
<i>Bulimina pagoda</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Bulimina sp.</i>	Hyaline	Triserial	SI/II	Oxic - Hypoxic	Detritivore
<i>Bulimina truncana</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Bulimina truncata</i>	Hyaline	Triserial	SI/II	Oxic - Hypoxic	Detritivore
<i>Buliminella elegantissima var limbosa</i>	Hyaline	Trochospiral	Epifaunal	Hypoxic - anoxic	Detritivore
<i>Buliminella elegantissima var tenuis</i>	Hyaline	Trochospiral	Epifaunal	Hypoxic - anoxic	Detritivore
<i>Buzasina ringens</i>	Agglutinated	Planispiral	Intermedia infaunal	Oxic	Detritivore
<i>Buzasina sp.</i>	Agglutinated	Planispiral	Intermedia infaunal	Oxic	Detritivore
<i>Cancris auricula</i>	Hyaline	Trochospiral	Intermedia infaunal	Hypoxic	Detritivore
<i>Cancris carmenensis</i>	Hyaline	Trochospiral	Intermedia infaunal	Hypoxic - Anoxic	Detritivore
<i>Cancris oblongus</i>	Hyaline	Trochospiral	Intermedia infaunal	Hypoxic	Detritivore
<i>Cancris sp</i>	Hyaline	Trochospiral	Intermedia infaunal	Hypoxic	Detritivore
<i>Cassidelina spinescens</i>	Hyaline	Biserial	Shallow Infaunal	Anoxic	Detritivore
<i>Cassidulina auka</i>	Hyaline	Planispiral	Intermedia infaunal	Hypoxic - Anoxic	Phytodetritivore
<i>Cassidulina carinata</i>	Hyaline	Planispiral	Intermedia infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Cassidulina laevigata</i>	Hyaline	Planispiral	Intermedia infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Cassidulina sp</i>	Hyaline	Planispiral	Intermedia infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Cassidulinoides parkeriana</i>	Hyaline	Planispiral	Intermedia infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Cassidulinoides sp</i>	Hyaline	Planispiral	Intermedia infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Chilostomella oolina</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Chilostomella ovoidea</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Chilostomella sp</i>	Hyaline	Globose, subglobose,	Intermedia infaunal	Oxic - Hypoxic	Detritivore

		ovoid or streptospiral			
<i>Cibicides pachyderma</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Cibicidoides cicatricosus</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Phytodetrivore
<i>Cibicidoides dispars</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Cibicidoides mckannai</i>	Hyaline	Trochospiral	Epifaunal	Oxic	Phytodetrivore
<i>Cibicidoides mundulus</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Phytodetrivore
<i>Cibicidoides robertonianus</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Cibicidoides sp</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Cibicidoides variabilis</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Cornuspira involvens</i>	Porcellaneous	Planispiral	Epifaunal - Shallow Infaunal	Oxic - Anoxic	Phytodetrivore
<i>Cribragoesella pacifica</i>	Agglutinated	Trochospiral	Intermediate infaunal	Oxic - Hypoxic	Detritivore
<i>Cribrostomoides jeffreysii</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Cribrostomoides nitidus</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Cribrostomoides sphaerilocula</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Cribrostomoides subglobosus</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Crithionina pisum</i>	Agglutinated	Spheric unilocular and elongated spheric	Shallow Infaunal-II	Oxic - Hypoxic	Detritivore/ Bacterivore
<i>Cyclammina cancellata</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Cyclammina trullissata</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Dentalina sp.</i>	Hyaline	Uniserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Discorbinella bertheloti</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Herbivore
<i>Discorbis peruvianus</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Herbivore
<i>Discorbis sp</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Herbivore
<i>Discorbis vilardeboanus</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic	Herbivore
<i>Dorothia bradyana</i>	Agglutinated	Biserial	Intermediate infaunal	Oxic	Detritivore
<i>Dorothia pseudoturris</i>	Agglutinated	Biserial	Intermediate infaunal	Oxic	Detritivore
<i>Dorothia scabra</i>	Agglutinated	Biserial	Intermediate infaunal	Oxic	Detritivore
<i>Eggerella bradyi</i>	Agglutinated	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore

<i>Eggerella sp</i>	Agglutinated	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Eggerelloides advena</i>	Agglutinated	Triserial	Infaunal	Oxic - Hypoxic	Detritivore
<i>Eggerelloides scabrum</i>	Agglutinated	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ehrenbergina compressa</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ehrenbergina trigona</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Epistominella afueraensis</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Epistominella exigua</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Epistominella obesa</i>	Hyaline	Trochospiral	Shallow Infaunal	Hypoxic - anoxic	Phytodetritivore
<i>Epistominella sp</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Eponides sp.</i>	Hyaline	Trochospiral	Epifaunal	Oxic	Phytodetritivore
<i>Eratidus foliaceus</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Ebuliminella basispinata</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ebuliminella curta</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Ebuliminella exilis</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Eusphaeroidina sp</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Evolutinella rotulata</i>	Agglutinated	Planispiral	Intermedia infaunal	Hypoxic	Detritivore
<i>Fissurina (Fissurina) ovata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina annectens</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina circularis</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina cucullata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina cucurbitasema</i>	Hyaline	Spheric unilocular	Shallow Infaunal	Oxic - Hypoxic	Detritivore

		and elongated spheric			
<i>Fissurina granifera</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina gravata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina laevigata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina marginata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina orbignyana</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina piriformis</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina sp.</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina submarginata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fissurina susanae</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Fursenkoina bradyi</i>	Hyaline	Biserial	Intermedia Infaunal	Hypoxic - anoxic	Detritivore
<i>Fursenkoina schreibersiana</i>	Hyaline	Biserial	Intermedia Infaunal	Hypoxic - anoxic	Detritivore
<i>Fursenkoina texturata</i>	Hyaline	Biserial	Intermedia Infaunal	Hypoxic - anoxic	Detritivore

<i>Galwayella trigonoornata</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic	Detritivore
<i>Glandulina laevigata</i>	Hyaline	Uniserial	Intermedia Infaunal	Oxic	Detritivore
<i>Glandulina ovula</i>	Hyaline	Uniserial	Intermedia Infaunal	Oxic	Detritivore
<i>Glandulina symmetrica</i>	Hyaline	Uniserial	Intermedia Infaunal	Oxic	Detritivore
<i>Glaphyrammina americana</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina affinis</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina auriculata</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Carnivore
<i>Globobulimina hoeglundi</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina notovata</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina pacifica</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina sp</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina spinifera</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina subaffinis</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Globobulimina turgida</i>	Hyaline	Triserial	Intermedia Infaunal	Oxic - Hypoxic	Carnivore
<i>Globocassidulina crassa</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic - Anoxic	Detritivore
<i>Globocassidulina minima</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Infaunal	Oxic - Anoxic	Detritivore
<i>Globocassidulina sp</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Infaunal	Oxic - Anoxic	Detritivore
<i>Globocassidulina subglobosa</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Epifaunal	Oxic - Anoxic	Detritivore
<i>Glomospira gordialis</i>	Agglutinated	Irregular	Epifauna/Shallow Infaunal	Hypoxic - anoxic	Phytodetritivore
<i>Gyroidina altiformis</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidina gemma</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidina lamarckiana</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore

<i>Gyroidina orbicularis</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidina polia</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidina rothwelli</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidina soldanii</i> <i>var multilocula</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Gyroidinoides</i> sp	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Hansenisca soldanii</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Anoxic	Detritivore
<i>Hansenisca</i> sp	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Hanzawaia boueana</i>	Hyaline	Trochospiral	Shallow Infaunal	Hypoxic	Phytodetritivore
<i>Hanzawaia mexicana</i>	Hyaline	Trochospiral	Shallow Infaunal	Hypoxic	Phytodetritivore
<i>Hanzawaia prona</i>	Hyaline	Trochospiral	Shallow Infaunal	Hypoxic	Phytodetritivore
<i>Haplophragmoides bulloides</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic	Phytodetritivore
<i>Haplophragmoides evoluta</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic	Phytodetritivore
<i>Haplophragmoides mexicanus</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic	Phytodetritivore
<i>Haplophragmoides planissima</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Haplophragmoides pusillus</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Haplophragmoides</i> sp	Agglutinated	Planispiral	Intermedia Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Haplophragmoides subsphaeroides</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Haplophragmoides wilberti</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic	Phytodetritivore
<i>Heterolepa bradyi</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Heterolepa broeckiana</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Heterolepa</i> sp	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Hoeglundina elegans</i>	Hyaline	Trochospiral	Epifaunal/SI	Oxic - Hypoxic	Omnivore
<i>Hormosina globulifera</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Hormosinella distans</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Hormosinelloides guttifer</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Hyalinonettrion clavatum</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Hyalinonettrion gracillimum</i>	Hyaline	Spheric unilocular and	Intermedia infaunal	Oxic - Hypoxic	Detritivore

		elongated spheric			
<i>Hyperammina</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Hyperammina echinata</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Hyperammina elongata</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Hyperammina friabilis</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Karreriella baccata</i>	Agglutinated	Biserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Karreriella bradyi</i>	Agglutinated	Biserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Labrospira crassimargo</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Labrospira robusta</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Laevidentalina baggi</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Laevidentalina communis</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Laevidentalina elegans</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Laevidentalina filiformis</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Laevidentalina subsoluta</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic	Detritivore/ Phytodetritivore
<i>Lagena aspera</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic	Detritivore
<i>Lagena hispidula</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic	Detritivore
<i>Lagena parvulipora</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic	Detritivore
<i>Lagena striata</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic	Detritivore
<i>Lagena substriata</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Lagena sulcata</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia Infaunal	Oxic - Hypoxic	Detritivore

<i>Lagenammina ampullacea</i>	Agglutinated	Spheric unilocular and elongated spheric	Epifaunal/SI	Oxic	Bacterivore
<i>Lagenammina difflugiformis</i>	Agglutinated	Spheric unilocular and elongated spheric	Epifaunal/SI	Oxic - Hypoxic	Bacterivore
<i>Lenticulina convergens</i>	Hyaline	Planispiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Lenticulina gibba</i>	Hyaline	Planispiral	Epifaunal/SI	Oxic - Hypoxic	Detritivore
<i>Lenticulina inornata</i>	Hyaline	Planispiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Lenticulina pliocaena</i>	Hyaline	Planispiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Lenticulina rotulata</i>	Hyaline	Planispiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Lenticulina sp</i>	Hyaline	Planispiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Lenticulina thalmani</i>	Hyaline	Planispiral	Epifaunal	Hypoxic	Detritivore
<i>Liebusella goesi</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Lobatula lobatula</i>	Hyaline	Trochospiral	Epifaunal	Oxic	Phytodetritivore
<i>Lobatula ungeriana</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Lobatula wuellerstorfi</i>	Hyaline	Trochospiral	Epifaunal	Oxic	Phytodetritivore
<i>Loxostomina limbata</i>	Hyaline	Biserial	Intermedia Infaunal	Anoxic	Phytodetritivore
<i>Lugdunum hantkenianum</i>	Hyaline	Biserial	Intermedia Infaunal	Hypoxic - Anoxic	Phytodetritivore
<i>Marsipella elongata</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic	Detritivore/ Bacterivore
<i>Martinottiella communis</i>	Agglutinated	Uniserial	Intermedia Infaunal	Hypoxic	Detritivore
<i>Martinottiella nodulosa</i>	Agglutinated	Uniserial	Intermedia Infaunal	Hypoxic	Detritivore
<i>Melonis affinis</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Melonis pompilioides</i>	Hyaline	Planispiral	SI/II	Oxic - Hypoxic	Detritivore
<i>Miliolinella subrotunda</i>	Porcellaneous	Miliolide	Epifaunal	Oxic	Herbivore
<i>Nodosinum gaussicum</i>	Agglutinated	Uniserial	Intermedia Infaunal	Oxic	Detritivore
<i>Nodulina dentaliniformis</i>	Agglutinated	Uniserial	Intermedia Infaunal	Oxic - Hypoxic	Detritivore
<i>Nonion commune</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic - Anoxic	Detritivore
<i>Nonion pacificum</i>	Hyaline	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Nonionella astricta</i>	Hyaline	Trochospiral	Epifaunal	Hypoxic	Detritivore
<i>Nonionella auris</i>	Hyaline	Trochospiral	Epifaunal	Hypoxic	Detritivore

<i>Nonionella sp</i>	Hyaline	Trochospiral	Epifaunal	Hypoxic	Detritivore
<i>Nonionella stella</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Anoxic	Detritivore
<i>Nonionoides grateloupii</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Anoxic	Phytodetritivore
<i>Nonionoides turgidus</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Phytodetritivore
<i>Nubeculina sp.</i>	Agglutinated	Uniserial	Intermedia Infaunal	Hypoxic - Anoxic	Detritivore
<i>Nummoloculina sp</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Phytodetritivore
<i>Nuttallides umbonifer</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Obesopleurostomella brevis</i>	Hyaline	Biserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Oolina globosa</i>	Hyaline	Spheric unilocular and elongated spheric	Intermedia infaunal	Hypoxic	Detritivore
<i>Oridorsalis pauciapertura</i>	Hyaline	Trochospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Oridorsalis tenerus subsp. Profundus</i>	Hyaline	Trochospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Oridorsalis umbonatus</i>	Hyaline	Trochospiral	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Palliolatella bradii</i>	Hyaline	Trochospiral	Intermedia infaunal	Oxic	Detritivore
<i>Paracassidulina minuta</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Parafissurina lateralis</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Hypoxic	Detritivore
<i>Parafissurina ovata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Detritivore
<i>Parafissurina sp</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Hypoxic	Detritivore
<i>Parafissurina ventricosa</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Detritivore
<i>Paratrochammina challengerii</i>	Agglutinated	Trochospiral	Epifaunal	Oxic - Hypoxic	Phytodetritivore / Bacterivore

