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# Bridging soil biogeochemistry and microbial communities (archaea and bacteria) in tropical seagrass meadows

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**Introduction:** Seagrass meadows are among the most valuable ecosystems, providing numerous ecosystem services and functions. Despite its importance, there is a lack of knowledge about soil's biogeochemical process variability, which can control microbiological communities. Thus, this study aimed to evaluate whether seagrass meadows in different geo-environments exhibit varying Fe and sulfate reduction intensities, shaping distinct archaea and bacteria communities.

**Methods:** Soil samples were collected in seagrass meadows under contrasting climatic, geological, vegetational and hydrological settings along the Brazilian coast (e.g., Semiarid Coast - SC, Southeastern Granitic Coast – GC, and Southern Quaternary Coast - QC). The soils were characterized by particle size, pH, redox potential (Eh), total organic C and total N content, acid-volatile sulfides (AVS), and simultaneously extracted Fe. Furthermore, a solid-phase Fe fractionation was performed to characterize the decomposition pathways in these soils, and the shifts in the microbial community along this spatial variation were analyzed using denaturing gradient gel electrophoresis.

**Results:** The studied soils presented a sandy texture (values ranging from 74  $\pm$  11.8 to 80.5  $\pm$  6.4%) caused by energetic hydrodynamic conditions. The pH values were circumneutral, while redox conditions presented significant distinction among the studied sites, ranging from anoxic to oxic (values ranging from -63 to +334 mV). The degree of pyritization (DOP) ranged from<

10% to values higher than 80%, highly influenced by rhizospheric oxidation, and higher AVS content was recorded for sites with lower DOP (i.e., GC and QC).

**Discussions:** Thus, biogeochemical processes in the seagrass soils present a wide variation in response to the geo-environmental settings. Plants influence the soil's geochemical and microbiological communities, retaining fine particles, promoting rhizosphere oxidation, and inducing anoxic conditions controlling the Fe and S forms. Moreover, the same plant species can result in distinct soil conditions and microbial communities due to geoenvironmental settings.

KEYWORDS

coastal wetlands, *Ruppia maritima*, *Halodule wrightii*, Fe fractionation, Metabolic pathways

#### **1** Introduction

Seagrass meadows consist of a worldwide distributed ecosystem, primarily composed of plant species that typically grow fully submerged in shallow coastal areas forming extensive meadows with high productivity and biodiversity (Olsen et al., 2016; Turschwell et al., 2021). The seagrasses are a phylogenetically related plant group (order Alismatales) with low taxonomic diversity (66 known species) assigned to six families (e.g., Cymodoceaceae, Hydrocharitaceae, Posidoniaceae, Zosteraceae; Ruppiaceae and Zannichelliaceae; Zidorn, 2016). These plant species interact with a wide range of megaherbivores (e.g., dugongs, manatees, green turtles) and smaller species (e.g., fishes, molluscs, crustaceans, sponges), supporting complex and diverse communities along the coastal zone of all continents except Antarctica (Nordlund et al., 2016; Unsworth et al., 2019).

This ecosystem can be found along the coast of 163 countries, covering 160,000 to 267,000 km<sup>2</sup> (McKenzie et al., 2020) and, thus, widespread over contrasting climatic, geological, vegetational and hydrological settings. However, estimations suggest that it may cover up to 600,000 km<sup>2</sup> (Fourqurean et al., 2012), corresponding to 0.2% of the ocean areas (Charette and Smith, 2010), which equals twice the area of mangroves forests (Siikamäki et al., 2013).

Seagrass meadows provide a wide range of ecological services, e.g., support marine food webs (Cui et al., 2021; Valentine and Heck, 2021); supply raw material and food (Connolly et al., 2005; Barbier et al., 2011); provide habitat and nursery (Orth et al., 2006; Liquete et al., 2016); water purification (Fernandes et al., 2009); coastal protection (Ondiviela et al., 2014); but also educational, recreational, spiritual and touristic services (de la Torre-Castro and Rönnbäck, 2004; Nordlund et al., 2016). As a result, seagrass meadows represent one of the most valuable ecosystems on the planet with values higher than U\$ 28,000.00 ha<sup>-1</sup> yr<sup>-1</sup> (2011 values; Barbier et al., 2011; Costanza et al., 2014).

Recently, numerous studies have been conducted to comprehend and highlight the role of seagrass meadows as globally significant carbon (C) sinks, storing significant amounts of C on their soils (Kennedy et al., 2010; Fourqurean et al., 2012; Rozaimi et al., 2016; Serrano et al., 2016; Ricart et al., 2020). The C storage in seagrass meadows results from high productivity (Duarte and Chiscano, 1999; Duarte et al., 2010), refractory plant tissues (Serrano et al., 2015; Kaal et al., 2016; Piñeiro-Juncal et al., 2020a), and the anoxic soil conditions, which may drive microbial communities and the anaerobic organic matter decomposition pathways of iron (Fe) and sulfate reduction (Fourqurean et al., 2012; Ugarelli et al., 2019; Sun et al., 2020).

In this ecosystem, due to the permanent flooding from seawater, with abundant sulfate supply, the organic matter decomposition occurs through Fe oxyhydroxides and sulfate reduction pathways since water column prevent oxygen diffusion through soil profile (Brodersen et al., 2017; Nóbrega et al., 2023). Consequently, Fe<sup>2+</sup> and sulfides (including HS- and polysulfides) are produced, which can eventually form iron sulfides such as FeS and FeS<sub>2</sub> (Otero and Macias, 2002; Queiroz et al., 2018). In anaerobic conditions, bioauthigenic pyrite (FeS<sub>2</sub>) emerges as the most stable product of sulfate reduction, holding diverse ecological significance (Otero et al., 2023).

In fact, the dynamics of Fe and sulfate in seagrass soils play an essential role in controlling several other processes that affect the very maintenance of this ecosystem, such as the bioavailability of nutrients (Delgard et al., 2016), trace metals (Tripathi et al., 2014), the fate of phosphorus (Brodersen et al., 2017), the geochemical control over dissolved sulfide (a highly phytotoxic compound for seagrass species (Azzoni et al., 2005; Holmer et al., 2005) and the C dynamics (Yu et al., 2021). Therefore, Fe and sulfate reduction, which are controlled by microbial processes (Brodersen et al., 2018; Piñeiro-Juncal et al., 2020b), play essential roles in the maintenance of different ecosystem services provided by seagrass meadows (e.g., carbon sequestration and pollutants immobilization) (Nordlund et al., 2016; York et al., 2017; Ramírez-Flandes et al., 2019; Ricart et al., 2020; Sun et al., 2020).

However, despite the ecological importance of Fe and sulfate biogeochemistry for seagrass meadows and other coastal wetlands (i.e., controlling pollutant, nutrient and C dynamics), there is still a global knowledge gap on its relationship with the microbial communities and how these groups are shaped in response to the

geo-environmental conditions (e.g., distinct climatic, geological conditions and vegetation), which may control the intensity of biogeochemical processes.

