



Article Diversity and Variation of Epiphytic Diatoms on *Ruppia maritima* L., Related to Anthropogenic Impact in an Estuary in Southern Brazil

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Abstract: Knowledge about the diversity and spatiotemporal variability of epiphytic diatom communities in estuarine meadows has great relevance for coastal ecology and, thus, contributes to understanding the impact of natural and anthropogenic changes on seagrass meadows. The community of epiphytic diatoms in Ruppia maritima L. meadows was investigated in two environments with different levels of anthropogenic impact and nutrient loads. Both impacted and non-impacted meadows had similar conditions in terms of water depth, temperature and transparency but distinct nutrient loads and salinity ranges. A total of 159 diatom taxa were found on Ruppia maritima leaves during the monitoring period, including freshwater (30.8%), marine (25.1%), brackish (9.4%) and cosmopolitan (8.8%) taxa. The most abundant species were C. placentula, T. tabulata, M. pumila and T. fasciculata, in addition to A. tenuissimus, C. adhaerens and M. moniliformis. Although present in both sites, C. placentula and T. tabulata were the dominant species in the impacted site. We found that 32% of the taxa were exclusive to the non-impacted site, 23% to the impacted site and 45% were common to both sites. The study sites showed marked differences in community attributes; i.e., higher richness, diversity and equitability and lower dominance were found in the non-impacted site, which is distant from anthropogenic sources of domestic and industrial sewage and has low concentrations of dissolved N and P in water and low values of sediment organic matter. Nutrient concentration and salinity were the main factors behind the spatial and temporal variability in the structure of the epiphytic community when all other environmental variables were similar (water depth, temperature, transparency and host plant). The influence of temperature and salinity on community structure was site-dependent. This study revealed the high richness and diversity of epiphytic diatoms in the meadows of the Patos Lagoon estuary (PLE) and the high spatial and temporal heterogeneity of the communities, and it shows the potential of epiphytic community studies for the assessment of environmental quality in seagrass meadow habitats.

Keywords: coastal lagoon; diatoms; diversity; eutrophication; submerged aquatic vegetation; spatial variability

1. Introduction

Epiphytic algae enhance seagrass-meadow primary productivity, biodiversity and structural complexity, sustaining a significant part of estuarine and marine food webs [1]. Epiphytes protect seagrasses from UV-B radiation and desiccation [2,3], are used as food by invertebrates and small fishes [4–6] and can be more productive than their macroscopic seagrass hosts [7].

In addition, epiphytic algae may contribute up to 50% of the carbon buried in seagrass soils [8]. In the face of natural and human-induced perturbations, changes in seagrass system biodiversity can influence resource use and productivity, alter trophic interactions and reduce the stability of estuarine and coastal systems [9]. Therefore, understanding the composition and abundance of epiphytic algae and how they vary according to abiotic



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). variables can contribute to the understanding of marine and coastal biodiversity but also to predictions of the impact of natural and anthropogenic changes on seagrass meadows.

Due to the faster growth rates and biomass turnover compared to their seagrass hosts and the specific community responses to changes in environmental conditions, epiphytic algae are excellent indicators of water quality, the integrity of seagrass meadows and the overall aquatic ecosystem [10,11]. The degradation of aquatic ecosystems causes changes in the structures of epiphytic algal communities, as well as subsequent reductions in taxonomic diversity and proliferations of toxic and opportunistic species [12,13]. However, few researchers have studied how epiphytic communities in general, and diatoms in particular, respond to changes in water quality, such as nutrient enhancement [14].

Among the epiphytic algae, diatoms are the dominant or the most abundant components in shallow estuarine areas [15,16]. The abundance and composition of diatom species are dependent on abiotic factors (nutrients, light availability, current velocity, wave action, temperature, salinity), biotic factors (especially herbivory action) [17–19] and intrinsic factors, such as motility [20].

The species of epiphytic diatoms on the same seagrass species may differ among different habitats [21]. For example, variability in spatial and temporal factors, such as salinity, nutrient availability and temperature, has been suggested to structure the epiphytic diatom community of *Thalassia testudinum* in Florida Bay, USA [22]. Another study [23] found that epiphyte composition was influenced mainly by environmental factors, such as wave energy, temperature, ammonium concentration and salinity compared to site (ocean *versus* bay) and host species. On the other hand, salinity has been shown to be the most important factor for diatom community dissimilarities between sites for both seagrasses [24] and macroalgal hosts [25,26]. Therefore, variations in the physicochemical characteristics of water have a significant influence on the epiphytic diatom community.

Despite their ecological relevance and their potential for water-quality monitoring, seagrass epiphytic algae are neglected in many areas where seagrass meadows occur mainly around reefs and within coastal lagoons; additionally, their conservation status is critical [13]. The few records of seagrass meadow epiphytes concern filamentous macroalgae on *R. maritima* [27] and *Halodule wrightii* [28], resulting in epiphytic diatoms remaining poorly studied [28,29].

Patos Lagoon in southern Brazil is a dominant feature of the coastal plain, connected to a large continental drainage basin (200.000 km²) and the Atlantic Ocean. The primary producers in the Patos Lagoon estuary (PLE) sustain a highly productive food web [30]. The PLE holds approximately 120 km² of shoals and shallow bays, which are under different levels of natural and anthropogenic impacts [31,32]. The distribution and abundance of seagrass meadows in Patos Lagoon have diminished since the 1990s due to a complex combination of natural and anthropogenic changes [33], with significant effects on the structure of benthic macrofauna, the abundance of resident estuarine fish [34], trophic links [35] and the fate of organic matter [36]. In the PLE, the composition, density and biomass of microalgae are directly linked to the salinity regime, which depends on the water circulation, driven by regional climate and local weather conditions [37], and can influence the distribution of nutrients in the estuary [38,39].

Despite their small size and very thin leaves, *Ruppia maritima* meadows in PLE hold an abundant epiphytic algal community [29,40] that can be visible to the naked eye during periods of higher productivity. Knowledge about seagrass epiphytes in the PLE is scarce, restricted to a single short-term study that followed the epiphytic colonization process on *R. maritima* leaves and which identified only two species and nine diatom genera [29] that are commonly found in many seagrass meadows around the world [20,22,29,41–49]. Epiphyte composition and diversity in mature seagrass meadows are unknown, limiting the understanding of the role of seagrass in estuarine biodiversity and the causes of local reductions. Knowledge about epiphyte algae diversity can be used for estuarine monitoring and conservation planning. The present study aimed to describe the temporal variability in epiphytic diatom abundance and community composition in *R. maritima* meadows at two distinct sites marked by different levels of anthropogenic impact and different abiotic variables.

Based on the premise that the spatial variability in community composition and abundance of epiphytic diatoms is structured mainly by the availability of nutrients and that temporal variability is mainly determined by salinity and temperature, our hypothesis states that the community attributes of the diatom assemblages will differ between the impacted and non-impacted areas, with higher values expected for the non-impacted site.