<i>Paratrochammina globorotaliformis</i>	Agglutinated	Trochospiral	Epifaunal	Oxic	Phytodetrivore / Bacterivore
<i>Planispirinoides bucculentus</i>	Porcellaneous	Miliolide	Intermedia Infaunal	Oxic	Detritivore
<i>Planulina ecuadorana</i>	Hyaline	Trochospiral	Epifaunal	Oxic	Phytodetrivore
<i>Polystomammina nitida</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Portatrochammina karica</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic	Phytodetrivore / Bacterivore
<i>Portatrochammina murrayi</i>	Agglutinated	Trochospiral	Epifaunal/SI	Oxic	Phytodetrivore /Bacterivore
<i>Portatrochammina pacifica</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic	Phytodetrivore /Bacterivore
<i>Praeglobobulimina primitiva</i>	Hyaline	Triserial	Intermedia Infaunal	Hypoxic	Detritivore
<i>Praeglobobulimina spinescens</i>	Hyaline	Triserial	Intermedia Infaunal	Hypoxic	Detritivore
<i>Procerolagena distoma</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Procerolagena gracilis</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Protoglobobulimina pupoides</i>	Hyaline	Triserial	Shallow Infaunal	Anoxic	Detritivore
<i>Psammotodendron arborescens</i>	Agglutinated	Irregular	Shallow Infaunal	Oxic	Bacterivore
<i>Psammosiphonella discreta</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Psammosphaera fusca</i>	Agglutinated	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Bacterivore
<i>Pseudobrizalina lobata</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Pseudonodosinella nodulosa</i>	Agglutinated	Uniserial	Intermedia Infaunal	Oxic	Detritivore
<i>Pseudoparrella pacifica</i>	Hyaline	Trochospiral	SI/II	Hypoxic - Anoxic	Omnivore
<i>Pseudoparrella subperuviana</i>	Hyaline	Trochospiral	SI/II	Hypoxic - Anoxic	Omnivore
<i>Pullenia bulloides</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Pullenia elegans</i>	Hyaline	Globose, subglobose,	Shallow Infaunal	Oxic	Detritivore

		ovoid or streptospiral			
<i>Pullenia quinqueloba</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Pullenia sp</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Pullenia subcarinata</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Pulleniatina obliquiloculata</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Pulleniella asymmetrica</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic	Detritivore
<i>Pyrgo depressa</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgo elongata</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgo murrhina</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgo sarsi</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgo sp</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgo subsphaerica</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgoella irregularis</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrgoella sphaera</i>	Porcellaneou s	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Pyrulina</i>	Hyaline	Biserial	Intermedia infaunal	Oxic	Detritivore
<i>Pyrulina angusta</i>	Hyaline	Biserial	Intermedia infaunal	Oxic	Detritivore
<i>Quinqueloculina seminulum</i>	Porcellaneou s	Miliolide	Epifaunal/SI	Oxic - Hypoxic	Herbivore
<i>Quinqueloculina sp</i>	Porcellaneou s	Miliolide	Epifaunal/SI	Oxic	Herbivore
<i>Quinqueloculina weaveri</i>	Porcellaneou s	Miliolide	Epifaunal/SI	Oxic	Herbivore
<i>Recurvoidella bradyi</i>	Agglutinated	Planispiral	Intermedia Infaunal	Oxic	Phytodetritivore
<i>Recurvoides contortus</i>	Agglutinated	Trochospiral	Shallow Infaunal	Oxic - Anoxic	Detritivore
<i>Recurvoides turbinatus</i>	Agglutinated	Trochospiral	Intermedia infaunal	Oxic	Detritivore
<i>Reophax apiculatus</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore

<i>Reophax bradyi</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax duplex</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax fusiformis</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax globosus</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax insectus</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax pilulifer</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax pisiformis</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax regularis</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax scorpiurus</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax sp</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reophax subdentaliniformis</i>	Agglutinated	Uniserial	SI/II	Oxic	Detritivore
<i>Reophax subfusiformis</i>	Agglutinated	Uniserial	Shallow Infaunal	Oxic	Detritivore
<i>Reusoolina apiculata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Detritivore
<i>Reusoolina laevis</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Rhabdammina abyssorum</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhabdammina pacifica</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhabdammina sp.</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhabdamminella cylindrica</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhizammina algaeformis</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhizammina sp</i>	Agglutinated	Tubular	Epifaunal	Oxic	Bacterivore
<i>Rhumblarella humboldti</i>	Agglutinated	Triserial	Epifaunal	Oxic - Anoxic	Detritivore
<i>Robertina sp</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Carnivore
<i>Robertinoides bradyi</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Carnivore
<i>Robertinoides oceanica</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic - Hypoxic	Carnivore
<i>Rosalina globularis</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Omnivore
<i>Rotalina pleurostomata</i>	Hyaline	Trochospiral	Shallow Infaunal	Oxic	Detritivore
<i>Rutherfordoides rotundatus</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Anoxic	Detritivore
<i>Saccamina sphaerica</i>	Agglutinated	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Bacterivore
<i>Saccorhiza ramosa</i>	Agglutinated	Tubular	Intermedia infaunal	Oxic	Bacterivore

<i>Sahulia conica</i>	Agglutinated	Biserial	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Saracenaria stolidota</i>	Hyaline	Planispiral	Intermedia infaunal	Hypoxic	Detritivore
<i>Sigmoilina sp</i>	Porcellaneous	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Siphouvigerina hispida</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Siphouvigerina proboscidea</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Sphaeroidina bulloides</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Sphaeroidina sp</i>	Hyaline	Globose, subglobose, ovoid or streptospiral	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Spirillina sp</i>	Hyaline	Planispiral	Epifaunal/Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Spiroloculina robusta</i>	Porcellaneous	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Spiroloculina sp</i>	Porcellaneous	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Spiroplectammina biformis</i>	Agglutinated	Biserial	Intermedia infaunal	Oxic - Hypoxic	Detritivore
<i>Spirosigmoilina tenuis</i>	Porcellaneous	Miliolide	Shallow Infaunal	Oxic	Herbivore
<i>Spirosigmoilinella sp.</i>	Agglutinated	Miliolide	Shallow Infaunal	Oxic	Detritivore
<i>Stainforthia fusiformes</i>	Hyaline	Biserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Suggrunda eckisi</i>	Hyaline	Biserial	Shallow Infaunal	Anoxic	Detritivore
<i>Suggrunda kleinPELLI</i>	Hyaline	Biserial	Shallow Infaunal	Anoxic	Detritivore
<i>Suggrunda porosa</i>	Hyaline	Biserial	Shallow Infaunal	Anoxic	Detritivore
<i>Takayanagia delicata</i>	Hyaline	Biserial	Intermedia infaunal	Oxic - Hypoxic	Omnivore
<i>Techitella melo</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic	Bacterivore
<i>Techitella sp.</i>	Agglutinated	Tubular	Shallow Infaunal	Oxic	Bacterivore
<i>Textularia earlandi</i>	Agglutinated	Biserial	SI/II	Oxic	Omnivore
<i>Textularia gramen</i>	Agglutinated	Biserial	SI/II	Oxic	Omnivore
<i>Textularia lythostrota</i>	Agglutinated	Biserial	SI/II	Oxic	Omnivore
<i>Textularia porrecta</i>	Agglutinated	Biserial	SI/II	Oxic	Omnivore
<i>Thurammina albicans</i>	Agglutinated	Spheric unilocular and elongated spheric	Shallow Infaunal	Oxic	Bacterivore
<i>Tolypammina vagans</i>	Agglutinated	Irregular	Shallow Infaunal	Oxic - Hypoxic	Bacterivore
<i>Trifarina angulosa</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - anoxic	Detritivore
<i>Triloculina tricarinata</i>	Porcellaneous	Miliolide	Epifaunal/SI	Oxic	Phytodetritivore

<i>Triloculina trigonula</i>	Porcellaneou s	Miliolide	Epifaunal/SI	Oxic	Herbivore
<i>Trochammina inflata</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Trochammina intermedia</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Trochammina nana</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Trochammina sp</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Trochammina squamata</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Phytodetrivore
<i>Trochammina triloba</i>	Agglutinated	Planispiral	Epifaunal/SI	Oxic	Phytodetrivore
<i>Uvigerina auberiana</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Uvigerina canariensis</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Uvigerina dirupta</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Hypoxic	Detritivore
<i>Uvigerina peregrina</i>	Hyaline	Triserial	Shallow Infaunal	Oxic - Anoxic	Omnivore
<i>Uvigerina semiornata</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic - Anoxic	Detritivore
<i>Uvigerina senticosa</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Uvigerina shiwoensis</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Uvigerina sp</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Uvigerina striata</i>	Hyaline	Triserial	Shallow Infaunal	Hypoxic	Detritivore
<i>Vaginulina americana</i>	Hyaline	Uniserial	Shallow Infaunal	Oxic	Omnivore
<i>Valvulineria araucana</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Valvulineria bradyana</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Valvulineria glabra</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Valvulineria inaequalis</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Valvulineria inflata</i>	Hyaline	Trochospiral	Epifaunal	Oxic - Hypoxic	Detritivore
<i>Vasicostella inflatiperforata</i>	Hyaline	Spheric unilocular and elongated spheric	Shallow Infaunal	Hypoxic	Detritivore
<i>Veleroninoides scitulus</i>	Agglutinated	Planispiral	Shallow Infaunal	Oxic	Detritivore
<i>Verneuilinulla propinqua</i>	Agglutinated	Triserial	Intermedia infaunal	Hypoxic	Detritivore

Appendix 6. Spearman's correlation coefficients between taxonomic and functional diversity and their  $p$ -value. Variables that are not significantly correlated are marked in bold. Variable "Taxa" = Species richness.

Variable 1	Variable 2	$\rho$	$p$ -value	Variable 1	Variable 2	$\rho$	$p$ -value
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Taxa	Simpson	0.639	p < 0.01	Pielou	Abundance	- 0.682	p < 0.01
Taxa	Shannon	0.794	p < 0.01	<b>Pielou</b>	<b>FRic</b>	<b>0.158</b>	<b>p ≥ 0.05</b>
<b>Taxa</b>	<b>Pielou</b>	<b>-0.040</b>	<b>p ≥ 0.05</b>	Pielou	FDiv	- 0.211	p < 0.05
Taxa	Fisher_a	0.631	p < 0.01	Pielou	FDis	0.532	p < 0.01
Taxa	Abundance	0.391	p < 0.01	Pielou	FEve	0.623	p < 0.01
Taxa	FRic	0.841	p < 0.01	Pielou	RaoQ	0.514	p < 0.01
<b>Taxa</b>	<b>FDiv</b>	<b>-0.057</b>	<b>p ≥ 0.05</b>	Fisher_a	Abundance	- 0.236	p < 0.05
Taxa	FDis	0.333	p < 0.01	Fisher_a	FRic	0.696	p < 0.01
Taxa	FEve	-0.257	p < 0.01	<b>Fisher_a</b>	<b>FDiv</b>	- <b>0.146</b>	<b>p ≥ 0.05</b>
Taxa	RaoQ	0.349	p < 0.01	Fisher_a	FDis	0.569	p < 0.01
Simpson	Shannon	0.962	p < 0.01	Fisher_a	FEve	0.337	p < 0.01
Simpson	Pielou	0.672	p < 0.01	Fisher_a	RaoQ	0.591	p < 0.01
Simpson	Fisher_a	0.834	p < 0.01	<b>Abundance</b>	<b>FRic</b>	<b>0.140</b>	<b>p ≥ 0.05</b>
<b>Simpson</b>	<b>Abundance</b>	<b>-0.179</b>	<b>p ≥ 0.05</b>	Abundance	FDiv	0.197	p < 0.05
Simpson	FRic	0.683	p < 0.01	Abundance	FDis	- 0.375	p < 0.01
Simpson	FDiv	-0.189	p < 0.05	Abundance	FEve	- 0.667	p < 0.01
Simpson	FDis	0.627	p < 0.01	Abundance	RaoQ	- 0.367	p < 0.01
Simpson	FEve	0.216	p < 0.05	<b>FRic</b>	<b>FDiv</b>	- <b>0.120</b>	<b>p ≥ 0.05</b>
Simpson	RaoQ	0.625	p < 0.01	FRic	FDis	0.494	p < 0.01
Shannon	Pielou	0.506	p < 0.01	<b>FRic</b>	<b>FEve</b>	- <b>0.012</b>	<b>p ≥ 0.05</b>
Shannon	Fisher_a	0.849	p < 0.01	FRic	RaoQ	0.526	p < 0.01
<b>Shannon</b>	<b>Abundance</b>	<b>-0.049</b>	<b>p ≥ 0.05</b>	<b>FDiv</b>	<b>FDis</b>	- <b>0.087</b>	<b>p ≥ 0.05</b>
Shannon	FRic	0.787	p < 0.01	<b>FDiv</b>	<b>FEve</b>	<b>0.080</b>	<b>p ≥ 0.05</b>
<b>Shannon</b>	<b>FDiv</b>	<b>-0.164</b>	<b>p ≥ 0.05</b>	<b>FDiv</b>	<b>RaoQ</b>	- <b>0.086</b>	<b>p ≥ 0.05</b>
Shannon	FDis	0.593	p < 0.01	FDis	FEve	0.414	p < 0.01
<b>Shannon</b>	<b>FEve</b>	<b>0.108</b>	<b>p ≥ 0.05</b>	FDis	RaoQ	0.990	p < 0.01
Shannon	RaoQ	0.602	p < 0.01	FEve	RaoQ	0.424	p < 0.01
Pielou	Fisher_a	0.586	p < 0.01				

Appendix 7. Niche parameters of all the species of benthic foraminifera in the marine surface sediments used in this study with a contribution > 0.05% in all the stations and their traits estimated by Outlying Mean Index (OMI). Inertia = total variability, OMI = Outlying Mean Index (i.e., marginality), Tol = Tolerance, RTol = Residual Tolerance, OMI% = percentage of variability given by OMI, Tol% = percentage of variability given by Tol, RTol% = percentage of variability given by RTol. Species in bold: *p*-value <0.05

\*Abbreviations of the traits. Shell: Group agglutinated (GA), Group hyaline (GH); Morpho: Morphology planispiral (MP), morphology tubular (MTub), morphology biserial (MB), morphology triserial (MT), morphology trochospiral (MTroc), morphology globose/subglobose/ovoid or streptospiral (MG), morphology uniserial (MU), morphology spheric unilocular and elongated spheric (MS). Microhabitat (Habit): group epifaunal (GE), group shallow infaunal (GSI) and group intermedia infaunal (GII). Trophic behaviour (Trophy): Detritivore (Det), Phytodetritivore (PhDet), Bacterivore (Bact) and Omnivore (Omni).