Because seagrass meadows colonize highly diverse environments (Gorman et al., 2020; Ricart et al., 2020; Nóbrega et al., 2023), the intensities of these geochemical processes may vary widely and shape different microbial communities. The present study tests the hypothesis that seagrass meadows at contrasting geoenvironments (e.g., climate, geology, and plant species and cover) will operate under different Fe and sulfate reduction intensities and, thus, will shape different microbiological communities. This is a novel study bridging the core soil processes of coastal wetlands (i.e., Fe and sulfate reduction) and microbial communities (bacteria and archaea) in diverse tropical geo-environments. Our results provide knowledge for a better understanding of the processes related to organic C accumulation, bioavailability of nutrients, and pollutant cycling in seagrass meadows.

#### 2 Material and methods

#### 2.1 Study site

Soil samples were collected in seagrass meadows located in three contrasting compartments along the 9,200 km of the Brazilian coast (Figure 1), presenting distinct climate (hot semiarid, humid tropical, and humid subtropical climates), geological setting (sedimentary and magmatic rocks), coastal conditions (estuary, beach, and coastal lagoon), and different plant species (Halodule wrightii. and Ruppia maritima) aiming to assess a maximum soil variability. Thus, seagrass soils from the semiarid coast (SC), granitic coast (GC) and quaternary coast (QC) compartments were samples and analyzed (Figure 1).

The semiarid coast (SC) extends from the states of Piauí to Pernambuco (Figure 1), marked by a semiarid climate (mainly classified tropical savanna and hot semi-arid climates; i.e., Aw and BSh climate types, respectively, according to Köppen-Geiger Climate Classification; Peel et al., 2007). The SC presents an annual rainfall (approximately 800 mm) lower than the evapotranspiration; and an average water temperature ranging from 27 to 29°C during the year (average minimum water temperature: 26°C and maximum: 30°C; seatemperature.org). In addition, this compartment is marked by close contact of the "Barreiras" group formation to the coastline, which results in sandy beaches and coral reefs close to the estuaries favouring seagrass beds formation (Copertino et al., 2016). The Barreiras group is a Miocene-Pliocene (Tertiary) transitional-coastal sedimentary deposit whose upper portion is composed dominantly of poorly sorted sand (Vilas Bôas et al., 2001; Arai, 2006). The seagrass meadow soils were collected in an estuary (at Ceará state, NE-Brazil) highly densely vegetated by H. wrightii Asch. (Figure 1); surrounded and close to (< 100 m) mangrove forests under low anthropogenic impacts due to the low occupation of the estuary (Silva and Souza, 2006), and found at water depths ranging from 1.7 to 2.9 m (water depth corrected to the mean water level).

The granitic coast (GC) covers the area from the Guanabara Bay (Rio de Janeiro state) to the Praia dos Sonhos (Santa Catarina state). In this compartment, narrow coastal plains are interrupted by the Serra do Mar's granitic-gneiss rock with small bays and beaches protected by rocky shores and almost 2,000 islands close to the shore (Schaeffer-Novelli et al., 1990). The humid tropical climate of the region (tropical rainforest climate - Af, according to Köppen-Geiger Climate Classification; Peel et al., 2007) presents an annual rainfall average higher than 1,600 mm during the entire year, with lower monthly precipitation higher than 60 mm, resulting in a hydric surplus, and the average monthly water temperature ranges from 21 and 27°C during the year (average minimum water temperature: 20°C and maximum: 28°C; seatemperature.org; Minuzzi et al., 2007). At this compartment, the representative H. wrightii. meadow was collected in an



FIGURE 1

Location of sampling sites along the Brazilian coast. In detail, plant species and canopy characteristics for the three studied sites. The x and y axes in the satellite images indicate the coordinates in UTM

urbanized beach area (at São Paulo state, Southeastern Brazil), presenting sparse vegetation with low plant density (Figure 1) at a water depth of 1.3 m (corrected to the mean water level).

The south quaternary coastal (QC) compartment extends 600 km from the Granitic coast to the border with Uruguay. This part of the coast presents a humid subtropical climate (Cfa according to Köppen-Geiger Climate Classification; Peel et al., 2007), with low annual temperature mean and a high-temperature oscillation (>10°C), annual precipitation of 1,300 mm, besides the hydric surplus (> 400 mm; Schaeffer-Novelli et al., 1990; Alvares et al., 2013). The average water temperature ranges from 13 to 24°C during the year (average minimum: 12°C and average maximum 26°C; seatemperature.org). The soil samples were collected at the *Lagoa dos Patos* (Patos Lagoon), at the *Rio Grande do Sul* State (S-Brazil), a 10,000 km<sup>2</sup> water body formed by sandy deposits during transgressive events during the Pleistocene (Toldo Jr. et al., 2000). The ephemeral and euryhaline *Ruppia maritima* seagrass meadow was collected at an average water depth of 1.5 m (corrected to the mean water level).

Seagrass meadows were observed and visually analyzed in each coast compartment to identify representative meadows (e.g., most frequent plant species, plant density and coverage), avoiding bare areas or seagrass meadows highly colonized with epiphytes or macroalgae.

# 2.2 Soil sampling procedure and physicochemical (Eh and pH) characterization

Three soil cores were randomly collected inside the representative seagrass meadows in each coast compartment using transparent polycarbonate tubes (60 mm diameter) attached to a submerged soil auger (Uwitec surface corer), which resulted in minimum sample compactation (Erich and Drohan, 2012). Soil samples were carefully extruded from the cores, partitioned into sections (0-3 cm, 3-6 cm, 6-10 cm, 10-13 cm, 13-16 cm, and 16-20 cm, with additional 5 cm increments) and, subsequently, pH and redox potential (Eh) values were promptly recorded to prevent oxidation. Following the characterization of physico-chemical conditions (Eh and pH), the samples were placed in plastic bags and transported to the laboratory under refrigeration (<4°C). In the lab, sub-samples (from the core's central region) were immediately frozen for subsequent analysis, while other sub-samples were dried (at 45°C until reaching a constant weight) and sieved through a 2 mm mesh.

The pH values were measured using a glass electrode calibrated using pH 4.0 and 7.0 buffer solution during core sectioning. The redox potential (Eh) was determined using a platinum electrode, with the final readings corrected by adding the potential of the calomel reference electrode (+ 244 mV) (Passos et al., 2016). All readings were taken after equilibrating the soil samples and the electrodes for several minutes (~2 min).

#### 2.3 Soil characterization

The soils from each seagrass meadow were characterized by particle size composition, total organic C, and total nitrogen.

The hydrometer method was used for the particle size analysis (Gee and Bauder, 1986) by a combination of physical (overnight shaking) and chemical (1 mol L<sup>-1</sup> NaOH + 0.015 mol L<sup>-1</sup> (NaPO<sub>3</sub>)<sub>6</sub>) dispersion after the organic matter elimination using hydrogen peroxide (30% vol). Total nitrogen (TN) and total organic carbon (TOC) were determined in an elemental analyzer (LECO 628 series). The TOC quantification was the difference between the total C and inorganic C contents (obtained after organic matter combustion in a muffle furnace at 550 °C for two hours; Howard et al., 2014).