2. Materials and Methods

2.1. Study Area

For the study of epiphytic diatoms, two shallow areas of the Patos Lagoon estuary were chosen (Figure 1): Saco da Mangueira (SM) 32°5′13.9704″ S, 52°7′14.9628″ W and Saco da Quitéria (SQ) 32°2′8.3688″ S, 52°14′53.052″ W. The hydrodynamics of the PLE mainly depend on the relationships between the variability in continental discharge, wind action and water exchange with the sea [50,51]. Shallow areas (<1.5 m) provide favorable conditions for the establishment and development of submerged aquatic vegetation [33].



Figure 1. Location of the study area: two meadows of *Ruppia maritima* in the Patos Lagoon estuary (PLE) in the state of Rio Grande do Sul, Brazil. SQ = Saco da Quitéria, SM = Saco da Mangueira. Based on Google Earth.

The SM is an environment with an area of approximately 23 km² that is approximately 10 km long and 3.5 km wide in the central part, which is the widest part. The SQ is a smaller environment with an area of approximately 4.4 km², approximately 3.5 km long and 1.7 km wide in more than half of its area, which tapers southwest to approximately 0.15 km wide. The SM is the largest shallow area of the PLE and is connected to this environment through an entrance of approximately 200 m, which is where the Ponte dos Franceses is located, a road that connects the city of Rio Grande to its industrial area. The SM is more heavily subject to anthropic action (release of effluents) and salt water than the SQ. The two sites were chosen for this study because they present a historical record of the presence of seagrass meadows, dominated by *Ruppia maritima*, and differ in terms of salinity and anthropogenic impact.

2.2. Epiphytic Diatom Sampling and Laboratory Procedures

Monthly monitoring of submerged vegetation was carried out (November 2015 to February 2018) in two *Ruppia maritima* meadows in the PLE located in SM and SQ.

At each site, vegetation samples were collected (N = 3) with a PVC extractor tube (10 cm in diameter and 78.5 cm² in area) buried 15 cm deep in the sediment. The samples were collected, pre-washed in the field to remove the sediment and stored in plastic bags. The material was sorted in the laboratory for the removal of epiphytes using tweezers, brushes and jets of filtered water from the site (pore size 0.45μ m). The removed epiphytes were placed in 100 mL flasks, and half of this volume was fixed with 4% formaldehyde. To identify and quantify diatoms, two permanent slides were made per sample (N = 6 slides per site) according to the oxidation technique from [52] using Naphrax[®] resin (Brunel Microscopes Ltd, Chippenhan, Wiltshire, UK), for a total of 180 slides. At least 400 diatom valves were counted per sample through counts of approximately 200 valves per permanent slide. To determine the relative abundance, the number of valves counted and the number of species found were considered. Abundant species were those with values greater than the average calculated for the sample [53]. The epiphytic diatoms were identified and photographed under optical (Olympus CX 40) and scanning electron (SEM) microscopes at the electron microscopy center (CEME-SUL) of the University Federal do Rio Grande (FURG).

In each meadow, temperature, salinity, depth, transparency (Secchi disk) and water samples (N = 3) were collected at monthly intervals for determination of suspended solids and dissolved inorganic nutrients. The water samples were filtered with pre-weighed cellulose acetate filters (0.45 μ m porosity), dried in an oven (60 °C for one hour) and reweighed to determine the material suspended in the water [54]. The water samples were filtered on the same day of collection and transported to the Ecology Center of the University Federal do Rio Grande dos Sul (CENECO-UFRGS) for analysis of dissolved inorganic nutrients (total ammoniacal nitrogen - TAN (NH₄+), nitrate (NO₃⁻), nitrite (NO₂⁻) and orthophosphate (PO₄⁻)). Dissolved silica in water (silicate) data were obtained from the database of the Long-Term Ecological Program of the Federal University of Rio Grande (PELD-FURG), originating from a sampling point (museum station) located between the two areas of this study.

Sediment samples (cores: 5 cm diameter, buried up to 5 cm in depth) were collected each month until February 2017; after this period, they were collected seasonally to analyze organic matter (OM) content. The samples were analyzed using the calcination method [55]. Data for precipitation (monthly accumulated) and continental discharge (sum of the Jacuí, Taquari and Camaquã Rivers) for 30 days before collection were obtained from the National Institute of Meteorology (INMET) and from the National Water Agency (ANA). Hourly wind data for three days before collection were obtained from the Praticagem of Rio Grande station (RGPilots).

2.3. Data Analysis

Differences in environmental variables between sites were analyzed using Student's *t*-test and the Mann–Whitney test after checking the assumptions of normality (Shapiro–Wilk and Kolmogorov–Smirnov) and homogeneity (Levene test) using Jamovi software [56].

Shannon's diversity (H' \log_2), equitability (J') and dominance (D') indices were calculated from the original matrix (159 taxa) using PAST software [57]. Diatom richness was treated here as the total number of taxa found. To investigate the spatial and monthly variation in richness, diversity, equitability and dominance, Student's *t*-test and the Mann–Whitney test were performed after verifying the statistical assumptions. Temporal comparisons were undertaken by considering the month of collection and the following period (n = 12).

To test the expected spatial distribution and explore the monthly variation in species abundance and composition, two-way PERMANOVAs were performed with a purified matrix based on periods (n = 13) that presented vegetation in both locations (April 2016 to February 2017, May 2017 and February 2018) composed of 80 taxa. The Bray–Curtis distance was calculated from transformed data, and 9999 permutations were used. To test for statistical significance, the p value of the Monte Carlo test was considered when the number of possible permutations was low (<100) for greater test precision [58]. To identify the taxa with the greatest contribution to the differentiation between the samples,

a multivariate analysis of percentage similarity (SIMPER) was performed using PRIMER[®] software, version 6 (PRIMER-e, Albany, Auckland, New Zealand. The ordination of the samples, based on differences in diatom abundance and composition, was analyzed using non-metric multidimensional scaling (NMDS) with the Bray–Curtis similarity index.

The relationships between the diatom community, the physical and chemical factors of the water of each meadow and meteorological variables were examined through canonical correspondence analysis (CCA) using matrices with monthly data for 104 taxa: 87 in the SQ (16 months) and 61 in the MS (14 months). For the analysis of CCA and PERMANOVAs (except composition), matrices with transformed data were used log (x+1), and rare taxa, defined as those whose frequency of occurrence was less than 5%, were excluded [59,60]. To identify relationships between the abiotic variables and the descriptor indices of the community structure (richness, diversity, equitability and dominance), simple linear regression analyses were performed.

3. Results

3.1. Environmental Variables

During the study period, the accumulated rainfall (30 days before each sampling date) ranged from 0 mm (June and July 2016 and September 2017) to 192.6 mm (March 2017) (Figure 2). The wind speed ranged between 2.6 ms⁻¹ (June 2016) and 9.6 ms⁻¹ (April 2017) (average 5.6 ms⁻¹). The fluvial discharge (average 1867 m³ s⁻¹) varied between 435.66 (June 2016) and 5444 m³ s⁻¹ (June 2017) (Figure 2).