Specie	Wal l	Code	Morph o	Habi t	Troph y	inerti a	OMI	Tol	Rtol	omi %	tol%	rtol %	<i>p</i> -valu e
<i>Ammobaculites agglutinans</i>	GA	Aaggl	MP	GSI	Bact	17.12	2.05	1.09	13.98	12	6.40	81.70	0.54
<i>Ammodiscus catinus</i>	GA	Acati	MP	GE	Det	17.98	12.53	1.63	3.82	69.70	9	21.30	0.11
<i>Astrorhizoides cornutus</i>	GA	Acorn	MTub	GE	Bact	11.54	11.54	0	0	100	0	0	0.50
<i>Astrononion echolsi</i>	GH	Aecho	MP	GSI	PhDet	5.85	5.85	0	0	100	0	0	0.92
<i>Ammoglobigerina globigeriniformis</i>	GA	Aglobi	MTroc	GSI	Det	12.62	3.05	3.01	6.56	24.20	23.80	52.00	0.10
<i>Ammoglobigerina globulosa</i>	GA	Aglobu	MTroc	GSI	Det	5.99	3.29	0.04	2.65	54.90	0.70	44.30	0.93
<i>Adercotryma glomeratum</i>	GA	Aglom	MG	GE	Herb	11.87	1.53	1.22	9.12	12.90	10.30	76.90	0.88
<i>Ammosphaeroidina grandis</i>	GA	Agrand	MG	GII	Det	29.54	29.54	0	0	100	0	0	0.13
<i>Astrorhiza granulosa</i>	GA	Agranu	MI	GII	Bact	11.47	8.21	1.74	1.52	71.60	15.20	13.20	0.67
<b><i>Ammodiscus incertus</i></b>	<b>GA</b>	<b>Aince</b>	<b>MP</b>	<b>GE</b>	<b>Det</b>	<b>20.10</b>	<b>4.77</b>	<b>3.79</b>	<b>11.53</b>	<b>23.70</b>	<b>18.90</b>	<b>57.40</b>	<b>0.04</b>
<i>Astacolus insolitus</i>	GH	Ainso	MP	GSI	Det	8.38	5.67	0.04	2.66	67.70	0.50	31.80	0.72
<i>Alveolophragmium orbiculatum</i>	GA	Aorb	MP	GSI	Det	5.32	5.23	0.01	0.08	98.40	0.10	1.50	0.65
<i>Ammodiscus planorbis</i>	GA	Aplan	MP	GE	Det	10.08	8.37	0.37	1.34	83	3.70	13.30	0.09
<i>Astrammmina rara</i>	GA	Arara	MI	GII	Car	17.34	8.14	2.80	6.41	46.90	16.10	36.90	0.38
<i>Astrononion stellatum</i>	GH	Astel	MP	GSI	PhDet	5.85	5.85	0	0	100	0	0	0.93
<i>Ammodiscus tenuis</i>	GA	Atenu	MP	GE	Det	7.58	2.21	0.62	4.76	29.20	8.10	62.70	0.95
<i>Bolivina alata</i>	GH	Balat	MB	GSI	PhDet	14.97	4.18	3.76	7.03	28	25.10	46.90	0.12
<i>Bathysiphon sp.</i>	GA	Bat	MTub	GSI	PhDet	18.02	18.02	0	0	100	0	0	0.25
<i>Bolivina (Loxostomum) boltovskoyi</i>	GH	Bbolt	MB	GE	PhDet	18.02	18.02	0	0	100	0	0	0.21
<b><i>Bathysiphon capillare</i></b>	<b>GA</b>	<b>Bcapi</b>	<b>MTub</b>	<b>GSI</b>	<b>PhDet</b>	<b>24.62</b>	<b>24.45</b>	<b>0</b>	<b>0.16</b>	<b>99.30</b>	<b>0</b>	<b>0.70</b>	<b>0.02</b>
<b><i>Bolivina costata</i></b>	<b>GH</b>	<b>Bcost</b>	<b>MB</b>	<b>GSI</b>	<b>PhDet</b>	<b>24.40</b>	<b>12.67</b>	<b>2.50</b>	<b>9.24</b>	<b>51.90</b>	<b>10.20</b>	<b>37.90</b>	<b>0.01</b>
<b><i>Buliminella elegantissima limbosa</i></b>	<b>GH</b>	<b>Belim</b>	<b>MTroc</b>	<b>GE</b>	<b>Det</b>	<b>31.44</b>	<b>31.02</b>	<b>0.03</b>	<b>0.40</b>	<b>98.60</b>	<b>0.10</b>	<b>1.30</b>	<b>0.01</b>

<i>Bulimina elongata</i>	GH	Belon	MT	GII	Det	10.63	10.6 3	0	0	100	0	0	0.69
<b><i>Buliminella elegantissima tenuis</i></b>	<b>GH</b>	<b>Belte</b>	<b>MTroc</b>	<b>GE</b>	<b>Det</b>	<b>30.96</b>	<b>30.5 0</b>	<b>0.0 2</b>	<b>0.45</b>	<b>98.50</b>	<b>0.1</b>	<b>1.4</b>	<b>0.01</b>
<i>Bathysiphon fusca</i>	GA	Bfusc	MTub	GSI	PhDet	10.12	10.1 2	0	0	100	0	0	0.65
<i>Bolivina humilis</i>	GH	Bhumi	MB	GSI	PhDet	8.31	8.31	0	0	100	0	0	0.81
<b><i>Bolivina interjuncta</i></b>	<b>GH</b>	<b>Binte</b>	<b>MB</b>	<b>GE</b>	<b>PhDet</b>	<b>18.74</b>	<b>5.64</b>	<b>1.9 2</b>	<b>11.1 8</b>	<b>30.1</b>	<b>10.3</b>	<b>59.7</b>	<b>0.02</b>
<i>Bulimina marginata</i>	GH	Bmarg	MT	GII	Det	7.46	6.40	0.0 1	1.04	85.9	0.1	14.0	0.51
<b><i>Bolivinita minuta</i></b>	<b>GH</b>	<b>Bminu</b>	<b>MB</b>	<b>GE</b>	<b>PhDet</b>	<b>15.21</b>	<b>9.00</b>	<b>0.3 3</b>	<b>5.88</b>	<b>59.2</b>	<b>2.2</b>	<b>38.7</b>	<b>0.01</b>
<i>Bolivina pacifica</i>	GH	Bpaci	MB	GE	PhDet	12.03	9.32	1.2 0	1.51	77.4	10	12.6	0.44
<b><i>Bulimina pagoda</i></b>	<b>GH</b>	<b>Bpago</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>20.44</b>	<b>17.9 9</b>	<b>0.4 9</b>	<b>1.95</b>	<b>88</b>	<b>2.4</b>	<b>9.6</b>	<b>0.05</b>
<i>Buccella peruviana</i>	GH	Bperu	MTroc	GSI	Det	20.24	14.5 9	1.1	4.55	72.1	5.4	22.5	0.10
<b><i>Bolivina plicata</i></b>	<b>GH</b>	<b>Bplic</b>	<b>MB</b>	<b>GE</b>	<b>PhDet</b>	<b>20.80</b>	<b>7.70</b>	<b>4.9 4</b>	<b>8.16</b>	<b>37</b>	<b>23.7</b>	<b>39.2</b>	<b>0.01</b>
<i>Bolivina (Loxostomum) salvadorensis</i>	GH	Bsalv	MB	GE	PhDet	18.02	18.0 2	0	0	100	0	0	0.32
<b><i>Bolivina seminuda</i></b>	<b>GH</b>	<b>Bsemi</b>	<b>MB</b>	<b>GE</b>	<b>PhDet</b>	<b>28.15</b>	<b>15.2 8</b>	<b>5.2 2</b>	<b>7.65</b>	<b>54.3</b>	<b>18.5</b>	<b>27.2</b>	<b>0.01</b>
<i>Bolivina spissa</i>	GH	Bspis	MB	GE	PhDet	11.53	2.20	1.6 9	7.65	19.1	14.6	66.3	0.10
<i>Bulimina striata</i>	GH	Bstri	MT	GSI	Det	7.58	3.93	0.0 3	3.62	51.9	0.4	47.8	0.19
<i>Bolivina subadvena</i>	GH	Bsuba	MB	GE	PhDet	23.42	23.4 2	0	0	100	0	0	0.15
<b><i>Bolivina subadvena subsp. Acuminata</i></b>	<b>GH</b>	<b>Bsubac u</b>	<b>MB</b>	<b>GE</b>	<b>PhDet</b>	<b>26.57</b>	<b>19.7 7</b>	<b>1.0 1</b>	<b>5.79</b>	<b>74.4</b>	<b>3.8</b>	<b>21.8</b>	<b>0.01</b>
<i>Bolivina subaenariensis</i>	GH	Bsubae	MB	GSI	PhDet	17.94	17.9 4	0	0	100	0	0	0.30
<i>Bulimina sp.</i>	GH	Bul	MT	GII	Det	7.71	7.71	0	0	100	0	0	0.90
<i>Buzasina sp.</i>	GA	Buz	MP	GII	Det	22.81	22.8 1	0	0	100	0	0	0.20
<i>Bolivina variabilis</i>	GH	Bvari	MB	GE	PhDet	10.63	10.6 3	0	0	100	0	0	0.63
<i>Cancris sp</i>	GH	Can	MTroc	GII	Det	9.73	9.73	0	0	100	0	0	0.78
<b><i>Cassidulina auka</i></b>	<b>GH</b>	<b>Cauka</b>	<b>MP</b>	<b>GII</b>	<b>PhDet</b>	<b>19.67</b>	<b>11.1 9</b>	<b>1.5 1</b>	<b>6.96</b>	<b>56.9</b>	<b>7.7</b>	<b>35.4</b>	<b>0.01</b>
<i>Cancris auricula</i>	GH	Cauri	MTroc	GII	Det	14.35	14.3 5	0	0	100	0	0	0.32
<i>Cyclammina cancellata</i>	GA	Ccanc	MP	GSI	Det	12.82	1.05	1.1 6	10.6 2	8.2	9	82.8	0.48
<i>Cassidulina carinata</i>	GH	Ccari	MP	GII	PhDet	9.94	2.20	1.4 8	6.27	22.1	14.9	63	0.15
<b><i>Cancris carmenensis</i></b>	<b>GH</b>	<b>Ccarm</b>	<b>MTroc</b>	<b>GII</b>	<b>Det</b>	<b>19.05</b>	<b>10.1 2</b>	<b>2.8 3</b>	<b>6.09</b>	<b>53.1</b>	<b>14.9</b>	<b>32</b>	<b>0.01</b>
<i>Cibicoides dispars</i>	GH	Cdisp	MTroc	GE	PhDet	11.27	11.2 7	0	0	100	0	0	0.63
<i>Cribrostomoides evolutus</i>	GA	Cevol	MP	GII	PhDet	11.20	3.52	0.2 7	7.41	31.4	2.4	66.2	0.49
<i>Chilostomella sp</i>	GH	Chilo	MG	GII	Det	10.63	10.6 3	0	0	100	0	0	0.62
<i>Cibicoides sp</i>	GH	Cibic	MTroc	GE	PhDet	8.95	8.95	0	0	100	0	0	0.71
<i>Cornuspira involvens</i>	GP	Cinvo	MP	GE	PhDet	7.88	6.99	0.0 2	0.87	88.7	0.2	11	0.74
<i>Cribrostomoides jeffreysii</i>	GA	Cjeff	MP	GSI	Det	5.85	5.85	0	0	100	0	0	0.95
<i>Cassidulina laevigata</i>	GH	Claev	MP	GII	PhDet	11.98	8.82	0.0 1	3.15	73.6	0.1	26.3	0.53
<i>Cibicoides mckannai</i>	GH	Cmcka	MTroc	GE	PhDet	15.88	15.8 8	0	0	100	0	0	0.38
<i>Cibicoides mundulus</i>	GH	Cmund	MTroc	GE	PhDet	10.27	5.74	1.0 6	3.47	55.9	10.3	33.8	0.07

<i>Cancris oblongus</i>	GH	Coblo	MTroc	GII	Det	12.14	3.91	0.0 3	8.2	32.2	0.2	67.6	0.17
<i>Chilostomella oolina</i>	GH	Cooli	MG	GII	Det	12.75	2.24	0.7 9	9.72	17.6	6.2	76.3	0.44
<i>Chilostomella ovoidea</i>	GH	Covoi	MG	GII	Det	14.01	8.14	1.1 1	4.76	58.1	7.9	34.0	0.09
<i>Cibicides pachyderma</i>	GH	Cpach	MTroc	GE	PhDet	8.79	7.47	0.2 2	1.1	84.9	2.5	12.6	0.23
<i>Cribragoesella pacifica</i>	GA	Cpaci	MTroc	GII	Det	10.34	10.3 4	0	0	100	0	0	0.63
<i>Cassidulinoides parkeriana</i>	GH	Cpark	MP	GII	PhDet	8.64	5.60	0.0 1	3.03	64.8	0.1	35	0.66
<i>Crithionina pisum</i>	GA	Cpisu	MS	GSI	Bact	9.01	7.32	0.2 4	1.45	81.2	2.7	16.1	0.41
<i>Cibicoides robertsonianus</i>	GH	Crobe	MTroc	GE	PhDet	10.63	10.6 3	0	0	100	0	0	0.63
<i>Cribrostomoides sphaerilocula</i>	GA	Cspha	MP	GSI	Det	14.45	5.19	1.1 8	8.08	35.9	8.2	55.9	0.49
<i>Cribrostomoides subglobosus</i>	GA	Csubg	MP	GSI	Det	11.48	9.31	0.0 1	2.15	81.2	0.1	18.8	0.36
<i>Cyclammina trullissata</i>	GA	Ctrul	MP	GSI	Det	9.73	9.73	0	0	100	0	0	0.77
<i>Cibicoides variabilis</i>	GH	Cvari	MTroc	GE	PhDet	4.11	4.11	0	0	100	0	0	1.00
<i>Discorbina bertheloti</i>	GH	Dbert	MTroc	GE	Herb	14.35	14.3 5	0	0	100	0	0	0.46
<i>Dorothia bradyana</i>	GA	Dbrad	MB	GII	Det	21.76	21.7 6	0	0	100	0	0	0.24
<i>Discorbis sp</i>	GH	Disc	MTroc	GE	Herb	10.31	9.53	0.1 5	0.64	92.4	1.4	6.2	0.43
<i>Discorbis peruvianus</i>	GH	Dperu	MTroc	GE	Herb	9.73	9.73	0	0	100	0	0	0.72
<i>Dorothia pseudoturris</i>	GA	Dpseu	MB	GII	Det	15.88	15.8 8	0	0	100	0	0	0.31
<b><i>Dorothia scabra</i></b>	<b>GA</b>	<b>Dscab</b>	<b>MB</b>	<b>GII</b>	<b>Det</b>	<b>22.26</b>	<b>20.2 7</b>	<b>0.0 1</b>	<b>1.98</b>	<b>91.1</b>	<b>0.0</b>	<b>8.9</b>	<b>0.02</b>
<b><i>eggerelloides advena</i></b>	<b>GA</b>	<b>Eadve</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>31.91</b>	<b>23.2 7</b>	<b>2.7 0</b>	<b>5.93</b>	<b>72.9</b>	<b>8.5</b>	<b>18.6</b>	<b>0.02</b>
<i>Eubuliminella basispinata</i>	GH	Ebasi	MT	GSI	Det	16.40	7.31	3.1 2	5.97	44.5	19.0	36.4	0.07
<b><i>eggerella bradyi</i></b>	<b>GA</b>	<b>Ebrad</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>16.31</b>	<b>6.72</b>	<b>0.3 1</b>	<b>9.28</b>	<b>41.2</b>	<b>1.9</b>	<b>56.9</b>	<b>0.02</b>
<i>Ehrenbergina compressa</i>	GH	Ecomp	MB	GSI	Det	13.19	8.68	0.1 7	4.34	65.8	1.3	32.9	0.09
<b><i>Eubuliminella curta</i></b>	<b>GH</b>	<b>Ecurt</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>32.05</b>	<b>31.6 7</b>	<b>0.0 3</b>	<b>0.35</b>	<b>98.8</b>	<b>0.1</b>	<b>1.1</b>	<b>0.02</b>
<i>Epistominella exigua</i>	GH	Eexig	MTroc	GSI	PhDet	18.26	2.7	3.4	12.1 6	14.8	18.6	66.6	0.41
<i>Eubuliminella exilis</i>	GH	Eexil	MT	GSI	Det	23.30	23.3	0	0	100	0	0	0.10
<i>Eratidus foliaceus</i>	GA	Efoli	MP	GSI	Bact	11.97	4.63	2.5 5	4.78	38.7	21.3	40	0.30
<b><i>Epistominella obesa</i></b>	<b>GH</b>	<b>Eobes</b>	<b>MTroc</b>	<b>GSI</b>	<b>PhDet</b>	<b>19.53</b>	<b>11.0 2</b>	<b>0.5 4</b>	<b>7.97</b>	<b>56.4</b>	<b>2.8</b>	<b>40.8</b>	<b>0.01</b>
<i>Epistominella sp</i>	GH	Epist	MTroc	GSI	PhDet	21.76	21.7 6	0	0	100	0	0	0.25
<i>Eponides sp.</i>	GH	Epon	MTroc	GE	PhDet	22.81	22.8 1	0	0	100	0	0	0.18
<i>Epistominella pulchella</i>	GH	Epulc	MTroc	GSI	PhDet	11.27	11.2 7	0	0	100	0	0	0.60
<i>Evolutinella rotulata</i>	GA	Erotu	MP	GII	Det	13.23	5.72	1.7 5	5.75	43.3	13.2	43.5	0.73
<b><i>eggerelloides scabrum</i></b>	<b>GA</b>	<b>Escab</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>17.62</b>	<b>5.56</b>	<b>0.9 2</b>	<b>11.1 4</b>	<b>31.5</b>	<b>5.2</b>	<b>63.2</b>	<b>0.05</b>
<i>eggerella sp</i>	GA	Esp	MT	GSI	Det	10.63	10.6 3	0	0	100	0	0	0.56
<i>Ehrenbergina trigona</i>	GH	Etrig	MB	GSI	Det	22.81	22.8 1	0	0	100	0	0	0.15
<b><i>Fissurina annectens</i></b>	<b>GH</b>	<b>Fanne</b>	<b>MS</b>	<b>GSI</b>	<b>Det</b>	<b>15.97</b>	<b>12.0 5</b>	<b>0.3 4</b>	<b>3.58</b>	<b>75.5</b>	<b>2.1</b>	<b>22.4</b>	<b>0.04</b>
<i>Fursenkoina bradyi</i>	GH	Fbrad	MB	GII	Det	18.02	18.0 2	0	0	100	0	0	0.25
<i>Fissurina gravata</i>	GH	Fgrav	MS	GSI	Det	7.71	7.71	0	0	100	0	0	0.86