#### 2.4 Fractionation of Fe and Sulfur

The partitioning of the solid phase of Fe was performed via a sequential extraction designed by the combination of methods proposed by Tessier et al. (1979); Huerta-Díaz and Morse (1990), and Fortin et al. (1993) using frozen samples, obtaining six operationally distinct fractions as follow: exchangeable Fe ( $Fe_{FX}$ ), extracted with 30 mL of MgCl<sub>2</sub> 1 mol L<sup>-1</sup> for 30 minutes; Fe associated with carbonates (Fe<sub>CA</sub>), extracted with 30 mL of 1 mol L <sup>1</sup> sodium acetate buffer solution (pH = 5.0) for 5 hours; ferrihydrite (Fe<sub>FR</sub>), extracted with 0.04 mol  $L^{-1}$  hydroxilamine + acetic acid 25% (v/v) at 30 °C for 6 hours; lepidocrocite (Fe<sub>IP</sub>), extracted with 0.04 mol  $L^{-1}$  hydroxilamine + acetic acid 25% (v/v) at 96 °C for 6 hours; crystalline Fe oxyhydroxides (Fe<sub>CR</sub>), extracted with 0.25 mol L<sup>-1</sup> sodium citrate + 0.11 mol L<sup>-1</sup> + 3 g of sodium dithionite for 30 minutes; and Fe associated with pyrite fraction (Fe<sub>PY</sub>), extracted with concentrated nitric acid for 2h, after the removal of silicate fraction (10 M HF during 16 hours, then added 5 grams of boric acid and agitated during 8 hours) and organic matter (concentrated H<sub>2</sub>SO<sub>4</sub> during 2 hours). For further details, please see Otero et al. (2009) and Osterrieth et al. (2016).

All solutions used for extraction were pretreatment for removing dissolved oxygen (i.e., purged with N<sub>2</sub> for 2 hours) to prevent possible Fe and S oxidation during the extraction steps. The extractants were collected between each step after centrifugation at 6,000 RPM (4 °C) for 30 minutes. The samples were washed with 20 mL of ultrapure water between each step and followed by centrifugation. The concentrations of Fe in each extract were quantified by flame atomic absorption spectrophotometry (Thermo Fisher Scientific, Waltham, MA, USA). Curve calibration solutions were prepared by diluting a certified standard solution (iron standard, TraceCERT®) to determine Fe concentration and a certified reference material (NIST SRM 2709a) was employed for quality control, with triplicate measurements yielding Fe concentration recovery values exceeding 92%. The results of the fractionation are presented as µmol g<sup>-1</sup> (micromole per gram) of dried soil, corrected by sample water content.

The degree of Fe pyritization (DOP), which determines the percentage of reactive Fe incorporated into the pyrite fraction, was calculated as follows: DOP (%) =  $100*Fe_{PY}/(Fe_{REACTIVE} + Fe_{PY})$  (Huerta-Diaz and Morse, 1992). The reactive fraction ( $Fe_{REACTIVE}$ ) was considered the sum of the fractions exchangeable to crystalline ( $\Sigma Fe_{EX} \rightarrow Fe_{CR}$ ), whereas the pseudo-total content was determined by the sum of the six fractions ( $\Sigma Fe_{EX} \rightarrow Fe_{PY}$ ).

The acid-volatile sulfides (AVS) fraction was extracted from frozen wet samples by acid distillation with HCl (20 mL, 3 mol L<sup>-1</sup>) in a gas-tight reaction flask after 40 minutes under continuous N<sub>2</sub> flow (Allen et al., 1993; Otero et al., 2009). The evolved H<sub>2</sub>S was trapped in a Zn acetate solution (3%). The concentration of sulfides was colourimetrically determined by the method proposed by Cline (1969), and the simultaneously extracted Fe (Fe-AVS) was quantified by flame atomic absorption spectrophotometry (Allen et al., 1993).

#### 2.5 Microbial molecular procedures

Frozen soil samples (400 mg) from different coastal compartments and depths (e.g., upper: 0-3 cm; intermediate: 16-20 cm; and deeper: 55-60 cm for SC and QC coast, and 35-40 cm for GC, due to soil depth limitation) were subjected to a total DNA extraction using the Power Soil DNA Isolation kit (MoBio Laboratories, Carlsbad, CA, USA) following the manufactory instructions. First, DNA extraction and integrity were assessed onto a 1.2% agarose gel electrophoresis with 1x TAE buffer (400 mmol L<sup>-1</sup> Tris, 20 mmol L<sup>-1</sup> acetic acid, and 1 mmol L<sup>-1</sup> EDTA). Subsequently, the gels were stained with GelRed<sup>TM</sup> (0.5  $\mu$ g mL<sup>-1</sup>), visualized, and photo-documented under ultraviolet light (DNR–Bio Imaging Systems/MiniBis Pro).

The amplification of the V6 region of ribosomal gene 16S rDNA for the bacterial community was performed with primers F968-GC and R1378, generating fragments of 410 bp (Heuer et al., 1997). The archaeal amplicons from rrs genes were obtained with primers Arch21F and Arch958R in the first reaction and Arch340FGC and Arch519R in the second nested-PCR reaction (Ovreås et al., 1997). The PCR reactions of bacterial communities were conducted in Veriti Thermal Cycler (Applied Biosystems, Waltham, USA), in a final volume of 50 µL, containing 1X Taq Buffer containing 2.50 mmol  $L^{-1}$  MgCl<sub>2</sub>, 0.25 mmol  $L^{-1}$  of each dNTP, 0.4 mmol  $L^{-1}$  of each primer, 1% formamide and 1U Taq DNA Polymerase (Fermentas, Burlington, Canada). For archaea communities, 1x Taq Buffer contained 3.0 mmol L<sup>-1</sup> MgCl<sub>2</sub>, 0.2 mmol L<sup>-1</sup> of each dNTP, 5 pmol L<sup>-1</sup> of each primer, and 1U Taq DNA polymerase (Fermentas, Burlington, Canada). The amplification reactions for bacterial DNA were performed in the following conditions: initial denaturation at 94°C and then 35 cycles of denaturation for 1 minute; annealing at 56°C for 1 min and elongation at 72°C for 2 min. At the end of 35 cycles, a final extension step was conducted at 72°C for 10 min. Amplification reactions for archaea communities were performed at 95°C for 5 minutes; and 30 cycles of 1 min at 95°C, 1 min at 55°C, 1 min at 72°C, and final extension for 10 min at 72°C.

The PCR products were loaded onto 6% (16S rRNA genes) and 8% (rrs genes) (w/v) polyacrylamide gels with denaturing gradients of 45 to 65% for bacterial and 30 to 55% for archaea community, using urea 7 mol  $L^{-1}$  and 40% deionized formamide as 100% denaturing solution. The gels were run for 16 h at 100 V at 60°C using the phorU2 systems (Ingeny International, Goes, Netherlands) and stained with SYBR Green I (Invitrogen, Breda, the Netherlands). The DGGE gels were photo-documented with Storm 845 (General Electric) and analyzed using the Image Quant

TL unidimensional (Amersham Biosciences, Amersham, UK, v.2003), where band patterns were converted into richness (number of bands) and abundance matrices.

#### 2.6 Statistical analysis

Differences between soil variables were analyzed by the nonparametric Kruskal-Wallis test, followed by the Bonferonni-Dunn method for multiple pairwise comparisons (XLSTAT, Addinsoft, New York, USA, 2014). Kruskal-Wallis text was used since it was not observed a normal distribution and equal variance. Since few assumptions are required for non-parametric analysis, this test is considered more robust for environmental data (Reimann et al., 2008). Pearson's correlation coefficient (R) established the relationships between different variables. Finally, a Discriminant Analysis (DA) was performed to develop a function that yields optimal discrimination of the study sites (SC, GC, and QC coasts). Through the DA, the relative contribution of the variables could be identified to the separation of the groups, i.e., the most relevant variables for separating the study sites (Reimann et al., 2008).