Figure 2. Average fluvial discharge, precipitation (30 days prior to each collection) and wind speed (3 days prior to collection) during the study period (November 2015 to February 2018).

The study sites showed differences regarding salinity and nutrient concentrations, but no differences were found regarding water depth, temperature, transparency or suspended material (Table 1). The water depth ranged between 0.37 and 1.17 m (mean 0.75) in the SM and between 0.28 and 1.30 m (mean 0.71) in the SQ. The water temperature ranged between 11.0 and 28.5 °C (mean 19.5) in the SQ and between 9.0 and 27.0 °C (mean 19.1) in the SM. The suspended material values ranged between 3.6 and 166.0 mg L⁻¹ (mean 32.6 mg L⁻¹) in the SQ and between 4 and 128.0 mg L⁻¹ (mean 31.8 mg L⁻¹) in the SM. Water transparency varied from 0.12 to 0.91 m (mean 0.48 m). The average salinity and range were significantly higher (Tukey, *p* = 0.02) in the SM (0.0–25.0, mean 8.8) than in the SQ (0.0–19.0, mean 5.5).

Variables/Periods and Sites		Temperature		ıre Salinity		De	pth	Transparency (Secchi Disk)		Total Suspended Solids		Phosphate		Nitrogen (TAN)		Nitrate + Nitrite		Organic Matter (Sediment)	
		SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM
	November	24.20	20.40	0.20	1.18	0.65	0.43	0.42	0.13	7.33	74.00	0.02	0.08	0.36	0.67	0.68	2.12	0.48	0.74
2015 —	December	21.70	26.50	1.60	6.80	0.55	0.63	0.38	0.20	32.00	42.83	0.01	0.03	0.50	1.28	0.54	0.39	0.70	0.50
-	January	27.60	24.80	0.12	6.61	0.99	0.53	0.35	0.23	23.33	68.93	0.02	0.05	0.07	0.27	0.32	0.39	0.73	0.33
	February	28.50	27.00	0.00	6.00	0.28	0.58	0.28	0.58	20.67	16.00	0.01	0.04	0.70	2.78	0.73	0.49	0.54	0.67
	March	23.50	24.00	17.00	15.00	0.38	0.37	0.38	0.37	38.33	21.87	0.01	0.07	3.16	4.85	0.63	0.49	0.92	0.73
	April	13.00	11.00	19.00	8.00	0.53	0.55	0.53	0.55	19.33	20.93	0.02	0.06	4.01	4.79	0.60	1.31	0.83	0.90
-	May	14.00	12.00	15.00	10.00	0.64	1.17	0.64	0.75	38.89	26.00	0.01	0.04	2.70	1.59	0.48	0.56	0.72	1.40
2016	June	13.00	12.00	5.00	5.00	0.65	0.61	0.65	0.49	29.33	66.67	0.01	0.07	0.71	1.69	0.60	0.61	1.45	1.12
2016 -	July	11.00	9.00	0.00	10.00	0.51	0.81	0.51	0.81	17.67	9.00	0.01	0.01	1.59	3.58	0.47	0.59	1.17	1.83
-	August	16.00	14.00	1.00	6.00	0.52	0.77	0.25	0.77	40.00	24.00	0.01	0.05	0.71	2.66	0.68	0.68	0.78	0.93
	September	19.00	17.00	6.00	6.00	0.63	0.68	0.63	0.25	19.67	10.83	0.01	0.07	2.08	1.66	0.60	0.67	0.82	1.38
	October	19.00	20.00	9.00	10.00	0.86	0.89	0.32	0.34	30.00	10.00	0.01	0.06	2.54	3.74	0.56	0.61	0.79	1.34
	November	19.00	19.50	1.00	5.00	0.65	0.53	0.12	0.53	70.00	6.00	0.04	0.05	1.09	1.67	3.44	1.04	0.59	0.91
	December	22.50	22.50	0.00	11.00	0.80	0.76	0.12	0.52	98.00	16.22	0.02	0.06	0.55	2.62	1.15	0.81	0.47	0.89

Table 1. Values of environmental variables (temperature (°C), salinity (‰), depth, transparency (m), total suspended solids, phosphate, total ammoniacal nitrogen (TAN), nitrate + nitrite (mg L^{-1}) and organic matter in the sediment (%)) during the study period (November 2015 to February 2018). SQ = Saco da Quitéria, SM = Saco da Mangueira.

Variables/Periods and Sites		Temperature		Temperature Salinity		De	Depth Transparency (Secchi Disk)		Total Suspended Solids		Phosphate		Nitrogen (TAN)		Nitrate + Nitrite		Organic Matter (Sediment)		
		SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM	SQ	SM
	January	25.50	24.00	5.00	8.00	0.58	0.74	0.58	0.72	15.33	29.67	0.01	0.04	2.24	2.13	0.90	0.68	0.56	1.35
	February	22.00	22.00	9.00	10.00	0.64	0.86	0.64	0.86	5.60	8.40	0.01	0.05	3.54	3.79	0.78	0.68	0.90	1.45
	March	ND	19.50	ND	7.00	ND	0.97	ND	0.97	ND	23.33	ND	0.06	ND	3.15	ND	0.57	ND	1.18
	April	ND	18.50	ND	25.00	ND	0.81	ND	0.38	ND	25.33	ND	0.11	ND	2.05	ND	0.90	ND	ND
	May	15.00	15.50	18.00	20.00	0.55	0.70	0.55	0.70	20.67	10.27	0.01	0.07	5.00	5.30	0.65	0.69	1.05	1.99
2017	June	13.00	17.00	0.00	0.00	1.29	0.81	0.35	0.81	29.67	19.33	0.02	0.07	0.98	0.70	1.34	1.01	ND	ND
2017	July	17.20	16.50	0.00	0.00	0.80	0.80	0.20	0.36	71.67	49.67	0.03	0.11	0.83	0.93	1.91	1.51	ND	ND
	August	15.50	15.00	4.00	6.00	0.85	0.60	0.85	0.60	7.00	14.00	0.01	0.07	0.52	3.11	1.16	0.93	0.78	1.08
	September	17.50	17.50	0.00	5.00	0.79	0.86	0.32	0.52	18.00	28.00	0.07	0.11	1.03	2.26	1.65	1.44	ND	ND
	October	18.50	18.50	0.00	0.00	1.30	0.98	0.15	0.17	154.00	105.33	0.06	0.04	0.60	0.61	1.95	1.57	ND	ND
	November	21.00	19.80	17.00	20.00	0.91	0.99	0.91	0.55	15.67	23.67	0.02	0.06	2.75	2.72	1.17	1.00	0.51	0.81
	December	21.50	22.00	1.00	6.00	0.69	0.89	0.69	0.15	15.67	74.67	0.03	0.05	0.44	1.77	0.89	0.97	ND	ND
	January	23.00	24.00	5.00	17.00	0.90	ND	0.90	ND	7.50	21.00	0.01	0.04	2.11	3.62	0.71	0.63	ND	ND
2018	February	25.50	26.00	10.00	15.00	0.50	0.90	0.50	0.23	15.00	23.67	0.01	0.01	0.70	4.96	0.80	0.65	0.78	1.56
Test		Stud	ent's t	Ma Whi	ann– itney	Stud	ent's t	Stude	ent's t	Ma Whi	nn– tney	Ma Whi	inn– itney	Ma Whi	inn– itney	Ma Whi	inn– itney	Ma Whi	inn– itney
<i>p</i> -value		0.	756	0.0	025	0.	505	0.6	529	0.8	340	<0.	.001	0.	01	0.	77	0.0	008

Table 1. Cont.