<i>Fissurina marginata</i>	GH	Fmarg	MS	GSI	Det	10.54	1.03	1.0 5	8.46	9.8	9.9	80.3	0.98
<i>Fissurina orbignyana</i>	GH	Forbi	MS	GSI	Det	11.22	9.27	0	1.95	82.6	0.0	17.4	0.28
<i>Fissurina (Fissurina) ovata</i>	GH	Fovat	MS	GSI	Det	6.58	6.58	0	0	100	0	0	0.91
<i>Fissurina piriformis</i>	GH	Fpiri	MS	GSI	Det	8.95	8.95	0	0	100	0	0	0.88
<i>Fursenkoina schreibersiana</i>	GH	Fschr	MB	GII	Det	12.86	12.8 6	0	0	100	0	0	0.49
<i>Fissurina sp.</i>	GH	Fsp	MS	GSI	Det	23.42	23.4 2	0	0	100	0	0	0.14
<i>Fissurina submarginata</i>	GH	Fsubm	MS	GSI	Det	15.42	11.0 3	0.4 7	3.92	71.5	3	25.4	0.16
<b><i>Globobulimina affinis</i></b>	<b>GH</b>	<b>Gaffi</b>	<b>MT</b>	<b>GII</b>	<b>Det</b>	<b>14.33</b>	<b>5.96</b>	<b>2.9</b>	<b>5.46</b>	<b>41.6</b>	<b>20.3</b>	<b>38.1</b>	<b>0.04</b>
<i>Gyroidina altiformis</i>	GH	Galti	MTroc	GE	Det	10.89	6.51	1.5 5	2.83	59.8	14.3	26	0.06
<i>Glaphyrammina americana</i>	GA	Gamer	MP	GSI	Det	11.34	1.86	1.0 0	8.47	16.40	8.90	74.7 0	0.22
<i>Globobulimina auriculata</i>	GH	Gauri	MT	GII	Car	7.42	5.01	0.1 2	2.29	67.60	1.60	30.8 0	0.70
<b><i>Globocassidulina crassa</i></b>	<b>GH</b>	<b>Gcras</b>	<b>MG</b>	<b>GSI</b>	<b>Det</b>	<b>17.38</b>	<b>8.65</b>	<b>1.7 0</b>	<b>7.04</b>	<b>49.80</b>	<b>9.80</b>	<b>40.5 0</b>	<b>0.01</b>
<i>Gyroidina gemma</i>	GH	Ggemm	MTroc	GE	Det	6.55	6.05	0.0 4	0.46	92.40	0.50	7.10	0.58
<i>Glomospira gordialis</i>	GA	Ggord	MI	GE	PhDet	14.47	10.9 3	0.0 0	3.54	75.50	0.00	24.5 0	0.23
<i>Globobulimina hoeglundi</i>	GH	Ghoeg	MT	GII	Det	14.46	14.4 6	0	0	100	0	0	0.29
<i>Glandulina laevigata</i>	GH	Glaev	MU	GII	Det	7.93	7.12	0.0 4	0.77	89.80	0.60	9.70	0.50
<i>Gyroidina polia</i>	GH	Gorbi	MTroc	GE	Det	7.17	4.62	0.1 9	2.37	64.40	2.60	33.0 0	0.81
<i>Glandulina ovula</i>	GH	Govul	MU	GII	Det	6.58	6.58	0	0	100	0	0	0.87
<i>Globobulimina pacifica</i>	GH	Gpaci	MT	GII	Det	16.07	6.13	4.8 0	5.15	38.10	29.8 0	32.0 0	0.16
<i>Gyroidina orbicularis</i>	GH	Gpoli	MTroc	GE	Det	7.71	7.71	0	0	100	0	0	0.75
<i>Gyroidina rothwelli</i>	GH	Groth	MTroc	GE	Det	21.76	21.7 6	0	0	100	0	0	0.18
<b><i>Gyroidina soldanii var multifocula</i></b>	<b>GH</b>	<b>Gsoldm</b>	<b>MTroc</b>	<b>GE</b>	<b>Det</b>	<b>17.58</b>	<b>10.1 1</b>	<b>0.5 4</b>	<b>6.93</b>	<b>57.5</b>	<b>3.1</b>	<b>39.4</b>	<b>0.01</b>
<i>Globobulimina subaffinis</i>	GH	Gsuba	MT	GII	Det	7.95	4.33	0.7 5	2.86	54.5	9.5	36	0.63
<i>Globocassidulina subglobosa</i>	GH	Gsubg	MG	GE	Det	10.33	2.84	2.8 3	4.65	27.5	27.4	45	0.44
<i>Galwayella trigonoornata</i>	GH	Gtrig	MS	GII	Det	7.71	7.71	0	0	100	0	0	0.88
<i>Globobulimina turgida</i>	GH	Gturg	MT	GII	Car	8.14	3.76	0.0 7	4.31	46.2	0.8	53	0.86
<b><i>Hanzawaia mexicana</i></b>	<b>GH</b>	<b>Hamex</b>	<b>MTroc</b>	<b>GSI</b>	<b>PhDet</b>	<b>29.86</b>	<b>21.2 0</b>	<b>3.0 9</b>	<b>5.56</b>	<b>71</b>	<b>10.3</b>	<b>18.6</b>	<b>0.05</b>
<i>Hanzawaia boueana</i>	GH	Hboue	MTroc	GSI	PhDet	17.94	17.9 4	0	0	100	0	0	0.26
<i>Haplophragmoides bulloides</i>	GA	Hbull	MP	GII	PhDet	5.85	5.85	0	0	100	0	0	0.90
<i>Hyalinonetrion clavatum</i>	GH	Hclav	MS	GII	Det	23.30	23.3 0	0	0	100	0	0	0.16
<i>Hormosinella distans</i>	GA	Hdist	MU	GSI	Det	11.96	5.86	1.1 9	4.91	49	10	41	0.08
<i>Heterolepa bradyi</i>	GH	Hebrad	MTroc	GSI	Det	8.94	4.33	0.2 0	4.41	48.4	2.3	49.3	0.29
<i>Hyperammina echinata</i>	GA	Hechi	MTub	GSI	Bact	13.42	13.4 2	0	0	100	0	0	0.44
<b><i>Hoeglundina elegans</i></b>	<b>GH</b>	<b>Heleg</b>	<b>MTroc</b>	<b>GE</b>	<b>Omni</b>	<b>11.83</b>	<b>1.65</b>	<b>1.4 2</b>	<b>8.76</b>	<b>13.9</b>	<b>12.0</b>	<b>74.1</b>	<b>0.03</b>
<i>Hyperammina elongata</i>	GA	Helon	MTub	GSI	Bact	9.64	9.23	0.0 7	0.34	95.7	0.8	3.5	0.23
<i>Hyperammina friabilis</i>	GA	Hfria	MTub	GSI	Bact	15.88	15.8 8	0	0	100	0	0	0.36
<b><i>Hormosina globulifera</i></b>	<b>GA</b>	<b>Hglob</b>	<b>MU</b>	<b>GSI</b>	<b>Det</b>	<b>13.68</b>	<b>6.88</b>	<b>0.4 5</b>	<b>6.35</b>	<b>50.3</b>	<b>3.3</b>	<b>46.4</b>	<b>0.05</b>

<b><i>Hyalinonettrion gracillimum</i></b>	<b>GH</b>	<b>Hgrac</b>	<b>MS</b>	<b>GII</b>	<b>Det</b>	<b>17.23</b>	<b>8.28</b>	<b>3.87</b>	<b>5.08</b>	<b>48.1</b>	<b>22.5</b>	<b>29.5</b>	<b>0.04</b>
<i>Hormosinelloides guttifer</i>	GA	Hgutt	MU	GSI	Det	9.97	8.22	0.88	0.87	82.5	8.8	8.7	0.41
<i>Haplophragmoides mexicanus</i>	GA	Hmexi	MP	GII	PhDet	14.62	9.82	0.05	4.75	67.2	0.3	32.5	0.31
<i>Haplophragmoides planissima</i>	GA	Hplan	MP	GII	PhDet	14.26	9.85	0.06	4.36	69	0.4	30.6	0.33
<b><i>Haplophragmoides pusillus</i></b>	<b>GA</b>	<b>Hpusi</b>	<b>MP</b>	<b>GII</b>	<b>PhDet</b>	<b>15.21</b>	<b>6.44</b>	<b>3.16</b>	<b>5.6</b>	<b>42.4</b>	<b>20.8</b>	<b>36.8</b>	<b>0.04</b>
<i>Hansenisca soldanii</i>	GH	Hsold	MTroc	GSI	Det	15.00	1.43	4.67	8.9	9.5	31.1	59.3	0.19
<i>Haplophragmoides sp</i>	GA	Hsp	MP	GII	PhDet	11.88	11.88	0	0	100	0	0	0.50
<i>Haplophragmoides subsphaeroides</i>	GA	Hsubs	MP	GII	PhDet	10.83	10.83	0	0	100	0	0	0.60
<i>Haplophragmoides wilberti</i>	GA	Hwilb	MP	GII	PhDet	11.54	11.54	0	0	100	0	0	0.58
<i>Hyperammina</i>	GA	Hyp	MTub	GSI	Bact	20.38	10.29	3.8	6.29	50.5	18.7	30.9	0.08
<i>Karrerella baccata</i>	GA	Kbacc	MB	GSI	Det	22.68	13.71	0.73	8.24	60.5	3.2	36.3	0.16
<i>Karrerella bradyi</i>	GA	Kbrad	MB	GSI	Det	17.34	17.34	0	0	100	0	0	0.27
<i>Lagena aspera</i>	GH	Laspe	MS	GII	Det	23.30	23.30	0	0	100	0	0	0.24
<i>Lagena substriata</i>	GH	Lasub	MS	GII	Det	13.42	13.42	0	0	100	0	0	0.36
<i>Laevidentalina baggi</i>	GH	Lbagg	MU	GSI	Det	21.76	21.76	0	0	100	0	0	0.25
<i>Laevidentalina communis</i>	GH	Lcomm	MU	GSI	Det	7.69	5.82	0	1.9	75.8	0.1	24.2	0.53
<i>Lenticulina convergens</i>	GH	Lconv	MP	GE	Det	21.77	14.50	2.22	5.05	66.60	10.20	23.20	0.07
<i>Labrospira crassimargo</i>	GA	Lcras	MP	GSI	Det	11.25	8.07	0.33	2.86	71.70	2.90	25.40	0.50
<i>Lagenammina diffugiiformis</i>	GA	Ldiff	MS	GE	Bact	14.28	4.20	1.26	8.82	29.40	8.80	61.80	0.06
<i>Laevidentalina filiformis</i>	GH	Lfili	MU	GSI	Det	35.27	35.27	0	0	100	0	0	0.07
<i>Lenticulina gibba</i>	GH	Lgibb	MP	GE	Det	23.42	23.42	0	0	100	0	0	0.16
<i>Liebusella goesi</i>	GA	Lgoes	MU	GSI	Det	12.36	10.22	0.01	2.13	82.70	0.10	17.20	0.43
<i>Lugdunum hantkenianum</i>	GH	Lhant	MB	GII	PhDet	15.88	15.88	0	0	100	0	0	0.32
<i>Lagena hispidula</i>	GH	Lhisp	MS	GII	Det	11.86	2.95	1.18	7.73	24.90	9.90	65.20	0.69
<i>Lenticulina inomata</i>	GH	Linom	MP	GE	Det	23.30	23.30	0	0	100	0	0	0.19
<i>Lobatula lobatula</i>	GH	Lloba	MTroc	GE	PhDet	11.31	5.14	1.31	4.85	45.40	11.60	42.90	0.47
<i>Lagena parvulipora</i>	GH	Lparv	MS	GII	Det	8.95	8.95	0	0	100	0	0	0.76
<b><i>Lenticulina pliocaena</i></b>	<b>GH</b>	<b>Lplio</b>	<b>MP</b>	<b>GE</b>	<b>Det</b>	<b>30.78</b>	<b>21.77</b>	<b>1.25</b>	<b>7.76</b>	<b>70.70</b>	<b>4.00</b>	<b>25.20</b>	<b>0.02</b>
<b><i>Labrospira robusta</i></b>	<b>GA</b>	<b>Lrobu</b>	<b>MP</b>	<b>GSI</b>	<b>Det</b>	<b>17.14</b>	<b>10.56</b>	<b>0.06</b>	<b>6.52</b>	<b>61.60</b>	<b>0.30</b>	<b>38.10</b>	<b>0.04</b>
<i>Lenticulina rotulata</i>	GH	Lrotu	MP	GE	Det	14.47	7.23	2.58	4.67	49.90	17.80	32.30	0.08
<i>Lenticulina sp</i>	GH	Lsp	MP	GE	Det	23.30	23.30	0	0	100	0	0	0.20
<i>Laevidentalina subsoluta</i>	GH	Lsubs	MU	GSI	Det	10.63	10.63	0	0	100	0	0	0.59
<i>Lagena sulcata</i>	GH	Lsulc	MS	GII	Det	13.42	13.42	0	0	100	0	0	0.48
<i>Lobatula ungeriana</i>	GH	Lunge	MTroc	GSI	Det	10.10	7.03	0.02	3.05	69.6	0.2	30.2	0.46
<i>Lobatula wuellerstorfi</i>	GH	Lwuel	MTroc	GE	PhDet	14.13	2.37	1.96	9.79	16.8	13.9	69.3	0.12
<b><i>Melonis affinis</i></b>	<b>GH</b>	<b>Maffi</b>	<b>MP</b>	<b>GSI</b>	<b>Det</b>	<b>11.82</b>	<b>2.25</b>	<b>1.17</b>	<b>8.40</b>	<b>19.1</b>	<b>9.9</b>	<b>71.1</b>	<b>0.02</b>