The bacterial and archaeal community structure changes among sites were obtained through beta-diversity estimations. The beta-diversity analysis compares the groups of microorganisms revealed by the DGGE band patterns among the samples, relying on the microorganisms' profile structure and relationship with the environmental soil parameters (Lozupone et al., 2007). For the beta-diversity calculation, similarity matrices were produced using the Bray-Curtis algorithm on square-root transformed data, allowing the observation of the distribution of the groups between the samples on a two-dimensional scale. PERMANOVA analyses, based on two factors (i.e., coastal compartment and depth of soil sample), were followed by a Monte Carlo method with fixed effects, sums of squares type III (partial) under a reduced model, and 999 permutations. The ANOSIM test followed a one-way analysis based on one factor allowing the comparison among treatments under 999 permutations based on a p-value<0.05. Finally, we performed a redundancy detrended analysis (RDA) regarding the environmental factors correlated to the bacterial distribution and sampling depth. The statistical tests regarding microbial communities were performed using the software PAST (Hammer et al., 2001), PRIMER 6+PERMANOVA (Clarke and Gorey, 2006), and Canoco (Canoco 4.5, Biometris, Wageningen, The Netherlands).

### **3** Results

# 3.1 Characteristics of the physicochemical environments

The particle size of soils from all studied sites presented a predominance of sand, with mean  $\pm$  standard deviation from 74.0  $\pm$  11.8% at the QC to 80.5  $\pm$  6.4% at the SC (Figure 2; Table 1). In contrast, silt content varied from 6.2  $\pm$  4.9% at SC to 12.2  $\pm$  5.4% at the QC coast, and clay varied from 8.1  $\pm$  4.1% at the GC to 14.1  $\pm$ 



2.3% at the SC coast. All three particle sizes presented a statistically significant difference (Table 1). Additionally, the soils from the three studied sites presented an erratic sand distribution throughout the soil profiles (Figure 2).

Regarding the physicochemical conditions, the pH values recorded for the three sites were circumneutral, with lower pH values recorded at the SC (mean value  $\pm$  standard deviation: 7.0  $\pm$  0.2), whereas GC (7.4  $\pm$  0.1) and QC (7.4  $\pm$  0.4) did not differ statistically (Table 1). Additionally, whereas the pH values of the GC coast soil did not present a significant variation through soil profiles, the soils from the SC coast present higher values at the deeper soil layers (> 10 cm), and the soils from the QC coast presented oscillating values, with a decrease at the intermediate soil layers (10-30 cm) followed by an increase at deeper layers (>30 cm) (Figure 3A).

The Eh values presented a wide variation, ranging from -63 to +334 mV, evidencing significantly contrasting conditions between the studied sites (Figure 3B; Table 1). Significantly lower Eh values were recorded for the SC (mean  $\pm$  standard deviation:  $+12 \pm 134$  mV), whereas the values recorded for QC ( $+166 \pm 136$  mV) and GC ( $+227 \pm 60$  mV) did not differ statistically. The redox contrasts were also evidenced in the surface soil horizons (down to 40 cm depth).

While the Eh values recorded at GC and QC evidenced an oxic environment (average mean for the surface soil horizons  $\pm$  standard deviation:  $\pm 209 \pm 113$  mV), significantly lower values were recorded at SC ( $-22 \pm 44$  mV), thus, evidencing an anoxic environment. As occurred to the pH, the Eh values presented different trends indepth among the different geoclimatic zones (Figure 3A). At the SC coast, the Eh values increased in-depth, whereas for both GC and QC coast, a decreasing trend in the recorded values (Figure 3B).

Significantly higher TOC contents were found at the SC coast (mean  $\pm$  standard deviation: 2.49  $\pm$  0.95%; Figure 3C) when compared to QC (0.53  $\pm$  0.17%) and GC (0.41  $\pm$  0.14%; Table 1). Similarly, significantly higher TN contents were recorded at the SC coast (0.14  $\pm$  0.11%; Figure 3D; Table 1), while the values recorded for QC (0.07  $\pm$  0.02%) and GC (0.05  $\pm$  0.01%) coast did not differ significantly (Figure 3D; Table 1). When considering the variation through the soil profile, the TOC and TN values recorded for the GC and QC coast presented a slight variation. On the other hand, for the SC coast, higher values were recorded for the upper (0-10 cm) compared to deeper (>35 cm) soil layers. The C:N ratios were also significantly different among the three sites, with lower values at QC (6.8  $\pm$  2.8), followed by GC (8.7  $\pm$  1.8) and SC (17.8  $\pm$  5.7; Figure 3C).

	p-value	Site				
Variable		Semiarid Coast (SC) (n=39)	Granitic Coast (GC) (n=30)	Quaternary Coast (QC) (n=39)		
Sand (%)	0.002	$79.7 \pm 5.1^{a}$	$80.5 \pm 6.4^{a}$	$74.0 \pm 11.8^{b}$		
Silt (%)	<0.01	$6.2 \pm 4.9^{\rm b}$	$11.4 \pm 5.3^{a}$	$12.2 \pm 5.4^{a}$		
Clay (%)	<0.01	$14.1 \pm 2.3^{a}$	$8.1\pm4.2^{\rm b}$	$13.8 \pm 6.8^{a}$		
рН	< 0.01	$7.0\pm0.2^{\mathrm{b}}$	$7.4 \pm 0.1^{a}$	$7.4 \pm 0.4^{a}$		
Eh (mV)	< 0.01	$+12 \pm 134^{b}$	$+227 \pm 60^{a}$	$+166 \pm 136^{a}$		
TOC (%)	< 0.01	$2.49 \pm 0.95^{a}$	$0.41 \pm 0.14^{\rm b}$	$0.53 \pm 0.17^{\rm b}$		
TN (%)	< 0.01	$0.14 \pm 0.11^{a}$	$0.05 \pm 0.01^{\rm b}$	$0.07\pm0.02^{\rm b}$		
C:N ratio	< 0.01	$17.8 \pm 5.7^{a}$	$8.7 \pm 1.8^{\rm b}$	$6.8 \pm 2.8^{\rm b}$		
Fe <sub>T</sub> (µmol g <sup>-1</sup> )	0.37	$78.1 \pm 29.6^{a}$	$73.3 \pm 19.9^{a}$	$63.7 \pm 26.1^{a}$		
Fe <sub>EX</sub> (µmol g <sup>-1</sup> )	< 0.01	$1.7 \pm 0.3^{a}$	$0.7\pm0.0^{ m b}$	$0.4 \pm 0.1^{\circ}$		
Fe <sub>CA</sub> (µmol g <sup>-1</sup> )	0.43	$0.8 \pm 0.1^{a}$	$1.0 \pm 0.3^{a}$	$0.6 \pm 1.2^{a}$		
Fe <sub>FR</sub> (µmol g <sup>-1</sup> )	< 0.01	$4.2 \pm 1.5^{\mathrm{b}}$	$19.4 \pm 15.4^{a}$	$2.6 \pm 2.5^{\rm b}$		
Fe <sub>LP</sub> (µmol g <sup>-1</sup> )	< 0.01	$14.7 \pm 5.2^{\rm b}$	$24.3 \pm 2.7^{a}$	$10.0 \pm 2.1^{\circ}$		
Fe <sub>CR</sub> (µmol g <sup>-1</sup> )	< 0.01	$4.8\pm1.4^{\rm b}$	$10.4 \pm 1.9^{a}$	$5.1 \pm 1.6^{b}$		
Fe <sub>OXIDES</sub> (µmol g <sup>-1</sup> )	< 0.01	$23.7 \pm 7.9^{\rm b}$	$54.0 \pm 16.0^{a}$	$17.7 \pm 4.4^{b}$		
Fe <sub>PY</sub> (µmol g <sup>-1</sup> )	< 0.01	$51.9 \pm 21.8^{a}$	$17.7 \pm 9.5^{\rm b}$	$45.0 \pm 26.5^{a}$		
DOP (%)	< 0.01	$62.8 \pm 4.0^{a}$	$23.3 \pm 10.6^{b}$	$58.8 \pm 27.4^{a}$		
AVS (µmol g <sup>-1</sup> )	0.04	$0.11 \pm 0.09^{b}$	$0.25 \pm 0.22^{a}$	$0.22 \pm 0.36$ <sup>a</sup>		
Fe-AVS (µmol g <sup>-1</sup> )	< 0.01	$11.1 \pm 7.5^{b}$	$30.2 \pm 5.7^{a}a$	$7.0 \pm 3.2^{b}$		