TAN and phosphate values were higher in the SM (p = 0.00). TAN values ranged between 0.20 and 5.44 mg L⁻¹ (mean 2.53 mg L⁻¹) in the SM and between 0.04 and 5.08 mg L⁻¹ (mean 1.60 mg L⁻¹) in the SQ. Phosphate values ranged between 0.01 and 0.11 mg L⁻¹ (mean 0.06 mg L⁻¹) in the SM and between 0.01 and 0.08 mg L⁻¹ (mean 0.02 mg L⁻¹) in the SQ. Nitrite and nitrate values did not differ (p = 0.65) between sites (SM: 0.36–2.18 mg L⁻¹, mean 0.86 mg L⁻¹; SQ: 0.26–3.48 mg L⁻¹, mean 0.98 mg L⁻¹). The sites differed in terms of sediment organic matter content (p = 0.00), which was higher in the SM (0.31–2.3%, mean 1.10%) than in the SQ (0.42–1.79%, mean 0.78%). The variability of the environmental variables is shown in Table 1.

3.2. Community Composition and Abundance

A total of 159 diatom taxa were found on *R. maritima* leaves during the monitoring period (102 in the SM and 126 in the SQ) (Table S1), including freshwater (30.8%), marine (25.1%), brackish (9.4%) and cosmopolitan (8.8%) taxa. Approximately 25.7% of the taxa were not identified at the species level; therefore, their habitats could not be defined. Comparing the sites (periods with vegetation), we found that 32% of taxa were exclusive to the SQ, 23% to the SM and 45% were common to both sites.

The taxa were distributed in 67 genera. The genera with the highest numbers of species were *Nitzschia* (19) and *Navicula* (12), followed by *Amphora* (8); *Tryblionella* (7); *Planothidium* (6); *Surirella* and *Diploneis* (5); *Cocconeis* and *Rhopalodia* (4); *Actinoptychus, Aulacoseira, Cyclotella, Fallacia, Fragilaria, Mastogloia, Pseudostaurosira, Skeletonema* and *Staurosirella* (3); *Achnanthes, Bacillaria, Caloneis, Desikaneis, Entomoneis, Eunotia, Gomphonemma, Halamphora, Hyalodiscus, Melosira, Pleurosigma, Seminavis, Tabularia* and *Terpsinoe* (2). Another 35 genera presented single species.

Throughout the study period, 31 taxa were found to be abundant in the SQ and 21 in the SM, with 14 of these taxa being common to both sites: *Ambo tenuissimus, Bacillaria paxillifera, Catenula adhaerens, Cocconeis placentula, Fistulifera saprophila, Melosira moniliformis, Navicula cruxmeridionalis, N. cryptotenella, N. frustulum, Pleurosira laevis, Pseudostaurosira brevistriata, P. clavatum, Skeletonema sp.3 and T. tabulata (Figures 3 and 4). While 17 abundant taxa were found only in the SQ (Aulacoseira granulata, Chaetoceros sp., Cyclotella meneghiniana, Cyclotella sp.1, Mastogloia pumila, Nitzschia eutinensis, N. filiformis, N. subcohaerens, Psedostaurosiropsis geocollegarum, Rhopalodia rumrichiae, Seminavis strigosa, Skeletonema potamos, Skeletonema sp.2, Staurosirella guenter-grassii, Staurosirella aff. ovata, Tabularia fasciculata and Tryblionella persuadens), only 7 abundant taxa were exclusive to the SM (Fragilaria sp.1, Navicula sp.3, Navicula sp.5, N. microcephala, Paralia sulcata, Planothidium sp. and Pseudopodosira echinus).*

The abundant taxa that were present continuously over time at both sites were *A. tenuissimus, C. adhaerens* and *M. moniliformis.* Three taxa were dominant in the SQ (*T. tabulata* in September and October 2016, *M. pumila* in May 2017 and *M. moniliformis* in February 2018) and three in the SM (*C. placentula* in January, February and May 2017; *T. tabulata* from April to July 2016; and *P. laevis* in September 2016) (Figure 5).



Figure 3. Melosira moniliformis var. octogona (1–14); Pleurosira leavis (15); Aulacoseira granulata (16–19); Chaetoceros sp. (20); S. potamos (21–22); Skeletonema sp.2 (23–24); Skeletonema sp.3 (25–27); Paralia sulcata (28–32); Cyclotella sp.1 (33–39); Cyclotella striata (40); Cyclotella meneghiniana (41–43); Pseudopodosira echinus (44–47); Ambo tenuissimum (48–60); Pseudostaurosira brevistriata (61–66); P. clavatum (67–69); Pseudostaurosiropsis geocollegarum (70–74); Staurosirella aff. ovata (75–82); S. guenter-grassi (83–84); Fragilaria sp.1 (85–86). Scale bar = 10 μm.



Figure 4. Tabularia tabulata (87–91); Cocconeis placentula (92–109); Fistulifera saprophila (110–117); Planothidium sp. (118–122); Navicula cryptotenella (123–129); Navicula cruxmeridionalis (130–133); Navicula sp.3 (134–139); Navicula sp.5 (140–142); Mastogloia pumila (143–146); Bacillaria paxillifera (147); Nitzschia filiformis (148); N. subcohaerens var. scotica (149); N. eutinensis (150–157); N. frustulum (158–164); N. microcephala (165–169); Tryblionella persuadens (170–175); Seminavis strigosa (176–179); Catenula adhaerens (180–185); Rhopalodia rumrichiae (186–188). Scale bar = 10 µm.



Figure 5. Distribution and proportions of the relative abundance (%) of epiphytic diatoms abundant in the non-impacted site (**A**) and impacted site (**B**). Data for diatoms that presented more than 10% abundance in at least one period.

In the sequence we present some information about the five dominant taxa. *Tabularia tabulata* (C. Agardh) Snoeijs

Dimensions: 185.0–490.0 µm long; 4.5–9.0 µm wide; 11–13 striae of 10 µm.

Marine and freshwater diatom, originally described by Agardh in 1983 as *Diatoma tabulatum* in the Adriatic Sea locality. The taxon was found in macroalgal samples in the Baltic Sea [61] at salinities of 4 to 7%. *Tabularia* is an epiphytic and epilitic genus and has a wide distribution worldwide [62]. The relative was similar at both sites (SQ: 0.5–78.7%, mean 19.0%; SM: 0.1–84.5%, mean 22.4%).