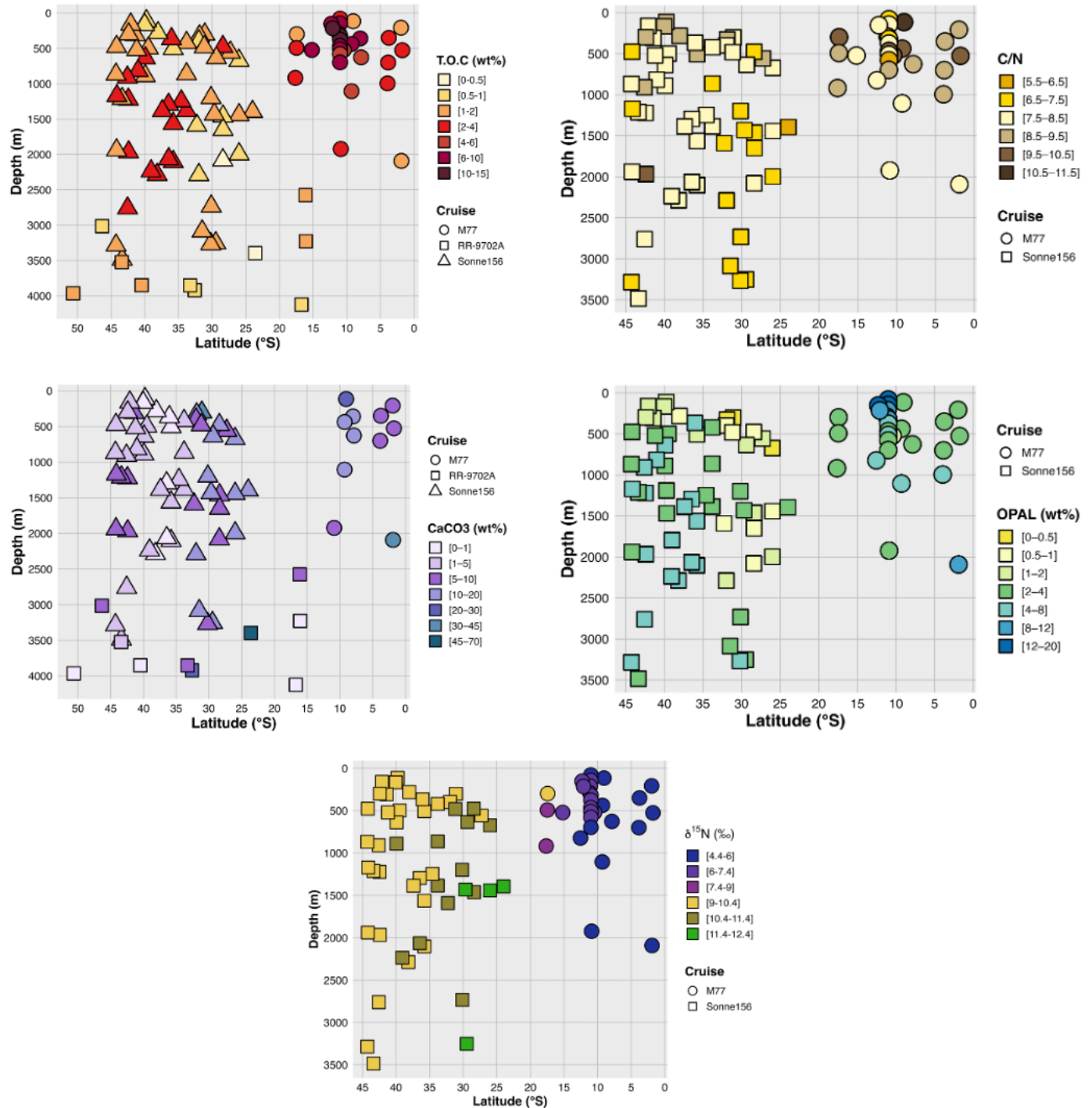
<i>Martinottiella communis</i>	GA	Mcomm	MU	GII	Det	24.47	22.0 3	0.0 3	2.40	90.1	0.1	9.8	0.02
<i>Marsipella elongata</i>	GA	Melon	MTub	GSI	Bact	9.23	7.00	0.7 7	1.45	75.9	8.4	15.7	0.04
<i>Martinottiella nodulosa</i>	GA	Mnodu	MU	GII	Det	14.59	8.70	1.2 5	4.64	59.7	8.6	31.8	0.57
<i>Melonis pompilioides</i>	GH	Mpomp	MP	GII	Det	13.17	10.3 3	1.1 9	1.64	78.5	9.1	12.5	0.19
<i>Miliolinella subrotunda</i>	GP	Msubr	MM	GE	Herb	12.63	11.5 2	0.1 1	1	91.3	0.8	7.9	0.17
<i>Nonionella astricta</i>	GH	Nastr	MTroc	GE	Det	31.30	31.3 0	0	0	100	0	0	0.11
<i>Nonion commune</i>	GH	Ncomm	MP	GSI	Det	12.47	1.62	1.4 8	9.37	13	11.9	75.1	0.56
<i>Nodulina dentaliniformis</i>	GA	Ndent	MU	GII	Det	13.83	1.77	1.8 1	10.2 6	12.8	13.1	74.1	0.17
<i>Nodosinum gaussicum</i>	GA	Ngaus	MU	GII	Det	6.58	6.58	0	0	100	0	0	0.92
<b><i>Nonionoides grateloupii</i></b>	<b>GH</b>	<b>Ngrat</b>	<b>MTroc</b>	<b>GSI</b>	<b>PhDet</b>	<b>16.02</b>	<b>5.63</b>	<b>4.2 6</b>	<b>6.13</b>	<b>35.2</b>	<b>26.6</b>	<b>38.2</b>	<b>0.05</b>
<i>Laevidentalina subsoluta</i>	GH	Nsubs	MU	GSI	PhDet	6.58	6.58	0	0	100	0	0	0.94
<i>Nonionoides turgidus</i>	GH	Nturg	MTroc	GSI	PhDet	9.73	9.73	0	0	100	0	0	0.65
<i>Obesopleurostomella brevis</i>	GH	Obrev	MB	GSI	Det	15.76	15.7 6	0	0	100	0	0	0.37
<i>Oridorsalis pauciapertura</i>	GH	Opauc	MTroc	GII	Det	13.42	13.4 2	0	0	100	0	0	0.43
<i>Oridorsalis tenerus subsp. Profundus</i>	GH	Otene	MTroc	GII	Det	11.88	11.8 8	0	0	100	0	0	0.52
<i>Oridorsalis umbonatus</i>	GH	Oumbo	MTroc	GII	Det	17.10	3.81	0.4 0	12.8 9	22.30	2.30	75.4 0	0.31
<i>Parafissurina sp</i>	GH	Paraf	MS	GSI	Det	13.57	13.5 7	0	0	100	0	0	0.38
<i>Psammotodendron arborescens</i>	GA	Parbo	MI	GSI	Bact	21.76	21.7 6	0	0	100	0	0	0.23
<i>Planispirinoides bucculentus</i>	GP	Pbucc	MM	GII	Det	22.81	22.8 1	0	0	100	0	0	0.18
<i>Pullenia bulloides</i>	GH	Pbull	MG	GSI	Det	12.24	3.58	0.5 1	8.15	29.20	4.20	66.6 0	0.25
<i>Paratrochammina challengerii</i>	GA	Pchal	MTroc	GE	Bact	10.01	6.10	0.6 5	3.26	60.90	6.50	32.5 0	0.41
<i>Pyrgo depressa</i>	GP	Pdepr	MM	GSI	Herb	7.71	7.71	0	0	100	0	0	0.90
<i>Psammosiphonella discreta</i>	GA	Pdisc	MTub	GE	Bact	11.83	5.53	0.5 4	5.76	46.70	4.60	48.7 0	0.24
<i>Planulina ecuadorana</i>	GH	Pecua	MTroc	GE	PhDet	21.63	21.6 3	0	0	100	0	0	0.26
<b><i>Pullenia elegans</i></b>	<b>GH</b>	<b>Peleg</b>	<b>MG</b>	<b>GSI</b>	<b>Det</b>	<b>18.52</b>	<b>10.6 7</b>	<b>0.3 1</b>	<b>7.54</b>	<b>57.60</b>	<b>1.70</b>	<b>40.7 0</b>	<b>0.01</b>
<i>Pyrgo elongata</i>	GP	Pelon	MM	GSI	Herb	11.54	11.5 4	0	0	100	0	0	0.55
<i>Procerolagena gracilis</i>	GH	Pgrac	MS	GSI	Det	23.30	23.3 0	0	0	100	0	0	0.17
<i>Pyrgoella irregularis</i>	GP	Pirre	MM	GSI	Herb	11.54	11.5 4	0	0	100	0	0	0.57
<i>Portatrochammina karica</i>	GA	Pkari	MTroc	GSI	Bact	13.57	13.5 7	0	0	100	0	0	0.46
<b><i>Pseudobrivalina lobata</i></b>	<b>GH</b>	<b>Ploba</b>	<b>MB</b>	<b>GSI</b>	<b>Det</b>	<b>22.02</b>	<b>13.2 4</b>	<b>0.3 3</b>	<b>8.45</b>	<b>60.10</b>	<b>1.50</b>	<b>38.4 0</b>	<b>0.03</b>
<i>Paracassidulina minuta</i>	GH	Pminu	MG	GSI	Det	6.58	6.58	0	0	100	0	0	0.90
<i>Pyrgo murrhina</i>	GP	Pmurr	MM	GSI	Herb	13.41	3.31	3.3 2	6.78	24.70	24.8 0	50.6 0	0.16
<i>Polystomammmina nitida</i>	GA	Pniti	MTroc	GSI	Det	9.66	5.89	0.0 2	3.76	61.00	0.20	38.9 0	0.61
<i>Pseudonodosinella nodulosa</i>	GA	Pnodu	MU	GII	Det	11.09	10.3 1	0.2 0	0.58	93.00	1.80	5.20	0.35
<i>Pulleniatina obliquiloculata</i>	GH	Pobli	MG	GSI	Det	8.95	8.95	0	0	100	0	0	0.76
<i>Portatrochammina murrayi</i>	GA	Pomur	MTroc	GE	Bact	10.63	10.6 3	0	0	100	0	0	0.66