Mean values followed by the same letter did not present a statistically significant difference.

#### 3.2 Fe fractionation

The values of pseudo-total Fe (Fe<sub>T</sub>) contents were statistically similar among the three sites (means  $\pm$  standard deviation for the SC:78.1  $\pm$  29.6 µmol g<sup>-1</sup>; GC: 73.3  $\pm$  19.9 µmol g<sup>-1</sup>; and QC: 63.7  $\pm$  26.1 µmol g<sup>-1</sup>; Figure 4; Table 1).

On the other hand, statistically significant differences were recorded for all Fe fractions (Figure 4; Table 1). At QC and SC, Fe<sub>PY</sub> was the most important fraction (mean ± standard deviation:  $51.9 \pm 21.8 \ \mu\text{mol g}^{-1}$  for SC coast and  $45.0 \pm 26.5 \ \mu\text{mol g}^{-1}$  for QC coast) representing more than 50% of the Fe<sub>T</sub> (DOP > 50%) whereas, for the GC, Fe<sub>PY</sub> represent less than 50% of the Fe<sub>T</sub> (DOP < 50%; Figure 4), with significantly lower contents ( $17.7 \pm 9.5 \ \mu\text{mol g}^{-1}$ ). Additionally, the DOP values presented a significant variation throughout the soil for GC and QC, with a lower DOP at the upper soil layers (0-10 cm; Figure 4). In contrast, the DOP at the SC did not oscillate with soil depth (Figure 4).

Both Fe<sub>EX</sub> and Fe<sub>CA</sub> were the fraction with a lower contribution to Fe<sub>T</sub>, with mean  $\pm$  standard deviation values of Fe<sub>EX</sub> contents oscillating from 0.4  $\pm$  0.1 to 1.7  $\pm$  0.3 µmol g<sup>-1</sup> for QC and SC coast, respectively; and mean values of Fe<sub>CA</sub> varying from 0.6  $\pm$  1.2 and 1.0  $\pm$  0.3 µmol g<sup>-1</sup> for the QC and GC coast, respectively (Figure 4).

Regarding the Fe oxides fraction (i.e.,  $Fe_{FR} + Fe_{LP} + Fe_{CR}$ ), significantly higher contents were quantified at GC (mean ± standard deviation: 54.0 ± 16.0 µmol g<sup>-1</sup>; Table 1), followed by SC (23.7 ± 7.9 µmol g<sup>-1</sup>), and QC coast (17.7 ± 9.5 µmol g<sup>-1</sup>). For the three sites, the lepidocrocite (Fe<sub>LP</sub>) was the most dominant oxyhydroxide fraction, with mean ( ± standard deviation) values ranging from 10.0 ± 2.1 to 24.3 ± 2.7 µmol g<sup>-1</sup> at the QC and GC coasts (Figure 4; Table 1). Moreover, for the GC, the Fe<sub>CR</sub> fraction was the less relevant oxyhydroxide fraction (10.4 ± 1.9 µmol g<sup>-1</sup>), whereas the ferrihydrite (Fe<sub>FR</sub>) was the less important Fe oxyhydroxide fraction for the other sites (4.2 ± 1.5 µmol g<sup>-1</sup> for NE and 2.6 ± 2.5 µmol g<sup>-1</sup> for QC coast; Figure 4).

# 3.3 Acid volatile sulfides (AVS) and simultaneously extracted Fe (AVS-Fe)

The AVS contents ranged from 0.05 to 2.03  $\mu$ mol g<sup>-1</sup> (Figure 3F) among the studied sites. Significantly, lower AVS contents were recorded at the SC (mean ± standard deviation: 0.11 ± 0.09  $\mu$ mol g<sup>-1</sup>), followed by QC (0.22 ± 0.36  $\mu$ mol g<sup>-1</sup>) and GC (0.25 ± 0.22  $\mu$ mol g<sup>-1</sup>; Figure 3F).



Regarding the AVS-Fe, statistically significant higher values were recorded for the GC (30.2  $\pm$  5.7  $\mu$ mol g<sup>-1</sup>), whereas the values recorded for SC (5.6  $\pm$  1.5  $\mu$ mol g<sup>-1</sup>) and QC (7.0  $\pm$  3.3  $\mu$ mol g<sup>-1</sup>) did not differ statistically (Figure 3G; Table 1).

#### 3.4 Discriminant analysis of soils variables

The discriminant analysis indicated the most relevant variables for the differentiation of the studied sites (Figure 5). For the soils from SC, the vectors position indicates an association with higher TOC, TN,  $Fe_{EX}$  contents, higher C:N ratio values, and lower pH, Eh, and AVS contents. The GC soils were associated with higher Fe oxyhydroxides ( $Fe_{FR}$ ,  $Fe_{LP}$ ,  $Fe_{CR}$ , and  $Fe_{OXIDES}$ ) and Fe-AVS as observed by the vectors positioned in its direction and lower DOP,  $Fe_{PY}$ , and clay contents as evidenced by the vectors positioned in the opposite direction (Figure 5). On the other hand, the vector's position indicates that soils from QC differ from the SC and GC by lower sand and  $Fe_{EX}$  contents and higher silt and pH (Figure 5). Furthermore, when comparing SC and QC, it can be stated that SC presents higher values of  $Fe_{PY}$  and DOP.

# 3.5 Bacteria and archaea community characteristics (PCR-DGGE Results)

PCR-DGGE results evidence a clear differentiation in the bacteria and *archaea* community structure among the coastal compartments and soil depths, shown by the PERMANOVA data analysis (Table 2). In fact, the coastal compartments played a higher influence in the distribution of the *archaea* (Pseudo-F = 15.657; p = 0.001) and bacteria (Pseudo-F = 8.097; p = 0.001) communities; followed by the influence of soil depth (Pseudo-F = 4.395 and 3.519 for *archaea* and bacteria communities, respectively; p = 0.001). In addition, the results demonstrated the integration of factors (coastal compartment vs. depth) in the distribution of the bacteria (Pseudo-F = 3.745; p = 0.001) and archaea (Pseudo-F = 6.294; p = 0.001) communities (Table 2).