Mastogloia pumila (Grunow) Cleve

Dimensions: 25.9–31.4 μ m long; 6.6–8.4 μ m wide; 26 striae of 10 μ m; 6–8 partectas on each side, 4 with sizes of 10 μ m.

Marine diatom described by Cleve for slightly brackish waters of the Baltic and Hawaii. The taxon can be epiphytic, epilithic and epipelic [63]. The genus is marine but can also occur in brackish and freshwater environments [64]. The mean relative abundance was higher in the SQ (0.1–57.9%, mean 8.4%) than in the SM (0.1–0.9%, mean 0.3%).

Melosira moniliformis var. octagona (Grunow) Hustedt

Dimensions: 12.3–33.3 μ m diameter; 22 areolae of 10 μ m on the valve face.

Freshwater diatom that is found in estuarine and marine environments. The species has been found in plankton and sediment samples from the Praia do Cassino and Patos Lagoon Estuary [63,65]. The variety has an octagonal shape in connective view, forming long chains in unoxidized material and double chains after oxidation. Under experimental conditions, it shows optimum growth in salinities between 20 and 30 [66]. The mean relative abundance was similar at both sites (SQ: 0.5–46.8%, mean 7.1%; SM: 0.7–26.6%, mean 8.6%).

Cocconeis placentula Ehrenberg

Dimensions: 17.0–30.5 µm long; 7.6–18.0 µm wide; 18–23 striae of 10 µm.

Freshwater diatom originally found in the roots of the aquatic macrophyte *Lemna* L. by Ehrenberg. It is found in phanerogams, such as *Ruppia maritima* [29], *Zostera noltei* [67] and *Posidonia oceanica* [48]. The genus and taxon are commonly found with epiphytic habits. The mean relative abundance was higher in the SM (2.3–75.1%, mean 27.8%) than in the SQ (0.3–18.5%, mean 6.1%).

Pleurosira laevis (Ehrenberg) Compère

Dimensions: 65.9–131.4 µm diameter; 11–16 areolae of 10 µm.

Brackish water diatom with cylindrical cells connected by ocelli, forming filaments in a zig zag pattern. It is also found in estuarine regions with freshwater, such as the Rio de La Plata estuary (salinity 0.5), where it was found to be epiphytic of *S. californicus* [68]. It has also been found as epiphytic [69], as well as in the plankton [64] and sediment [70], in the PLE. The genus is probably one of the few that has found its ideal niche in brackish waters, such as estuaries [62]. The mean relative abundance was higher in the SM (0.1–53.7%, mean 9.9%) than in the SQ (0.1–4.5%, mean 1.1%).

3.3. Spatial and Temporal Variability

Community Attributes: Diversity, Equitability and Dominance

The mean values of the community attributes indicated variability across months and significant differences between sites for all community attributes (Figure 6). In general, higher richness (19–41), diversity (1.4–4.4; mean 3.3) and equitability values (0.3–0.8; mean 0.7) were found in the SQ, and lower dominance (0.1–0.6; mean 0.2) was found compared to the SM. For the SM, the ranges and average values for those attributes were: richness 8.0–30; diversity 0.9–3.7, mean 2.2; equitability 0.3–0.8, mean 0.5; dominance 0.1–0.7, mean 0.4. We found significant spatial differences for all community descriptor indices.

Richness and diversity differed between sites in most comparisons, except in August and November 2016 and February and May 2017. The sites were similar in terms of equitability and dominance in the periods of August, September and November 2016 and in February and May 2017. In December 2016 and February 2018, the sites were similar only in terms of equitability.

In the visual analysis of the NMDS graphs (Figure 7), as seen in both Figure 7A (abundance) and 7B (composition), it was possible to observe that most of the samples were distributed on opposite sides, revealing that the composition and abundance of the PLE diatom community in the more heavily impacted site (SM) differed from the non-impacted site (SQ).



Figure 6. Monthly variability in richness (**a**), diversity (**b**), equitability (**c**) and dominance (**d**) for the epiphytic diatom community of the Saco da Quitéria (SQ) and Saco da Mangueira (SM) during the study period.

The results of the NMDS were confirmed statistically with PERMANOVAs, which revealed that meadows were distinct in terms of diatom composition and abundance (p < 0.05). According to the Monte Carlo test, the meadows were similar in terms of composition only in the May 2016 and January 2017 periods (p > 0.05). At both sites, temporal variability in the diatom composition and abundance was detected (p < 0.05). In pairwise comparisons, when analyzing the month of collection and the following month (N = 12), we observed distinctions regarding composition (SQ: N = 6; SM: N = 3) and abundance (SQ: N = 10; SM: N = 8). At both sites, the abundance had a more pronounced variability than the composition, which varied more in the SQ than in the SM (Table 2). The differences between the sites in terms of abundance and composition were also observed in the NMDS analyses (Figure 7).



Figure 7. Results of non-metric multidimensional scaling (NMDS) analysis of samples from both sites during the study period. (**A**) abundance and (**B**) composition (presence and absence). Matrix composed of 80 sp. in both analyses. Q =Saco da Quitéria; M =Saco da Mangueira.

Table 2. Summarized results from the PERMANOVAs. Spatial and temporal variation (monthly) in the abundance and composition of the epiphytic diatom community in *R. maritima* meadows (Saco da Quitéria = SQ, Saco da Mangueira = SM). The months are represented in abbreviated form. For the Monte Carlo test, a *p* value < 0.05 = significant difference.

PERMANOVA Results												
	Spatial V	ariation		Temporal Variation								
	Sites	Abundance	Composition	Marchler	Abun	dance	Composition					
S	Q imes SM	p = 0.0001	p = 0.0001	Monutry	<i>p</i> = 0	.0001	<i>p</i> = 0	.0001				
	Comparisons (Monte	e Carlo test <i>p</i> -val	ue)	Comparisons (Monte Carlo test <i>p</i> -value)								
$\mathrm{SQ}\times\mathrm{SM}$	April 2016	0.0010	0.0115		SQ	SM	SQ	SM				
$\mathrm{SQ}\times\mathrm{SM}$	May 2016	0.0084	0.0568	April $ imes$ May 2016	0.0155	0.1127	0.0485	0.2005				
$SQ \times SM$	June 2016	0.0015	0.0049	May \times June 2016	0.0180	0.0326	0.0455	0.0999				
SQ imes SM	July 2016	0.0052	0.0183	June \times July 2016	0.0181	0.0115	0.0168	0.0307				
SQ imes SM	August 2016	0.0187	0.0216	July $ imes$ August 2016	0.0691	0.0131	0.0892	0.0904				
SQ imes SM	September 2016	0.0061	0.0201	August \times September 2016	0.0228	0.0135	0.0529	0.1170				
SQ imes SM	October 2016	0.0061	0.0076	September \times October 2016	0.0394	0.0216	0.0321	0.1540				
SQ imes SM	November 2016	0.0104	0.0236	October \times November 2016	0.0243	0.0106	0.0625	0.0165				
$\mathrm{SQ}\times\mathrm{SM}$	December 2016	0.0050	0.0128	November \times December 2016	0.0341	0.0412	0.1373	0.0281				
$\mathrm{SQ}\times\mathrm{SM}$	January 2017	0.0400	0.0565	December \times January 2017	0.0523	0.1226	0.1529	0.1506				
$\mathrm{SQ}\times\mathrm{SM}$	SM February 2017 0.0126 0.0446		0.0446	January $ imes$ February 2017	0.0298	0.2053	0.0622	0.2385				
$\mathrm{SQ}\times\mathrm{SM}$	Q × SM May 2017 0.0078 0.0298		0.0298	February \times May 2017 0.0		0.1874	0.0127	0.6075				
$\mathrm{SQ}\times\mathrm{SM}$	February 2018	0.0063	0.0105	May $ imes$ February 2018	0.0051	0.0196	0.0139	0.0735				