<i>Pseudoparrella pacifica</i>	<b>GH</b>	<b>Ppaci</b>	<b>MTroc</b>	<b>GII</b>	<b>Omni</b>	<b>12.40</b>	<b>5.53</b>	<b>1.64</b>	<b>5.23</b>	<b>44.60</b>	<b>13.20</b>	<b>42.10</b>	<b>0.03</b>
<i>Protoglobulimina pupoides</i>	<b>GH</b>	<b>Ppupo</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>23.23</b>	<b>13.40</b>	<b>4.35</b>	<b>5.48</b>	<b>57.70</b>	<b>18.70</b>	<b>23.60</b>	<b>0.01</b>
<i>Pullenia quinqueloba</i>	GH	Pquin	MG	GSI	Det	12.26	4.85	0.79	6.61	39.60	6.50	53.90	0.13
<i>Praeglobulimina spinescens</i>	GH	Pspin	MT	GII	Det	8.06	6.60	0.40	1.06	81.90	4.90	13.20	0.73
<i>Pullenia subcarinata</i>	GH	Psubc	MG	GSI	Det	8.27	1.82	1.29	5.16	22.10	15.60	62.40	0.49
<i>Pseudoparrella subperuviana</i>	GH	Psubp	MTroc	GII	Omni	19.15	14.16	1.56	3.43	73.90	8.20	17.90	0.07
<i>Pullenia sp</i>	GH	Pulle	MG	GSI	Det	8.00	8.00	0	0	100	0	0	0.80
<i>Pyrgo sp</i>	GP	Pyrgo	MM	GSI	Herb	11.66	9.91	0.21	1.54	85.00	1.80	13.20	0.26
<i>Quinqueloculina seminulum</i>	GP	Qsemi	MM	GE	Herb	20.02	15.93	0.98	3.11	79.60	4.90	15.50	0.06
<i>Quinqueloculina sp</i>	GP	Qsp	MM	GE	Herb	11.54	11.54	0	0	100	0	0	0.56
<i>Rhabdammina abyssorum</i>	<b>GA</b>	<b>Rabys</b>	<b>MTub</b>	<b>GE</b>	<b>Bact</b>	<b>9.80</b>	<b>8.81</b>	<b>0.15</b>	<b>0.83</b>	<b>89.9</b>	<b>1.6</b>	<b>8.5</b>	<b>0.05</b>
<i>Rhizammina algaeformis</i>	GA	Ralga	MTub	GE	Bact	5.58	4.73	0.00	0.85	84.8	0	15.2	0.79
<i>Reophax apiculatus</i>	GA	Rapic	MU	GSI	Det	12.86	12.86	0	0	100	0	0	0.47
<i>Reophax bradyi</i>	<b>GA</b>	<b>Rbrad</b>	<b>MU</b>	<b>GSI</b>	<b>Det</b>	<b>11.35</b>	<b>7.24</b>	<b>0.42</b>	<b>3.69</b>	<b>63.80</b>	<b>3.70</b>	<b>32.50</b>	<b>0.05</b>
<i>Recurvoides contortus</i>	GA	Rcont	MTroc	GSI	Det	15.71	3.12	3.86	8.74	19.80	24.60	55.60	0.15
<i>Rhabdammina cylindrica</i>	GA	Rcyli	MTub	GE	Bact	12.09	12.09	0	0	100	0	0	0.50
<i>Reophax duplex</i>	GA	Rdupl	MU	GSI	Det	11.91	9.65	0.09	2.17	81.00	0.70	18.20	0.19
<i>Reusoolina apiculata</i>	GH	Reapi	MS	GSI	Det	13.57	13.57	0	0	100	0	0	0.38
<i>Recurvoidella bradyi</i>	GA	Recbrad	MP	GII	PhDet	10.83	10.83	0	0	100	0	0	0.67
<i>Reophax sp</i>	GA	Reoph	MU	GSI	Det	12.35	4.73	0.43	7.19	38.30	3.50	58.20	0.50
<i>Reophax fusiformis</i>	GA	Rfusi	MU	GSI	Det	9.61	6.04	0.53	3.05	62.80	5.50	31.70	0.31
<i>Rhabdammina sp.</i>	GA	Rhabda	MTub	GE	Bact	10.62	5.29	0.48	4.85	49.80	4.50	45.70	0.12
<i>Rhizammina sp</i>	GA	Rhiza	MTub	GE	Bact	9.37	7.40	0.08	1.89	79.00	0.90	20.20	0.13
<i>Rhumlerella humboldti</i>	GA	Rhumb	MT	GE	Det	32.47	29.35	0.68	2.45	90.40	2.10	7.50	0.09
<i>Reusoolina laevis</i>	<b>GH</b>	<b>Rlaev</b>	<b>MS</b>	<b>GSI</b>	<b>Det</b>	<b>17.84</b>	<b>10.48</b>	<b>1.42</b>	<b>5.94</b>	<b>58.70</b>	<b>8.00</b>	<b>33.30</b>	<b>0.05</b>
<i>Robertinoides oceanica</i>	GH	Rocea	MTroc	GSI	Car	15.88	15.88	0	0	100	0	0	0.33
<i>Rosalina globularis</i>	GH	Roglo	MTroc	GSI	Omni	12.90	12.90	0	0	100	0	0	0.48
<i>Rhabdammina pacifica</i>	GA	Rpaci	MTub	GE	Bact	11.22	7.94	0.7	2.58	70.8	6.3	23.0	0.06
<i>Reophax pilulifer</i>	GA	Rpilu	MU	GSI	Det	22.81	22.81	0	0	100	0	0	0.25
<i>Reophax pisiformis</i>	GA	Rpisi	MU	GSI	Det	22.81	22.81	0	0	100	0	0	0.15
<i>Rotalina pleurostomata</i>	GH	Rpleu	MTroc	GSI	Det	12.26	12.26	0	0	100	0	0	0.53
<i>Rutherfordoides rotundatus</i>	GH	Rrotu	MB	GSI	Det	4.53	4.53	0	0	100	0	0	0.97
<i>Reophax scorpiurus</i>	<b>GA</b>	<b>Rscor</b>	<b>MU</b>	<b>GSI</b>	<b>Det</b>	<b>13.80</b>	<b>3.27</b>	<b>2.89</b>	<b>7.64</b>	<b>23.70</b>	<b>20.90</b>	<b>55.40</b>	<b>0.01</b>
<i>Reophax subdentaliniformis</i>	GA	Rsubd	MU	GII	Det	8.00	8.00	0	0	100	0	0	0.83
<i>Reophax subfusiformis</i>	GA	Rsubf	MU	GSI	Det	13.57	13.57	0	0	100	0	0	0.36
<i>Recurvoides turbinatus</i>	GA	Rturb	MTroc	GII	Det	15.88	15.88	0	0	100	0	0	0.34

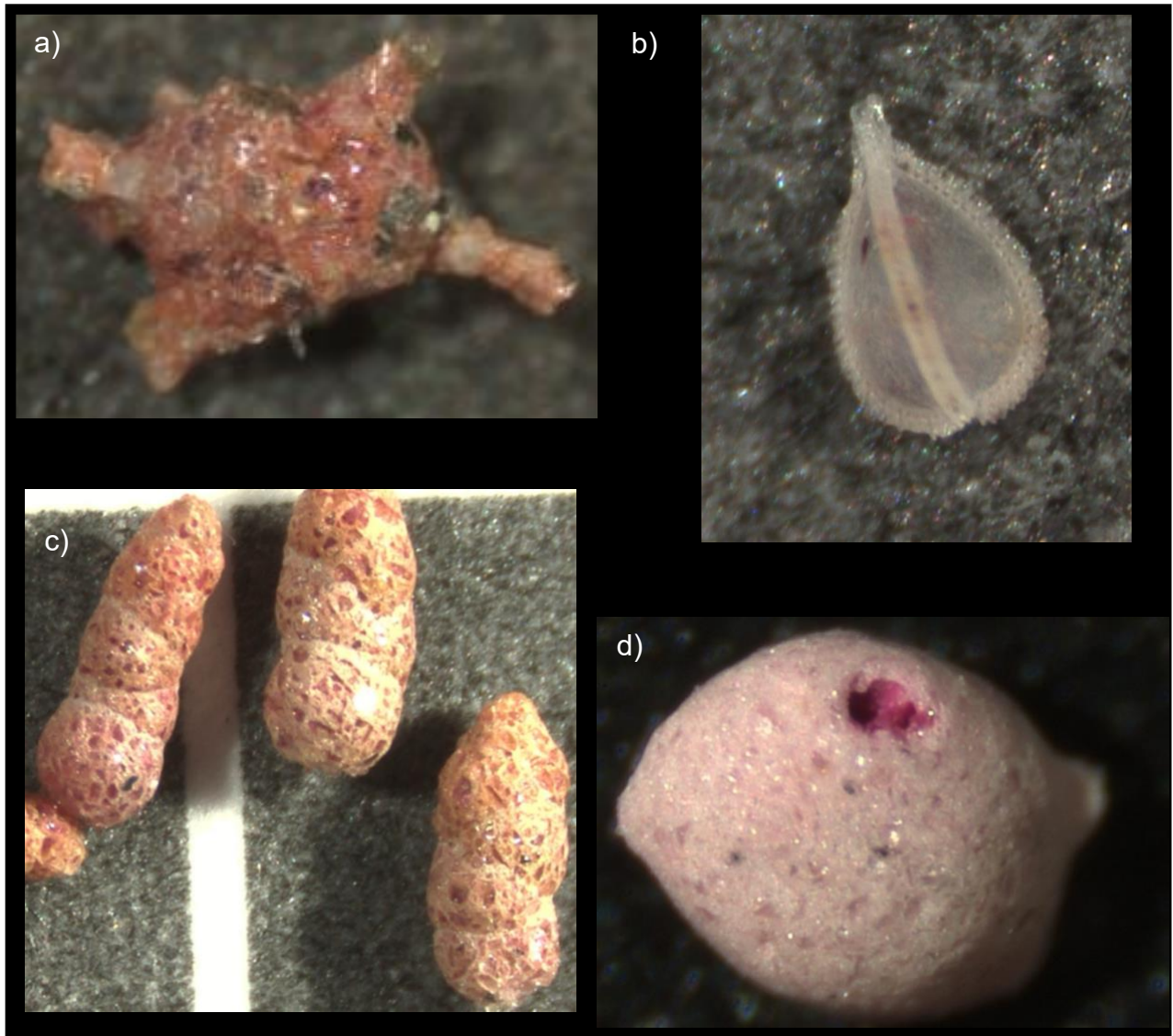
<i>Sphaeroidina bulloides</i>	GH	Sbull	MG	GSI	Det	9.39	5.26	0.58	3.55	56	6	38	0.09
<i>Sahulia conica</i>	GA	Sconi	MB	GII	Det	22.67	11.75	0.62	10.3	51.8	2.7	45.4	0.06
<i>Suggrunda eckisi</i>	GH	Secki	MB	GSI	Det	18.02	18.02	0	0	100	0	0	0.27
<b><i>Stainforthia fusiformis</i></b>	<b>GH</b>	<b>Sfusi</b>	<b>MB</b>	<b>GSI</b>	<b>Det</b>	<b>19.69</b>	<b>10.91</b>	<b>0.84</b>	<b>7.94</b>	<b>55.40</b>	<b>4.30</b>	<b>40.30</b>	<b>0.01</b>
<i>Siphouvigerina hispida</i>	GH	Shisp	MT	GSI	Det	12.26	12.26	0	0	100	0	0	0.53
<i>Suggrunda kleinpelli</i>	GH	Sklei	MB	GSI	Det	29.54	29.54	0	0	100	0	0	0.13
<i>Siphouvigerina proboscidea</i>	GH	Sprob	MT	GSI	Det	9.15	8.69	0.06	0.40	95	0.70	4.40	0.28
<b><i>Saccorhiza ramosa</i></b>	<b>GA</b>	<b>Sramo</b>	<b>MTub</b>	<b>GII</b>	<b>Bact</b>	<b>12.91</b>	<b>4.28</b>	<b>1.51</b>	<b>7.12</b>	<b>33.2</b>	<b>11.7</b>	<b>55.1</b>	<b>0.02</b>
<i>Saccamina sphaerica</i>	GA	Sspha	MS	GSI	Bact	7.94	3.05	1.14	3.74	38.5	14.4	47.1	0.60
<i>Saracenaria stolidota</i>	GH	Sstol	MP	GII	Det	12.86	12.86	0	0	100	0	0	0.46
<i>Spirosigmollina tenuis</i>	GP	Stenu	MM	GSI	Herb	7.71	7.71	0	0	100	0	0	0.87
<i>Thurammina albicans</i>	GA	Talbi	MS	GSI	Bact	22.81	22.81	0	0	100	0	0	0.18
<b><i>Trifarina angulosa</i></b>	<b>GH</b>	<b>Tangu</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>19.17</b>	<b>5.75</b>	<b>1.07</b>	<b>12.36</b>	<b>30.0</b>	<b>5.6</b>	<b>64.5</b>	<b>0.01</b>
<b><i>Takayanagia delicata</i></b>	<b>GH</b>	<b>Tdeli</b>	<b>MB</b>	<b>GII</b>	<b>Omni</b>	<b>16.29</b>	<b>6.82</b>	<b>2.9</b>	<b>6.57</b>	<b>41.9</b>	<b>17.8</b>	<b>40.3</b>	<b>0.02</b>
<i>Textularia earlandi</i>	GA	Tearl	MB	GII	Omni	13.42	13.42	0	0	100	0	0	0.47
<i>Trochammina inflata</i>	GA	Tinfl	MP	GSI	PhDet	12.20	8.31	1.8	2.09	68.1	14.8	17.1	0.43
<i>Techitella melo</i>	GA	Tmelo	MTub	GSI	Bact	11.18	9.97	0.28	0.93	89.2	2.5	8.3	0.25
<i>Trochammina nana</i>	GA	Tnana	MP	GSI	PhDet	7.71	7.71	0	0	100	0	0	0.86
<i>Textularia porrecta</i>	GA	Tporr	MB	GII	Omni	22.81	22.81	0	0	100	0	0	0.25
<i>Trochammina sp</i>	GA	Troch	MP	GSI	PhDet	7.71	7.71	0	0	100	0	0	0.80
<i>Techitella sp.</i>	GA	Tsp	MTub	GSI	Bact	15.05	8.11	2.86	4.08	53.9	19	27.1	0.31
<i>Trochammina squamata</i>	GA	Tsqua	MP	GSI	PhDet	22.16	4.57	4.19	13.39	20.6	18.9	60.5	0.22
<i>Triloculina tricarinata</i>	GP	Ttric	MM	GE	PhDet	8.63	8.15	0.04	0.45	94.4	0.4	5.2	0.44
<i>Triloculina trigonula</i>	GP	Ttrig	MM	GE	Herb	11.54	11.54	0	0	100	0	0	0.58
<i>Trochammina triloba</i>	GA	Ttril	MP	GE	PhDet	18.75	15.68	0.09	2.98	83.6	0.5	15.9	0.09
<i>Tolypammina vagans</i>	GA	Tvaga	MI	GSI	Bact	10.12	10.12	0	0	100	0	0	0.68
<b><i>Uvigerina auberiana</i></b>	<b>GH</b>	<b>Uaube</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>9.42</b>	<b>3.90</b>	<b>0.96</b>	<b>4.56</b>	<b>41.4</b>	<b>10.2</b>	<b>48.4</b>	<b>0.01</b>
<b><i>Uvigerina canariensis</i></b>	<b>GH</b>	<b>Ucana</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>16.16</b>	<b>13.38</b>	<b>0.12</b>	<b>2.66</b>	<b>82.8</b>	<b>0.8</b>	<b>16.5</b>	<b>0.03</b>
<i>Uvigerina peregrina</i>	GH	Upere	MT	GSI	Omni	12.34	0.50	2.70	9.15	4	21.8	74.1	0.09
<b><i>Uvigerina semiornata</i></b>	<b>GH</b>	<b>Usemi</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>10.85</b>	<b>9.84</b>	<b>0.12</b>	<b>0.89</b>	<b>90.7</b>	<b>1.1</b>	<b>8.2</b>	<b>0.05</b>
<i>Uvigerina senticosa</i>	GH	Usent	MT	GSI	Det	9.63	9.44	0.01	0.17	98.1	0.2	1.8	0.22
<i>Uvigerina sp</i>	GH	Usp	MT	GSI	Det	9.44	9.44	0	0	100	0	0	0.81
<b><i>Uvigerina striata</i></b>	<b>GH</b>	<b>Ustri</b>	<b>MT</b>	<b>GSI</b>	<b>Det</b>	<b>16.35</b>	<b>10.96</b>	<b>0.55</b>	<b>4.84</b>	<b>67.0</b>	<b>3.4</b>	<b>29.6</b>	<b>0.01</b>
<b><i>Vaginulina americana</i></b>	<b>GH</b>	<b>Vamer</b>	<b>MU</b>	<b>GSI</b>	<b>Omni</b>	<b>35.27</b>	<b>35.27</b>	<b>0</b>	<b>0</b>	<b>100</b>	<b>0</b>	<b>0</b>	<b>0.03</b>
<i>Valvulineria araucana</i>	GH	Varau	MTroc	GE	Det	11.54	11.54	0	0	100	0	0	0.60
<i>Vasicostella inflatiperforata</i>	GH	Vasin	MS	GSI	Det	14.46	14.46	0	0	100	0	0	0.41
<i>Valvulineria bradyana</i>	GH	Vbrad	MTroc	GE	Det	9.73	9.73	0	0	100	0	0	0.72