Moreover, the separation of the *archaea* community is also demonstrated by the ANOSIM-test (coastal compartment: Global R = 0.947; p = 0,001 and soil depth: Global R = 0.786; p = 0.001). Regarding the bacterial community, the ANOSIM-test demonstrated a clear separation among the sites (Coastal compartment: Global R = 0.967; p = 0,001). However, some



profile overlap was found regarding the soil depth (Global R = 0.58; p = 0.001), indicating some similarity among the bacterial group sampled in distinct soil depths.

According to the redundancy analysis, the soils from SC archaea communities were positively associated with TOC,  $Fe_T$ , TN, DOP, and clay, whereas the GC were associated with  $Fe_{OXIDES}$  and sand (Figure 6A). The upper horizons of the QC archaea communities

were positively associated with higher Eh and pH, whereas the intermediate and deeper soil layers were negatively associated with  $Fe_{OXIDES}$  (Figure 6A). The intermediate and deeper soil layers from GC were, comparatively, more associated with the deeper layers from SC soils. The GC bacterial communities were positively associated with  $Fe_{OXIDES}$ , whereas QC communities were negatively associated with  $Fe_{OXIDES}$ , (Figure 6B). The SC



#### FIGURE 5

Discriminant analysis results of seagrass meadows' soils variables in the three studied geo-environment (i.e., SC, GC, and QC) along the Brazilian coast. TOC, Total organic carbon contents; TN, total nitrogen contents; AVS, acid volatile sulfides; AVS-Fe, Fe associated with AVS fraction; FeEX, soluble and exchangeable Fe; FeCA, Fe bounded to carbonates; FeFR, Fe associated with ferrihydrite; FeLP, Fe associated with lepidocrocite; FeCR, Fe associated with crystalline Fe oxyhydroxides; FePY, Fe associated with pyrite; Fe<sub>OXIDES</sub>, the sum of FeFR; FeLP; and FeCR; and DOP, degree of Fe pyritization. TABLE 2 Permanova analysis of the in a Monte Carlo test arrangement with 999 permutations, under a p-value MC< 0.05.

Factors	Permanova table						
Factors	df F test Permutations		Permutations	p-value (MC)			
Archaea							
Study Sites (SS)	2	15.657	999	0.001			
Soil Depth (SD)	3	4.395	997	0.002			
ES x SD	3	6.294	999	0.001			
Bacteria							
Study Sites (SS)	2	8.097	997	0.001			
Soil Depth (SD)	3	3.519	998	0.001			
SS x SD	3	3.745	999	0.001			

The results evaluated the significance of the role of each factor and its combinations on the variation of the microbial community structure.

communities were positively associated with TOC, TN,  $Fe_T$ , DOP, and clay and negatively associated with Eh and pH (Figure 6B).

### 4 Discussion

# 4.1 Physicochemical diversity of seagrass meadows and its effects on soil biogeochemical processes

Diverse studies have shown that soils of coastal wetlands (e.g., mangroves, salt marshes, and hypersaline tidal flats) can be very dynamic and spatially variable, resulting from an interaction between biotic (e.g., plant species and bioturbation; Araújo Júnior et al., 2012; Cabral et al., 2020), and abiotic factors (i.e., physiographic position, flooding frequency, geological bedrocks, and climate; Seybold et al., 2002; Du Laing et al., 2009; Albuquerque et al., 2014; Ferronato et al., 2016; Ferreira et al., 2022).

Our findings indicate that the distinct geo-environments and observed vegetation characteristics (e.g., plant biomass, density, and species) resulted in differentiated soil physico-chemical conditions at the studied sites (Figures 2 and 3; Table 1). Overall, the predominance of sand in the three studied sites reflects a high hydrodynamics energy which influences the particle size distribution in the studied sites. A more energetic hydrodynamic condition favours fine particle transportation and coarse particle accumulation (Flemming, 2000; Pejrup, 1988). Accordingly, it can be stated that a more energetic condition was recorded for soils from GC, where the seagrass meadows were in a beach area with coarse soil textures, and from estuarine soils at SC. Conversely, QC seagrass meadows are located in a closed estuary, i.e., a choked lagoon (Figure 1; Toldo Jr. et al., 2000; Lanari and Copertino, 2017), and thus present characteristics of a slightly lower energetic environment according to Flemming (2000) (i.e., higher contents of fine particles such as silt and clay; Figure 2; Table 1).



Results of redundancy detrended analysis regarding the environmental factors correlated to (A) archaea and (B) bacteria distribution and depth. TOC: Total organic carbon contents, TN: total nitrogen contents, Fe<sub>OXIDES</sub>: the sum of FeFR, FeLP, and FeCR, Fe<sub>T</sub>: pseudo total Fe, and DOP: degree of Fe pyritization. On the other hand, SC soils present finer soil texture (mean clay content:  $14.1 \pm 2.3\%$ ; Table 1), mainly at the soil surface, compared to GC soils. These characteristics can be attributed to the increased plant biomass within the meadow in SC, which has developed over coarser soils (Figure 1), which attenuates the effects of turbulence and wave action, promoting sedimentation of finer suspended particles and decreasing its resuspension in the upper soil layers (Gacia et al., 1999; Ondiviela et al., 2014). On the other hand, specifically for the QC samples, the higher sand contents on the upper soil layers (down to 10 cm depth) may result from a recently enhanced erosive event that may have delivered coarse particles to the lagoon (Toldo Jr. et al., 2000).

The contrasting geo-environments reflect significantly distinct redox conditions (Figure 3). For example, at the SC coast, anoxic conditions (Eh< +100 mV; Otero et al., 2009) were predominant, whereas sub-oxic conditions (+100 mV< Eh< +350 mV; Otero et al., 2009) predominate at GC and QC, despite anoxic conditions recorded at deeper layers (deeper than 30 cm) of the QC (Figure 3). Furthermore, despite the SC and GC seagrass meadows presenting similar plant species (*H. wrightii*; Figure 1), the redox conditions recorded for the GC were closely related to the QC (vegetated by *Ruppia maritima*; Figure 3). This fact highlights that seagrass plant species do not solely drive redox conditions, since same plant species may present distinct redox conditions. Additionally, compared to seagrass meadows from other sites (Table 3), the redox potentials at GC and QC were consistently higher, indicating the importance of abiotic factors (e.g., particle size distribution, hydrodynamic) for seagrass meadows' soils characteristics.

In this sense, the different plant biomass among sites (Figure 1) and the potentially higher input of labile organic matter from the surrounding mangroves and seagrass meadows in the SC soil produced lower Eh values. This explanation corroborates the opposite association observed in the discriminant analysis (Figure 5) and the significant negative correlation between TOC and Eh (Figure 7A). As labile organic matter is produced, exuded, or trapped by seagrasses vegetation, there is an intensification of the microbial respiration processes (Kaldy, 2012; Barrón et al., 2014), depleting the dissolved oxygen concentrations, leading to the consumption of other electron acceptors substituting  $O_2$  and resulting in lower Eh values (Canfield et al., 1993; Miller et al., 2007).