According to the SIMPER analysis, the species with the greatest contributions to the dissimilarities between sites in terms of abundance were: *T. fasciculata, M. pumila* and *R. rumrichiae* (April 2016); *B. paxillifera, T. fasciculata* and *A. tenuissimus* (May 2016); *N. eutinensis, M. pumila* and *P. brevistriata* (June 2016); *B. paxillifera, M. pumila* and *T. fasciculata* (July 2016); *R. rumrichiae, N. cryptotenella* and *M. pumila* (August 2016); *P. laevis, T. tabulata* and *P. brevistriata* (September 2016); *T. tabulata, C. adhaerens* and *P. sulcata* (October 2016); *Skeletonema* sp.2, *P. laevis* and *Navicula* sp.3 (November 2016); *P. laevis, P. brevistriata* and *S. guenter-grassii* (December 2016); *T. tabulata, T. fasciculata* and *P. brevistriata* (January 2017); *C. placentula, T. fasciculata* and *T. tabulata*, *P. laevis* and *C. placentula*, *M. pumila, T. tabulata* and *C. placentula* (May 2017); *N. microcephala, P. laevis* and *P. sulcata* (February 2018).

The periods with the highest mean dissimilarities (more than 70.0) were July 2016 and May 2017. The taxa that most contributed to dissimilarities between sites in July 2016 (*B. paxillifera* (6.2%), *M. pumila* (6.2%), *T. fasciculata* (5.4%)) had greater abundances in the SQ. In May 2017, *M. pumila* (12.6%) and *T. tabulata* (9.6%) had higher abundances in the SQ, while *C. placentula* (9.4%) had higher abundances in the SM. In general, most taxa that contributed to the dissimilarity had higher abundances in the SQ. The first three taxa of the SIMPER analysis contributed approximately 30% of the dissimilarity between sites. Only three periods had mean dissimilarities below 50 (May 2016, June 2016 and August 2016), again confirming that the sites were distinct in terms of abundance (Table 3).

For the temporal variability of the community, among the comparisons made (n = 12), periods of distinction regarding richness occurred in all the comparisons (SQ: n = 3; SM: n = 6), and they were also noted for diversity (SQ: n = 7; SM: n = 5), equitability (SQ: n = 6; SM: n = 4), dominance (SQ: n = 6; SM: n = 4), abundance (SQ: n = 10; SM: n = 8) and composition (SQ: n = 6; SM: n = 3).

Table 3. SIMPER analysis with the three taxa that most contributed (%) to the dissimilarities between
the impacted and non-impacted areas. AD = average dissimilarity.

Month	Average Dissimilarity		Taxa and Contribution (%)	
April 2016	AD = 52.56	T. fasciculata (12.51)	M. pumila (10.6)	R. rumrichiae (9.75)
May 2016	AD = 48.57	B. paxillifera (7.86)	T. fasciculata (7.39)	A. tenuissimus (6.69)
June 2016	AD = 46.37	N. eutinensis (10.28)	M. pumila (9.54)	P. brevistriata (8.74)
July 2016	AD = 71.71	B. paxillifera (6.17)	M. pumila (6.17)	T. fasciculata (5.41)
August 2016	AD = 45.41	R. rumrichiae (6.14)	N. cryptotenella (6.01)	M. pumila (5.9)
September 2016	AD = 67.21	P. laevis (8.26)	T. tabulata (6.91)	P. brevistriata (4.39)
October 2016	AD = 61.63	T. tabulata (8.81)	C. adhaerens (7.09)	P. sulcata (6.44)
November 2016	AD = 61.15	Skeletonema sp.2 (8.44)	P. laevis (6.04)	Navicula sp.3 (6.02)
December 2016	AD = 66.15	P. laevis (8.25)	P. brevistriata (7.9)	S. guenter-grassii (6.76)
January 2017	AD = 52.80	T. tabulata (8.52)	T. fasciculata (4.88)	P. brevistriata (4.5)
February 2017	AD = 54.17	C. placentula (11.85)	T. fasciculata (10.38)	T. tabulata (9.07)
May 2017	AD = 70.73	M. pumila (12.65)	T. tabulata (9.64)	C. placentula (9.36)
February 2018	AD = 53.12	N. microcephala (9.81)	P. laevis (5.44)	P. sulcata (5.39)

3.4. Correspondence between Epiphytic Diatom Assemblages and Environmental Variables

Canonical correspondence analysis (CCA) showed that the percentage of variance explained by the relationship of the physical and chemical variables with the taxa was 64% (sum of axes 1, 2 and 3) (Figure 8).





Figure 8. Canonical correspondence analysis (CCA) of the epiphytic diatom community in relation to environmental variables across the entire study period (November 2015 to February 2018) at both sites (Saco da Quitéria and Saco da Mangueira). Taxa with a frequency greater than 5%. Transformed data log (x + 1).

The CCA differentiates the sites, with most samples located on the positive side of axis 1 being from the SQ and those on the negative side from the SM. Axis 1 was mainly positively correlated with phosphate and organic matter and negatively correlated with suspended material. Axis 2 was positively related to salinity and suspended material and negatively related to temperature and nitrate + nitrite.

In the SM, the taxa with the highest abundances in periods with the highest phosphate concentrations were: *P. leavis, C. radiatus, T. circumsuta, N. cruxmeridionalis* (September 2016) and *C. placentula* (May 2017). May 2017 was also the period with the highest organic matter, TAN and salinity values. The taxa *T. americana, T. compressa, M. moniliformis* and *N. microcephala* had their highest abundances in a period with lower phosphate concentrations (February 2018). Despite the low concentration of phosphate, the period of February 2018 had a high concentration of TAN and high salinity. *Navicula* sp.3 was more abundant in the period with lower suspended material (November 2016). In the SQ, the taxa with the highest abundances in periods with lower salinities were *C. crucicula* (November 2015), *C. meneghiniana* (November 2016) and *Cyclotella* sp.1 (December 2016). November 2016 was also the period with the highest concentration of nitrate + nitrite, phosphate and suspended material. The suspended material had a negative relationship with the water transparency in the SQ (R² = 0.58).