<i>Valvulineria glabra</i>	GH	Vglab	MTroc	GE	Det	14.06	1.15	4.4 1	8.51	8.2	31.3	60.5	0.44
<i>Valvulineria inaequalis</i>	GH	Vinae	MTroc	GE	Det	12.89	12.2 8	0.1 4	0.47	95.3	1.1	3.6	0.16
<i>Valvulineria inflata</i>	GH	Vinfl	MTroc	GE	Det	16.14	6.44	3.5 2	6.19	39.9	21.8	38.3	0.06
<i>Verneuilinulla propinqua</i>	GA	Vprop	MT	GII	Det	23.30	23.3 0	0	0	100	0	0	0.14
<i>Veloroninoides scitulus</i>	GA	Vscit	MP	GSI	Det	23.42	23.4 2	0	0	100	0	0	0.19

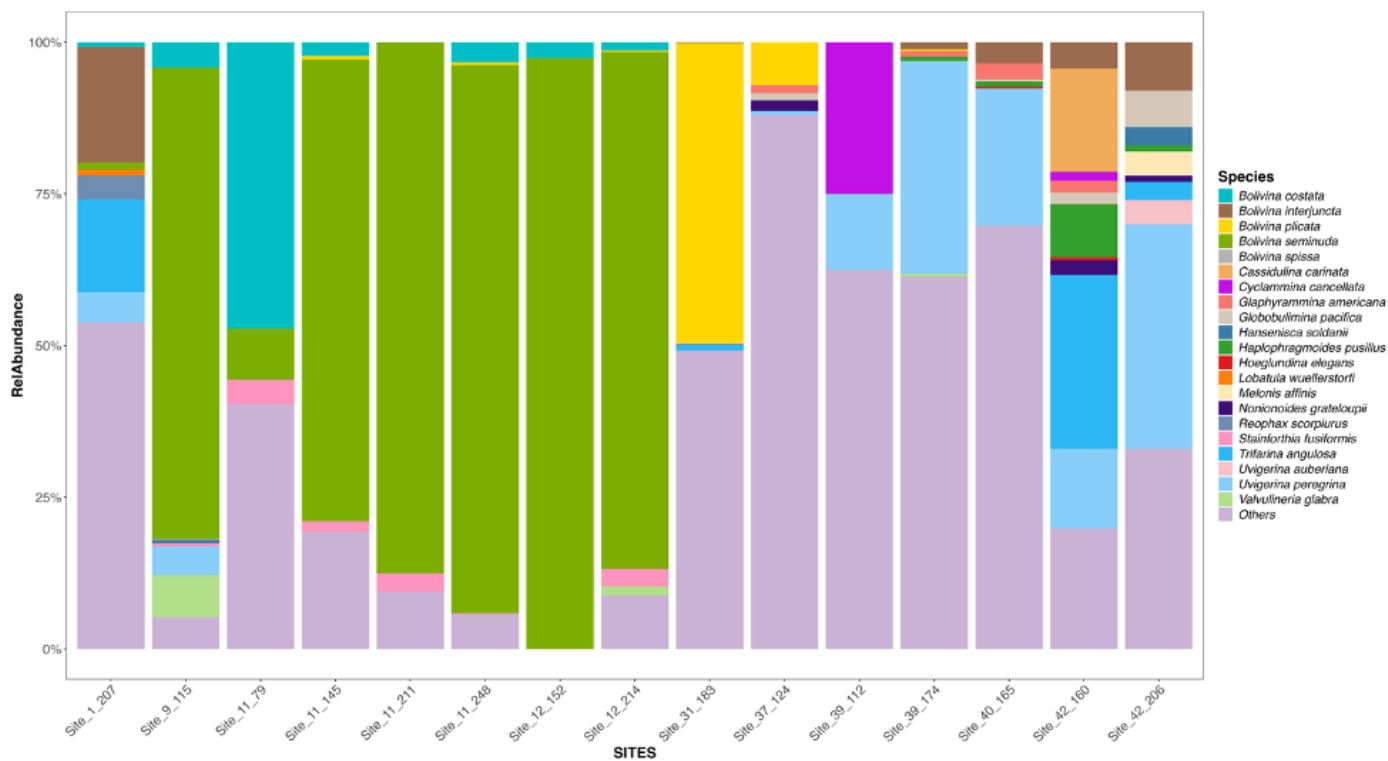
## Supplementary Figures



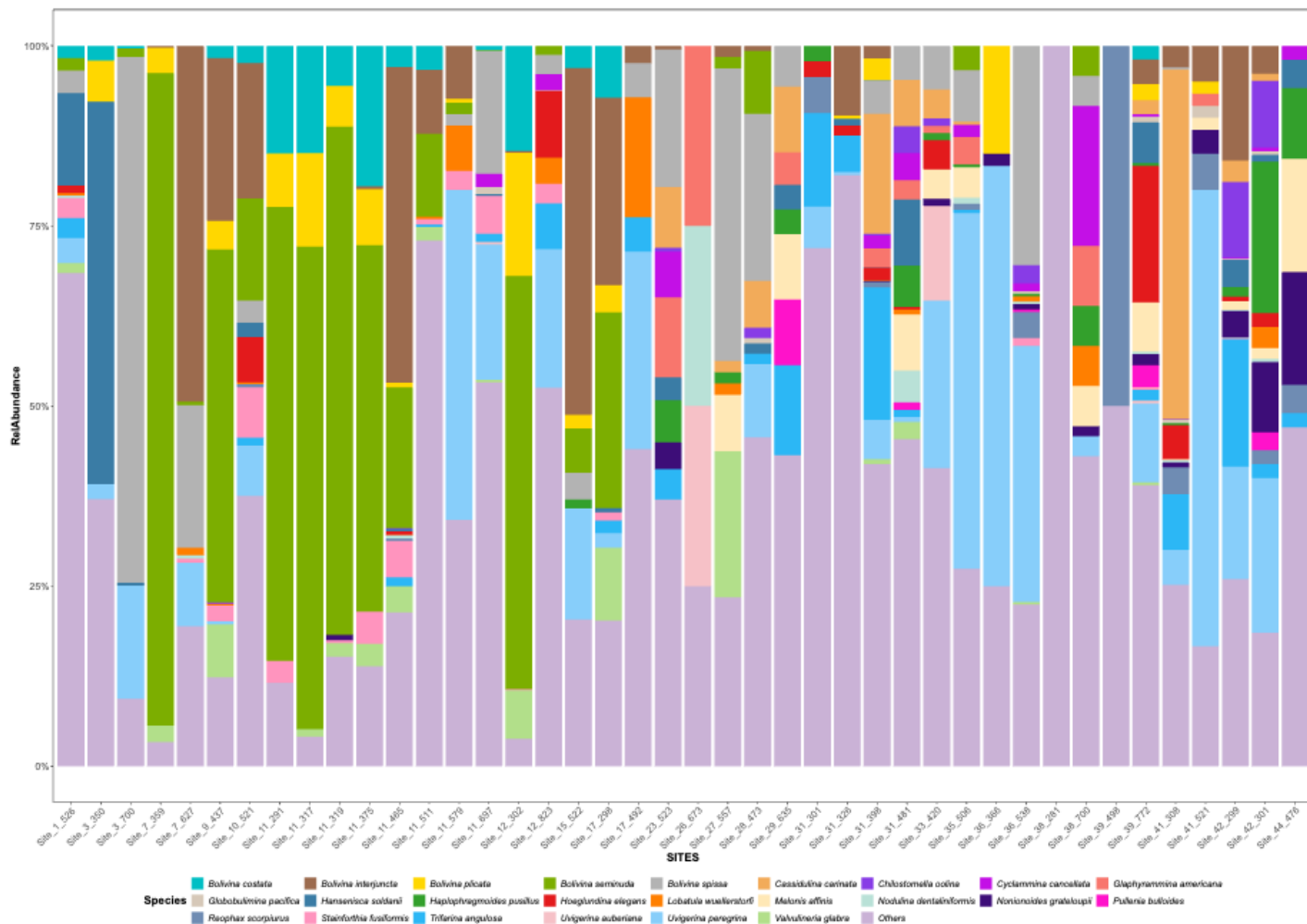
**Figure S1.** Distribution of bulk sediment composition across the Southeastern Pacific. **a)** Total organic carbon (TOC, wt %), **b)** ratio between organic carbon and total nitrogen content (C/N), **c)** calcium carbonate (CaCO<sub>3</sub>, wt %), **d)** biogenic opal (Opal, wt %), and nitrogen isotope ( $\delta^{15}\text{N}$ ). The symbols refer to the expeditions of M77, SO161, and RR-9702A.



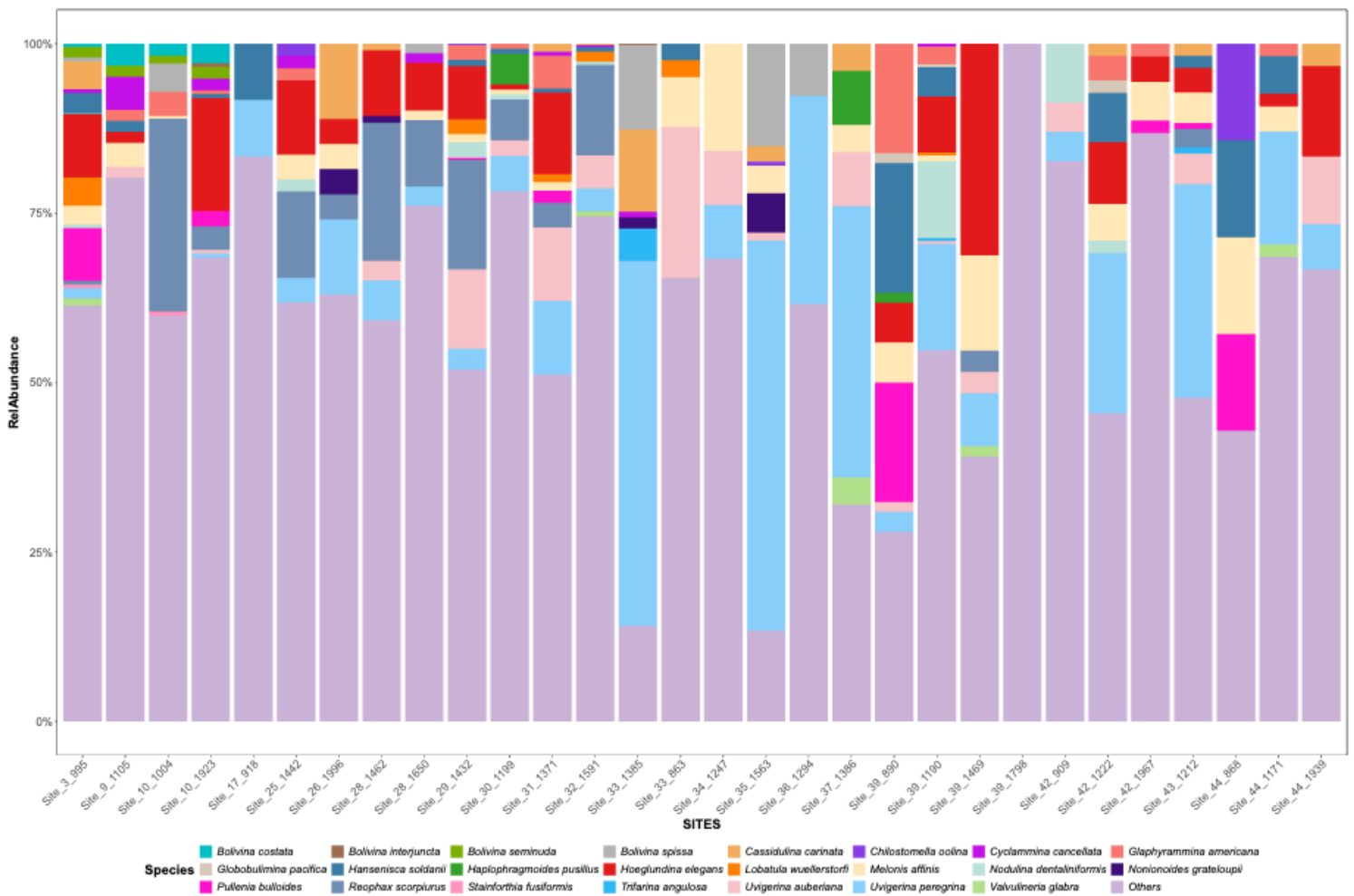
**Figure S2.** New records of living benthic foraminifera from the Southeastern Pacific. a) *Astrorhizoides cornutus*, b) *Galwayella trigonoornata*, c) *Liebusella goesi*, and d) *Crithionina pisum*.



**Figure S2.** Relative abundance of the species with more than 20% of contribution in all marine surface sediment samples < 250 m along the Southeastern Pacific. *Nodulina dentaliniformis* and *Pullenia bulloides* were removed from the graph because they have no presence in the sites.