TABLE 3	Physicochemical	(Eh and pH), sand,	and total organic carbon (	TOC) contents quantified fo	r different seagrass meadows are	ound the globe.
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Site	Seagrass specie	Eh	рН	Sand	ТОС	Reference	
		(mV)		(%)			
Global data					2.5 ± 0.1	Fourqurean et al., 2012	
Moreton Bay Australia	Zostera muelleri* and others			$100 \pm 0$	$0.1 \pm 0.1$	Samper-Villarreal et al.,	
woreton Day, Australia	Zostera muelleri <sup>+</sup> and others			91 ± 6	$0.5 \pm 0.4$	2016	
Rhode Island, USA	Z. marina		7.3 to 8.5	14 to 97	0.2-30	Bradley and Stolt, 2003	
Cockburn Sound, Western Australia	Posidonia sinuosa				0.4 - 1.3	Serrano et al., 2015	
Port Curtis, Australia	Z. muelleri				1.5 - 2.0		
Clinchaum Mahaus Diand Danish Baltis	Z. marina				$0.2 \pm 0.1$		
Supshavn, Nyborg Fjord, Danish Banic					$2.1 \pm 0.1$		
Sint Annaland, Eastern Scheld, The Netherland	Z. marina				$0.7 \pm 0.1$	Ricart et al., 2015	
Anna Iran Dara Darana	Z. marina				0.3 ± 0.01		
Arcachon bay, France	Z. noltii				1.8 ± 0.3		
Palk Bay, India		-52 to +56	7.2 to 8.2	39-90		Thangaradjou et al., 2015	
Awerange Bay, Indonesia	Halodule uninervis* and others	-300 to 0	6.4 to 7.2		$1.2 \pm 0.1$ to $4.3 \pm 0.8$	Alongi et al., 2008	
Tanahara da Pataran Maratan Arata la	Ruppia megacarpa	-120 to +49		54 to 62	3.8 to 4.9	Kilminster, 2013	
Leschenault Estuary, western Austrana	Halophila ovalis	+40 to +70		42 to 61	2.5 to 3.7		
Fangan bay, Australia	Z. muelleri			72 to 93	1.7 ± 0.8	Macreadie et al., 2014	
Blanes Bay, Spain		-74 to +396				Marbà and Duarte, 2001	

On the other hand, it must be considered the role of seagrass plants diffusing photosynthetic-generated O<sub>2</sub> to protect the basal meristem from the damages arising from sulfides (Brodersen et al., 2015). This fact would explain the higher Eh values found in the uppermost soil layers from the GC and QC (Figure 3B). These roots have an oxidative effect resulting from an effective photosynthetic system (Olsen et al., 2016), and its well-developed aerenchyma enables rapid gas diffusion to roots, rhizomes, basal meristem, and soil (Koren et al., 2015). However, the oxidation of the seagrass rhizosphere can be diminished by an intense soil microbial metabolism, which rapidly consumes the diffused oxygen. As extensive mangrove forest borders the seagrass meadows at the SC sampling site, an intensification of the labile organic matter input (dissolved and particulate; Bouillon et al., 2007; Maher et al., 2013), stimulating microbial respiration, may have prevented the establishment of sub-oxic and oxic conditions at the upper-most soil layers as occurred at GC and QC coasts (Figure 3B).

The circumneutral pH values recorded for the studied sites (Figure 3A) are also related to the reductive processes of Fe and sulfate reduction in these soils. Both geochemical processes consume  $H^+$  or produce alkalinity (Equation 1 and 2; Otero et al., 2009). Conversely, the lower pH values at the upper layers of SC (0-10 cm depth) may result from intense rhizospheric activity since the higher roots' density may result in higher organic acid exudation to increase nutrients bioavailability, especially phosphorus (Long et al., 2008).

 $4FeOOH + CH_2O + 7H^+ \rightarrow 4Fe^{2+} + HCO_3 - +6H_2O$ 

$$SO_4^{2^-} + 2CH_2O \rightarrow HS^- + H^+ + 2HCO_3^-$$
 (2)

The soils' organic C and N contents were significantly higher on the SC, presumably derived from the higher plant biomass and the input from surrounding mangroves (Figure 1). The higher C:N ratio recorded for the SC coast (Figure 3) supports a hypothesis for C inputs from mangrove forests bordering the seagrass (Kennedy et al., 2010; Garcias-Bonet et al., 2019). Furthermore, there is a significant correlation between TOC and TN for all studied sites (Figure 7B), also represented by the strong association of the TOC and TN by the discriminant analysis results (Figure 5).

This association indicates that the soil organic matter is the source of N in these areas and is possibly attributed to the N fixation and assimilation processes promoted by sulfate-reducing bacteria (Welsh, 2000; Mohr et al., 2021). Additionally, the higher TOC content in the SC results from the less energetic metabolic pathway, corroborated by the lower Eh recorded in this site (Figure 3B). Thus, the intensity of anaerobic microbial metabolism associated with primary productivity may vary and is responsible for the C accumulation in seagrass meadow soils. In this case, the lower metabolic yield (e.g., SC) promotes soil organic matter accumulation, generally with low humification and unsaturation degree but enriched in H, N, and alcoholic and methoxyl groups (Neue et al., 1997; Kristensen et al., 2008).

Compared to seagrass meadows located in other parts of the world, the TOC contents were lower than the global average (average: 2.5%, median 1.8%; Fourqurean et al., 2012; Armitage and Fourqurean, 2016) but comparable to other sites around the

(1)

12

globe (Table 3). However, a marked variability was observed between the studied sites, probably associated with the interaction of multiple environmental factors (e.g., seagrass species, landscape configuration, and soil redox conditions; Ricart et al., 2015; Ricart et al., 2020). However, our study evidence that geo-environmental factors controlling microbial metabolisms may be more relevant for C accumulation than plant species since the meadows formed by the same species present significant TOC contents (e.g., SC and GC). Additionally, TOC contents were a significant variable for differentiating microbial communities among studied sites (Figure 7). Thus, more studies must be conducted to improve comprehension of the spatial variability of seagrass meadows, improving the accuracy of the C stored in these ecosystems (Ricart et al., 2020), and how the effects of organic C control microbial communities.

# 4.2 Fe and S dynamic and metabolic pathways

Despite the differentiated geological setting of the studied sites, with the influence of granitic-gneiss rocks at the GC (Valeriano et al., 2016) and the predominance of quaternary sandy deposits at the SC and QC (Toldo Jr. et al., 2000; Vilas Bôas et al., 2001), the Fe<sub>T</sub> contents did not differ statistically (Table 1). However, the Fe fractionating presented significant differences among the sites. Accordingly, the intensity of Fe and S metabolic pathways contrasted among the studied sites.