Comparing the community structure with the environmental variables, in the SM, richness, diversity and equitability were positively (but weakly) related to temperature ($R^2 = 0.24$, 0.20 and 0.17) and negatively related to dominance ($R^2 = 0.20$); in the SQ, richness was negatively related to organic matter and TAN ($R^2 = 0.12$ for both variables). Salinity was negatively related to richness and diversity ($R^2 = 0.30$ for both descriptors).

Our results supported the hypothesis that the community attributes of the diatom assemblages would differ between the impacted and non-impacted areas, with higher values found in the non-impacted site and higher dominance in the impacted site.

4. Discussion

The results of the present study revealed that meadows of *Ruppia maritima* in PLE sustain rich and diverse epiphytic diatom communities. The total number of diatom taxa found in our study (68 genera and 159 sp) was much higher than values reported for other *Ruppia* meadows from estuarine areas [40,44]. The richness and composition found here were similar to those found for local brackish marsh plants located in a pristine site [69]. This study also revealed that community structure and composition vary according to season and between sites, driven mainly by changes in salinity and nutrient conditions.

The study sites were marked by significant differences in salinity, nutrients (phosphate and TAN) and sediment organic matter. The higher values of salinity, nutrients and organic matter found in the SM compared to the SQ may have been related to proximity to the ocean and eutrophication sources, entailing receipt of both domestic waste and industrial effluents [71]. Considering that the study sites were classified as brackish water—i.e., intended for the preservation and balance of aquatic communities—the values of nitrogen (TAN) and nitrate found were above the limits established by Brazilian legislation (Nacional Environmental Concil—CONAMA 357/2005 [72]): on average, 6.3 and 4.0 times higher in the SM and SQ for TAN and, in terms of means, 1.2 and 1.4 times higher in the SM and SQ for nitrate.

Values within the limit defined by CONAMA were found only in the period of November 2015 in the SQ (TAN) and January 2016 in both sites (TAN and nitrate). Phosphate concentrations were compared to values considered normal in unpolluted estuaries (0.03 mg L⁻¹; [73,74]), and they were higher in the SQ only in the periods of September and October 2017. In the SM, there were lower values only in the periods of December 2015, July 2016 and February 2018.

According to our results, *R. maritima* is an important host for epiphytic diatoms, both in impacted and non-impacted areas. The fact that the same species could be studied at both sites implied that there was similar availability of microhabitats for epiphytes due to their

similar morphologies. The abundance and diversity of epiphytic communities depend on the host plant, since different seagrass morphologies tend to promote distinct microhabitats for epiphytes [49]. Another study, however, found no differences in community structure among seagrass species during a short sampling period, which agrees with the neutral substrate hypothesis [42]. According to this hypothesis, the substrate does not stimulate or prevent the growth of epiphytes. However, while some macrophytes can be neutral hosts for their epiphytes, others can profoundly affect their production and, in both cases, the community composition can vary significantly between hosts [75]. Despite the similar availability of microhabitats, the differences in the epiphytic diatom communities between the sites did not reveal specificity for *R. maritima*, which would have indicated the neutrality of this substrate. However, the anthropogenic impact showed a greater influence on the epiphytic diatom community than the morphological similarity of the substrate.

While host preference is an unresolved question in structuring epiphytic diatoms [76], environmental conditions may have equal or greater importance. Despite colonizing the same substrate (*R. maritima*), the studied epiphytic diatom community showed high heterogeneity between sites. The greater richness, diversity and equitability and higher percentage of exclusive taxa at the SQ site compared to the SM site reflect a better preserved environment and higher water quality. In contrast, high nutrient loads decrease diatom species diversity and promote the dominance of a few taxa [77–79], as we observed in the SM. Epiphytic diatoms in sites with high levels of nutrients have a composition that favors dominant species, while sites with lower levels of nutrients, although containing these same species, have lower abundances [80]. Spatial differences in nutrient availability may exert the greatest control over regional variations in epiphyte loads in seagrasses [81].

In addition to nutrient availability, salinity may be a structuring factor in the spatial distribution of the seagrass epiphytic diatom community [22]. We found spatial differences in composition and abundance (CCA) marked by both nutrient availability and salinity (higher in the SM). While salinity negatively affected richness, diversity and equitability in the SQ, the SM community structure was not affected by the variability in salinity, indicating that diatoms may be more generalist and tolerant to salinity variability. In tropical seagrass meadows, differences between communities of epiphytic diatoms growing on *T. testudinum* were related to salinity levels (marine *versus* hypersaline) [82]. In the hypersaline site, there were greater numbers of rare taxa and lower richness and diversity than in the more stable marine environment. In our study, the degree of anthropogenic impact and proximity to the ocean influenced the epiphytic communities at the study sites.

In a study with epiphytic diatoms from the Baltic Sea, no reduction in richness related to salinity decrease was observed [26]. Another study found that, in the absence of anthropogenic impacts, stable marine environments enhance algal species' richness and diversity in *R. cirrohosa* meadows [83]. In our study, the variability in salinity was very high throughout the year (from 0 to 30), depending mainly on the seasonal cycle of precipitation and fluvial discharge. The autumn months were marked by higher values. Temporal stability and range variability may be more important structuring factors than the average salinity.

Seasonal changes in the epiphytic diatom community were found to be largely related to temperature [22], which was also the main factor responsible for changes in the epiphytic diatom community in subtropical mangroves in South China [84], while lower richness and diversity in diatom epiphytes in marsh plants has been found during winter [85]. For seagrasses, epiphytic biomasses are usually higher in summer months [49,86]. Additionally, some authors found that temperature was positively related to the richness of algal species associated with *R. cirrhosa* [83], and periods of increase in the diversity of epiphytic diatoms accompanied increases in temperature [87]. The low diversity of epiphytic diatoms during the winter and spring periods and the high diversity during the autumn and summer have been attributed to plant substrate abundance, influenced by freshwater discharge [88]. In addition to seasonality factors (temperature, salinity) and plant substrate availability, the variability in the nutrient pulses was also found to influence the high variability in the composition and abundance of the community over the months.

In our study, seasonality appeared to have distinct effects on community attributes. While it may have affected community composition, our analysis showed that temperature did not have significant effects on diatom community diversity and equitability in the non-impacted site SQ. Except for the reduction during the spring period (September and October 2016), diversity and equitability in the SQ remained high during most of the study period. However, in the impacted site SM, temperature appeared to be related to community variability. In addition, months with similar average temperatures had similar richness, diversity, equitability and dominance (April and May 2016, May and June 2016 (except for equitability) and June and July 2016; and November and December 2016, December 2016 and January 2017 (except richness) and January and February 2017) (Figure 6a–d).