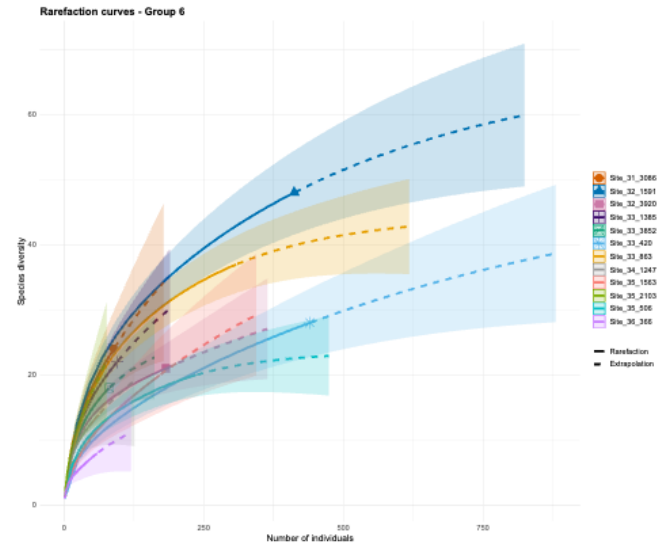
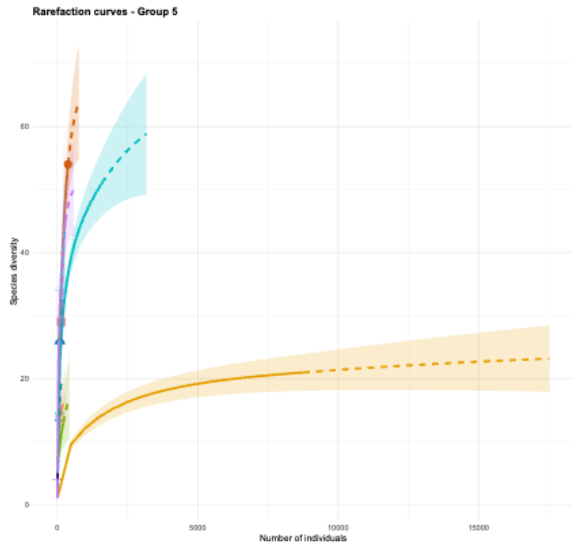
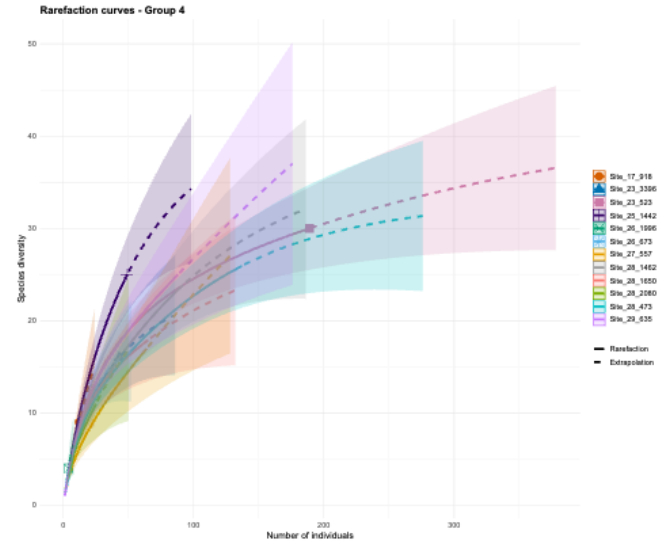
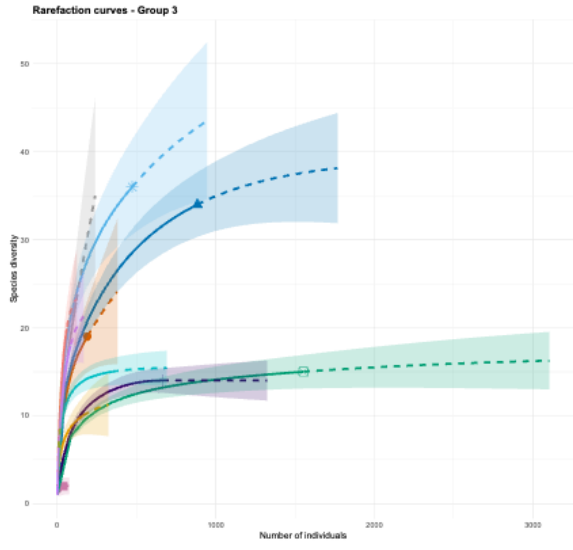
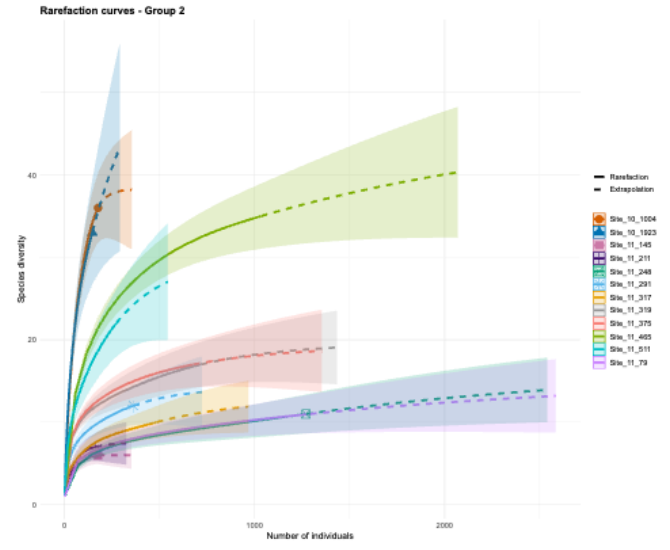
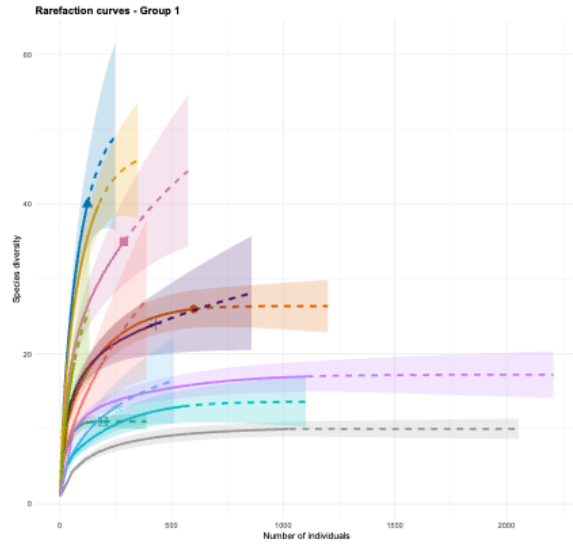
**Figure S3.** Relative abundance of the species with more than 20% of presence in the in all marine surface sediment samples >250 to 850 m along the Southeastern Pacific.



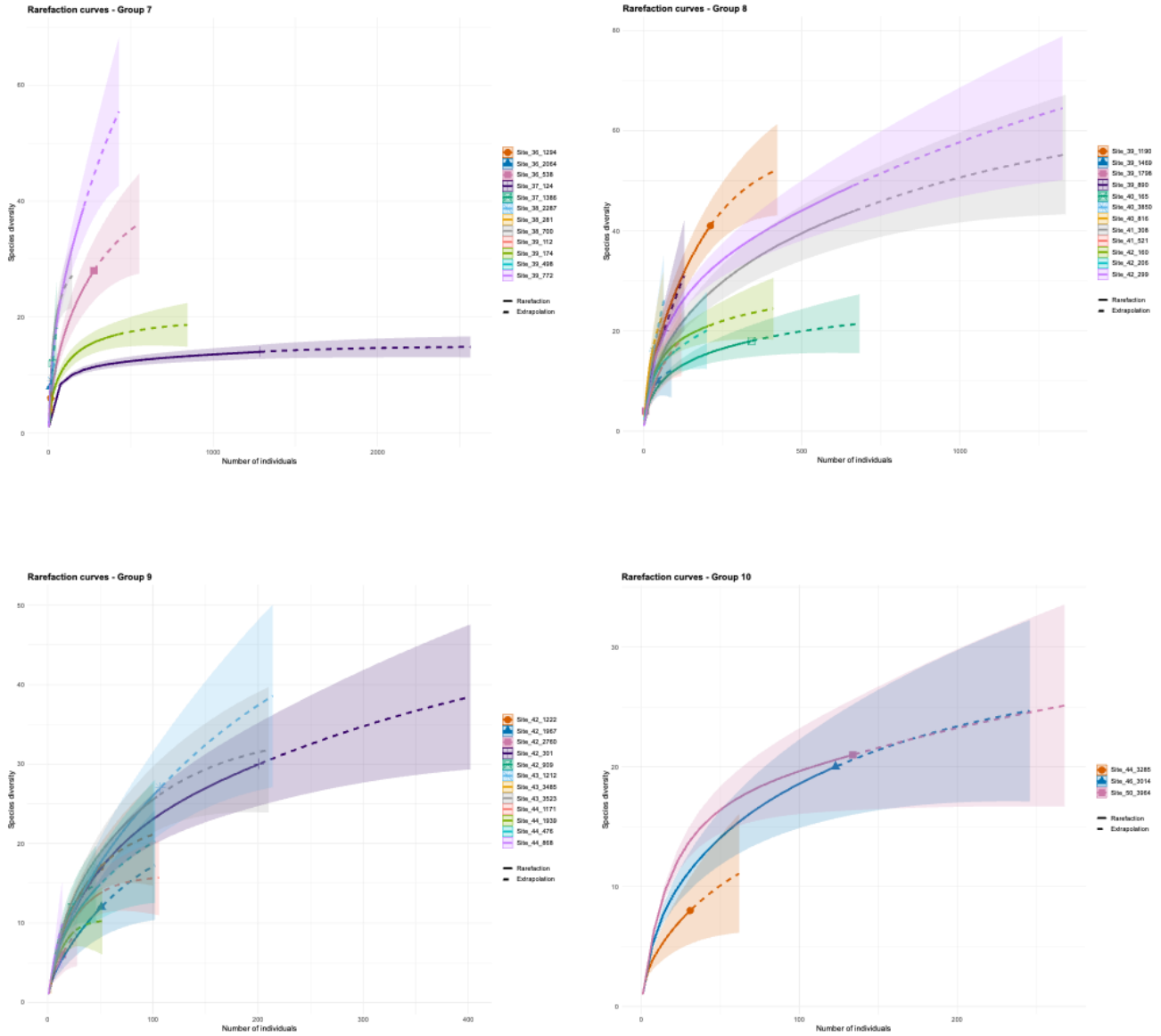
**Figure S4.** Relative abundance of the species with more than 20% of presence in the marine surface sediment samples >850 to 2000 m along the Southeastern Pacific. *Bolivina plicata* was removed from the graph due to its presence at any sample.



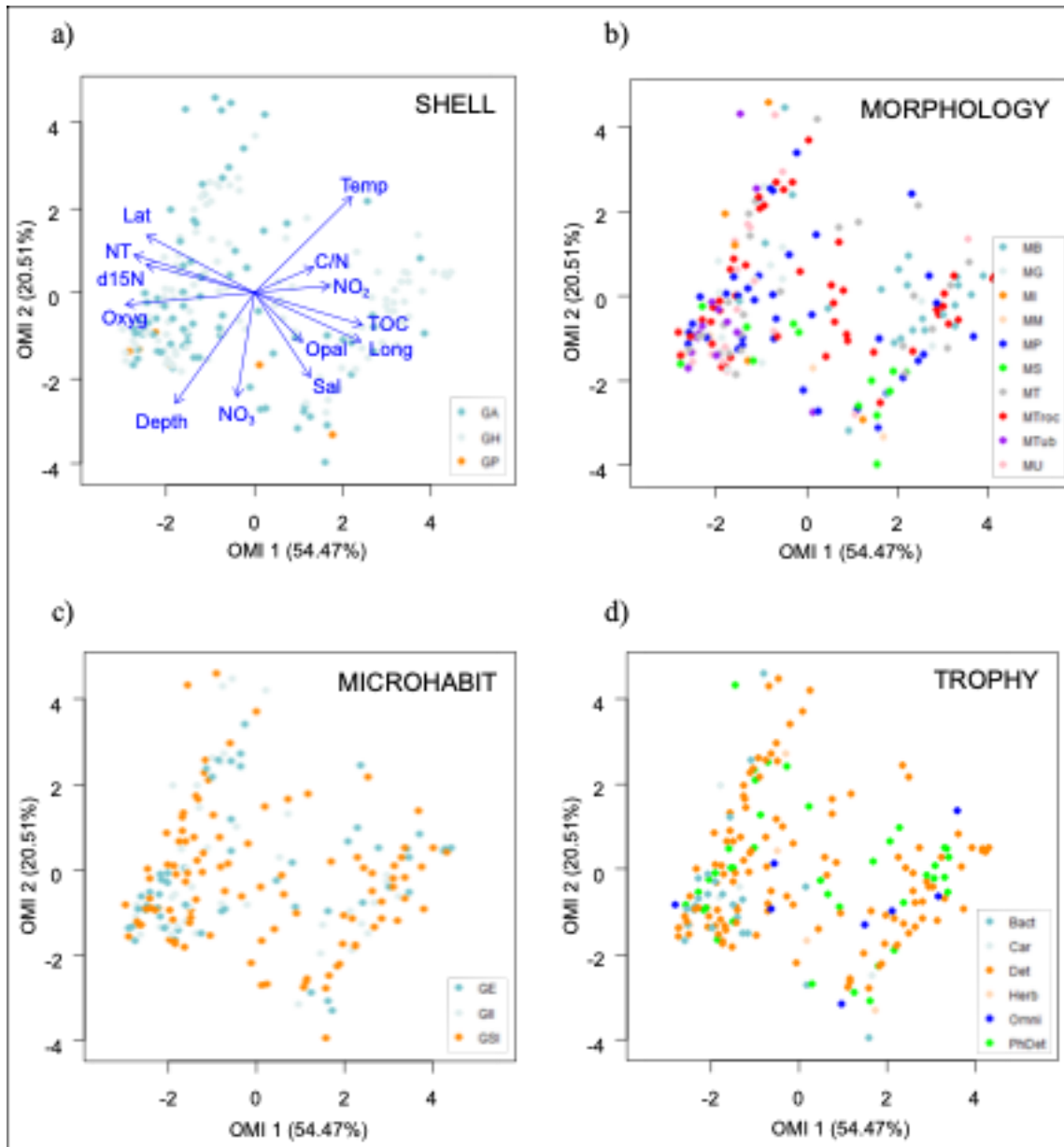
**Figure S5.** Relative abundance of the species with more than 20% of presence in the marine surface sediment samples > 2000 m along the Southeastern Pacific. *Bolivina costata*, *B. plicata*, *C. carinata*, *Haplophragmoides pusillus*, *Nonionoides grateloupii*, *Stainforthia fusiformis* and *Trifarina angulosa* were removed from the graph due they showed no presence in the samples.



**Figure S6.** Rarefaction curves of site groups along the western South American margin: a) 1° - 9°S. b) 10° - 11°S. c) 11° - 17°S. d) 17°S – 29°S. e) 29° - 31°S. f) 31° - 36°S. g) 36° - 39°S. h) 39° - 42°S. i) 42° - 44°S. j) 44° - 50°S.



**Figure S7.** Rarefaction curves of site groups along the western South American margin: a) 36° - 39°S. b) 39° - 42°S. c) 42° - 44°S. d) 44° - 50°S.



**Figure S8.** Outlying mean index (OMI) of the influence of the environmental variables on the functional traits of the benthic foraminifera species. **a)** Influence of the environmental variables on the type of test, GA: group agglutinated, GH: group Hyaline, GP: group porcellanaceous. **b)** Influence of the environmental variables on the morphology, MB: Morphology biserial, MG: Morphology globular, MI: Morphology irregular, MM: Morphology miliolide, MP: Morphology planispiral, MSp: Morphology spheric, unilocular and elongated spheric. MT: Morphology triserial, MTroc: Morphology trochospiral, MTub: Morphology tubular, MU: Morphology uniserial. **c)** Influence of the environmental

variables on the microhabitat, GE: group epifaunal, GII: Group intermedia infaunal and GSI: Group shallow infaunal. **d)** Influence of the environmental variables on the trophic behaviour, Bact: bacterivore, Car: carnivore, Det: detritivore, Herb: Herbivore, Omni: omnivore and PhDet: phytodetritivore. Letters and arrows in blue refer to the environmental variables, Temp: Temperature, Oxyg: oxygen, Sal: salinity, Lat: Latitude, Long: Longitude, NO<sub>3</sub>: Nitrate, NO<sub>2</sub>: Nitrite, TOC: Total Organic Carbon, NT: nitrogen total and d<sup>15</sup>N: <sup>15</sup>δN).