The seagrass meadow located at the SC differed from other sites by the significantly higher DOP values, which remained above 60% (Figure 4). Under the anoxic conditions at the SC (Table 1), microbiologically mediated Fe<sup>3+</sup> reduction decreased Fe oxyhydroxide contents while increasing the soluble Fe<sup>2+</sup> and Fe associated with sulfides (e.g., pyrite). On the other hand, at GC, the higher Eh values indicate an unfavourable environment for Fe reduction (i.e., oxic conditions) and the maintenance of Fe in oxyhydroxide forms (Figure 4). In such environments with redox oscillations between suboxic to oxic and circumneutral pH conditions (e.g., GC; Table 1), soluble Fe<sup>2+</sup> is oxidized to Fe<sup>3+</sup> and precipitated as oxyhydroxides (Ferreira et al., 2021; Ferreira et al., 2022) For the QC, fairly high DOP values occurred (>60%, especially at soil layers deeper than 10 cm; Figure 4) despite the high redox potentials (suboxic environment). In this case, the presence of pyrite could result from former (or transient) environmental conditions favourable to pyrite formation (Ferreira et al., 2015). Remnant pyrites may be protected by Fe oxyhydroxide coatings, preventing complete pyrite oxidation (Huminicki and Rimstidt, 2009). Furthermore, the presence of poorly crystalline Fe oxyhydroxides, especially FeLP, at the upper layers of GC and QC could be considered a result of the oxidation of sulfides by the rhizosphere (Azzoni et al., 2001; Brodersen et al., 2015) since lepidocrocite is the main product of Fe oxidation in saline environments (Equation 3; Otero et al., 2009).

$$\text{FeS}_2 + 3.5\text{O}_2 + \text{H}_2\text{O} \rightarrow \text{Fe}(\text{OH})_3 + 2\text{SO}_4^{2-} + 4\text{H}^+$$
 (3)

The higher content of crystalline Fe oxyhydroxides (i.e.,  $Fe_{CR}$ ) at the GC could also explain the lower DOP registered at this site since crystalline Fe forms are less prone to reduction due to their mineralogical features such as lower surface area, reactivity, and crystal structure (i.e., rhombohedral and hexagonal or irregular plates) when compared to poorly crystalline Fe oxyhydroxides (Cornell and Schwertmann, 2003). Conversely, the higher susceptibility of poorly crystalline Fe oxyhydroxides to microbial Fe reduction could also explain the negative correlation between DOP vs.  $Fe_{FR}$  and vs.  $Fe_{LP}$  (Figures 5, 7C, D).

Moreover, the AVS and AVS-Fe contents also varied among the studied geo-environments (Figures 3F, G), corroborating its effects on Fe and S dynamics. These fractions correspond to a complex mixture of aqueous H<sub>2</sub>S and its dissociated form, Fe-S complexes, polysulfides, and solid-phase mackinawite and greigite (Rickard et al., 2017), with significant ecological importance once it may control trace metals dynamics and bioavailability (MaChado et al., 2010). Thus, the higher AVS content in the GC compared to SC and QC reinforces the different intensities of Fe and S metabolic pathways among the studied sites. Furthermore, due to its phytotoxicity, high sulfide production has been associated with declining seagrass meadows(Holmer et al., 2005). Accordingly, the higher AVS content at the GC, especially on surface soil layers, may be related to the lower plant density at this site (Figure 1), especially compared to the SC vegetated by the same plant species, since lower plant density and lower organic C content prevents Fe oxyhydroxides reduction (and lower DOP).

Regarding microbial communities, we observed a positive correlation with higher  $Fe_{OXIDES}$  and sand contents, higher Eh values, and negatively correlated with TOC at GC site (Figure 7). This fact indicates that the low plant density and high Fe inputs from the Fe-rich geological surroundings affected the microbial structure. A recent study has reported that forming distinct Fecycling bacterial genera (e.g., Lewinella and Woeseia) communities in seagrass meadows positively correlates with greater Fe contents in its soils (Martin et al., 2022). On the other hand, previous studies have reported that sandy sediments (e.g., GC) with high sulfide production favour the dominance of archaea communities (Llobet-Brossa et al., 2002; Buhring et al., 2005). Specifically, high sulfide production increases genes related to sulfate reduction in a *Posidonia* spp. ecosystem (Fraser et al., 2023).

The higher plant cover in SC possibly changes microbial activity in the seagrass rhizosphere (Jankowska et al., 2015) since we observed a clear cluster in microbial communities, mainly driven by clay content, TOC, TN, and DOP (Figure 6). Accordingly, previous studies have reported that gross primary production in tropical seagrass bacteria communities increases the amount of C available for sulfate-reducing bacteria metabolism (Pollard and Moriarty, 1991; Williams et al., 2009). Moreover, Zhang et al. (2022) revealed that higher clay contents play an important role in seagrass soil biogeochemistry. For example, these authors reported that seagrass soils with a predominance of fine particles have a higher capacity for root exudate retention, increasing its concentration around the roots and enhancing the abundance of sulfate-reducing bacteria (see DOP values in Table 1). Interestingly, in a *Thalassia hemprichii* system it was demonstrated that induced carbon addition changed the bacterial community, mainly those functional groups related to N cycle (Zhang et al., 2021). A higher TOC content in SC could increase N uptake and, consequently, boost photosynthetic performance of seagrass and modified rhizosphere microbial communities.

Changes in seagrass microbial communities seem to be contextdependent since microbial groups changed according to geographical location. More importantly, the same plant species (*H. wrightii.* at SC and GC) shapes different microbial communities in the soil (Figure 6). Thus, probably seagrass microbial communities change according to plant type, but microbial rhizosphere core strongly depends on soil geochemical properties (Cúcio et al., 2016). For example, Vogel et al. (2021) demonstrated that the main drivers of microbial community structure in *Thalassia testudinum* seagrass were local geo-environmental conditions.

Therefore, our findings reveal that seagrass meadows soils at contrasting geo-environments (e.g., climate, geology, and plant species and cover) present different intensities of Fe and sulfate reduction, ultimately shaping different microbiological communities. More importantly, we provide evidence that bacteria and archaea communities responded similarly to geoenvironmental conditions, indicating that both could be equally important to microbial activity and nutrient cycling in seagrass ecosystems. These results shed light on the knowledge gap about the intensities of geochemical processes involving Fe and S dynamic and metabolic pathways in seagrass meadows' soils.

#### **5** Conclusions

In this study, we identify that seagrass meadows are a heterogeneous ecosystem in which the variability of soil composition affects the intensities of geochemical processes due to distinct geo-environments (e.g., climate, geology, salinity, plant species and coverage, and anthropic influence). For example, in soils under higher plant cover that enhances the C contents and geology that favors higher Fe contents, there are higher contents of fine particles (e.g., clay), an abundance of sulfate-reducing bacteria, and microbial-mediated processes involving Fe and S. This environment favors metabolic pathways with low oxidation rates (e.g., sulfate reduction), Fe sulfides formation, and ultimately, highcapacity C sequestration. On the other hand, in seagrass meadow soils with higher contents of sand and low plant density, a more oxidizing environment is established that favors Fe oxyhydroxides and AVS formation. This contrasting environment presents a lower capacity for organic C accumulation and contaminants immobilization. In addition, the differentiated geo-environmental conditions markedly influenced microbial communities, even when vegetated with the same plant species. Our findings revealed novel insights into the heterogeneity of seagrass meadow soils and a better understanding of the biogeochemical processes inherent to these ecosystems.

### Data availability statement

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

### Author contributions

GN and TF: Funding acquisition, Investigation, Formal analysis, Conceptualization, Methodology, Validation, Writing original draft, review and editing. PA, HQ, AP, MC, DG, WZ, JS, FA, XO: Investigation, Formal analysis, Data validation, Writing original draft and editing. All authors contributed to the article and approved the submitted version.

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### Conflict of interest

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

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