Dominant diatom species are the main food source for grazers [89], and the reduced abundance of high-profile dominant taxa, such as *T. tabulata* (August 2016) and *P. laevis* (October 2016), may have been caused by grazing. A previous study [90] found higher densities of benthic macrofauna in the PLE in the summer period, which was related to temperature, but, in February 2018, the non-dominance of *C. placentula*, observed in the previous period, reflected a community that was more developed and, possibly, experiencing low herbivory. In the SM, abundance in most periods was different but similar in periods of similar temperature (April and May 2016, December 2016 and January 2017 and January and February 2017). The taxa with similar abundances were *T. tabulata* and *C. placentula*, which were dominant in some periods. With regard to composition, most periods were similar. The difference in composition between the periods of June and July 2016 may have been due to the different concentrations of nutrients (phosphate), whereas the periods of June and July 2016, October and November 2016 and November and December 2016 may have been distinct due to the reduction in salinity in November 2016.

Increases in TAN concentration and salinity may have altered the community in the SQ, as they coincided with periods of richness and diversity reduction (March, April and October 2016, February and May 2017 and February 2018), and some species were probably not tolerant. Since salinity and TAN concentration varied together, it was difficult to determine which of these factors was most important, but we found that they had no positive effect on the community. The high levels of variability in salinity and TAN concentration can be considered stress factors for the community, but some taxa, such as T. fasciculata and R. runrichiae, were tolerant to variations and were present in abundance under different salinity and TAN conditions. Periods of similar diversity, equitability and dominance (April and May 2016, July and August 2016, November and December 2016 and December 2016 and January 2017) had similar salinities. In July 2016, an increase in diversity occurred together with a reduction in salinity to zero, which may explain the significant difference found in relation to the previous period (June). The observation of differences between the periods in both abundance and composition (April and May 2016, May and June 2016, June and July 2016, September and October 2016, February and May 2017 and May 2017 and February 2018) showed that epiphytic community taxa are constantly dynamic, possibly as a result of the variability of environmental conditions, such as salinity and nutrient availability.

According to the CCA results, the community attributes were significantly related to salinity variability (Figure 8). Freshwater taxa, such *S. potamos*, *N. eutinensis*, *P. brevistriata*, *P. clavatum* and *R. runrichiae*, showed higher abundances in the SQ. The negative relationship of *S. potamos* with salinity was observed by [91] in phytoplankton samples from Patos Lagoon and confirmed in our study. Some marine taxa had higher abundances in the SM (*C. jonesiana* and *P. sulcata*), and also in the SQ, during periods of higher salinity (*N. obtusa*). A greater distance from the estuarine mouth and high variability in salinity (from 0 up to 25), as observed at the SQ site, favor the development of abundant freshwater species, which developed mainly during winter and spring. With increased salinity (summer and autumn), freshwater diatom species were reduced in abundance. At high salinity, freshwater diatoms may not have the ability to produce the osmolarity necessary to generate the same turgor

pressure as at low salinity [92]. However, some freshwater taxa negatively related to salinity were still present in lower abundances at intermediate salinities (17 and 19), which shows that they can tolerate salinity variability. As richness and diversity in the SQ were negatively related to salinity and TAN, and these factors vary together, it was difficult to distinguish the main influencing factor. Perhaps environmental fluctuations can lead to the coexistence of various species based on different responses to environmental gradients [66].

The results of the CCA revealed that part of the community variability was also related to nutrient concentrations (Figure 8). Some abundant taxa tolerated the high N and P concentrations found in the SM. Some of these taxa have already been recorded in environments with high nutrient loads, such as C. meneghiniana, N. cryptotenella [93], F. saprophila [94], N. microcephala [95], P. sulcata [96] and P. laevis [97]. Cocconeis placentula also showed tolerance to the highest concentrations of nutrients in the SM, being dominant in May 2017, the period with the highest concentration of phosphate and organic matter. The greater availability of nutrients and lower competition for light in the biofilm may have stimulated the growth of the taxon. In a study with epiphytic diatoms on Z. marina, a positive response from *C. placentula* to nitrate enrichment was found [98]. Its dominance in the SM may have been the result of a disruption of the community involving a reduction in the abundance of larger species, such as *M. moniliformis* and *P. leavis*, which can overlay the innermost layers of the biofilm, where C. placentula is found. This taxon represents the pioneer stage of succession in R. maritima [29] but remained abundant throughout the study period in both study sites. According to [18], low-stature and prostrated taxa, such as C. placentula, are the dominant forms at low nutrient levels. Additionally, in the periods when this taxon was dominant (January, February and May 2017), there was a reduction in the diversity of the community. A previous study [99] also found a reduction in the diversity value when C. placentula was dominant. Some abundant taxa, such as M. pumila and T. fasciculata, were negatively related to nutrients (phosphate and nitrate + nitrite) and their highest abundances occurred in the SQ. Tabularia fasciculata showed a positive relationship with the increase in nutrients in an experimental study, regardless of the presence of grazers [100]; however, the taxon has already been recorded in abundance as an epiphytic in a preserved environment of the PLE [69]. Mastogloia pumila lives in saline environments, such as ponds, estuaries and salt lakes and salt marshes [69,70,101], and we did not find records of its presence as epiphytic in eutrophic environments. Suspended material was also important in the CCA, and higher amounts represented a reduction in water transparency, confirmed by the negative relation found in the SQ. Diatoms can adapt to low light levels and are able to survive in dark estuaries [102]. Some abundant SQ taxa, such as *Cyclotella* sp.1, S. guenter-grassi, P. brevistriata, S. potamos and T. persuadens, which have positive relations to suspended material, showed that they are also able to develop in higher turbidity in the water column.

5. Conclusions

This study presents, for the first time, a detailed analysis of the spatial and temporal variability of the epiphytic diatom community in *Ruppia maritima* meadows in Brazil. The meadows of the Patos Lagoon estuary sustain abundant and diverse epiphytic assemblages of diatoms, which are important primary producers and food sources for local economic species, such as invertebrates (shrimp) and small-sized fish (such as juveniles of the Mugilidae family). The distinction between the communities in the impacted and non-impacted sites showed the influence of the anthropogenic nutrient load on epiphytic diatoms, which are less diverse under environmental impacts. The temporal variability of the community structure was influenced by abiotic factors, such as salinity and temperature. The present study contributes to the knowledge about benthic diatoms, one of the most abundant and important groups of epiphytic organisms. Reductions in seagrass meadows, as observed during recent decades, may decrease the complexity of benthic habitats, which is likely to change the abundance and diversity of epiphytic diatoms and their ecological functions, and these are still poorly understood. In addition to studies on other epiphytic groups, future

directions should integrate studies on the effect of nutrient increase and grazer pressure on the abundance and diversity of epiphytic algae and the consequences for seagrass abundance.

Supplementary Materials: The following supporting information can be downloaded at: https: //www.mdpi.com/article/10.3390/d14100787/s1, Table S1. Complete list of epiphytic diatoms found on *Ruppia maritima* meadows of Patos Lagoon Estuary between November 2015 and February 2018, registered montly in two studied sites: Saco da Quitéria (SQ) and Saco da Mangueira (SM). The relative abundance (%) of each taxa is indicated for each month. Abundant taxa (•).

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