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Anthropogenic Impacts on Coral Reef Harpacticoid Copepods

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Abstract: The number of studies demonstrating the susceptibility of benthic reef communities to anthropogenic impacts is growing. However, for some of the components of reef fauna, such as meiobenthic harpacticoid copepods, information is still lacking. Here, different diversity and taxonomic distinctness indexes and multivariate analyses were used to test whether the assemblage of harpacticoid copepods colonizing Artificial Substrate Units (ASUs) is an appropriate tool for the identification of reefs subjected to different levels of anthropogenic pressure. Furthermore, we also evaluate if diffused, persistent, anthropogenic impacts generate the homogenization and simplification of Harpacticoida assemblages. Six reefs were organized into two groups along the coast, depending on their proximity to very large urban centers. ASUs were used for meiofauna colonization and, for each reef, 320 Harpacticoida individuals were separated for identification at the species level. Abiotic parameters were analyzed, and significant differences were found between the two groups of reefs, with an increase in dissolved inorganic nutrients found in areas near large urban centers. Both the multivariate analyses and the indexes of diversity showed a clear separation between the reefs closer to the urban zones and those further away, as a response to the anthropogenic pressure. As hypothesized, in the impacted reef areas, there was a strong simplification and homogenization of the harpacticoid copepod assemblages. However, the results of the indexes, based on taxonomic distinctness, suggest that there was no phylogenetic signal of anthropogenic impact on coral reef harpacticoid copepods.

Keywords: meiobenthos; diversity indices; phytal; urban pollution

1. Introduction

Anthropogenic impact is changing our planet. Human pressure continues to affect the ability of ecosystems to provide essential services [1]. Understanding how ecosystems and ecological communities respond to global changes is one of the main challenges faced by ecologists [2]. As a result, impact/environmental quality indicators and organism models are sought after and tested in the most diverse marine ecosystems [2].

On a global scale, coastal, estuarine and transition ecosystems are strongly impacted by human activities [3], such as overfishing, eutrophication, tourism, and engineering works [4,5]. This is due to the development of human populations historically being concentrated in coastal areas, causing profound alterations in these environments [6]. Recently, studies have highlighted the negative

effects of anthropogenic disturbances on coastal benthic communities, such as increased seawater temperatures in tropical environments [7], ocean acidification [8,9], and diffused source pollution [10]. In addition, during the last few decades, scientific literature has pointed out that human activities are not random in their negative and positive impacts on the communities in question [11–14]. Evidence shows that most species show a decline in population size when subjected to anthropogenic impacts (“losers”), and are being replaced by a much smaller number of resistant/opportunistic species that thrive in environments impacted by human activity (“winners”) [15]. The result is a homogenization of communities, with less diversity on both regional and global scales [15].

The only reef formations in the southwest Atlantic Ocean are found in Brazil, and they are among the most prominent marine ecosystems in tropical Brazil. Furthermore, due to their distribution along 3000 km of coastline, they are found within some of the most biologically diverse marine ecosystems [16–18]. In addition to their use as breeding and nursery areas, they shelter and feed a huge variety of marine organisms [19]. Despite the importance of goods and ecological services provided by reef systems, such as fishery production and reef tourism [19,20], the extension of coastal reefs in northeast Brazil shows signs of being strongly impacted as a result of uncontrolled urban development, tourist activities, and the eutrophication of water in association with pollution [16–18,21]. Furthermore, in zones located closest to large urban centers, it has been found that anthropogenic impacts are more aggravated, with an increase in sedimentation due to the removal of fragments from the Atlantic Forest and an increase in the release of effluents, such as waste, which appear to be the main factors responsible for this alarming situation [22].

The use of artificial substrates units (ASU) in studies evaluating environmental quality has been recommended in recent years as a potential solution to bypass the effects of natural variation in substrate structure or characteristics [23–26]. This reduces the heterogeneity between replicates of the same association and, as a result, reduces the effort needed to detect impacts. For ASUs previously tested in studies with meiofauna, abundance values of more than 3000 individuals per ASU were usually found (e.g., References [23,25]), which allows ASUs to be implemented as appropriate tools in environmental monitoring studies.

Harpacticoid copepods have a high diversity in phytal environments [27], fast life cycles, a large ecological importance in the transfer of energy to higher trophic levels [28,29], and are sensitive to anthropogenic impacts [30,31]. These characteristics make the group a very good indicator in studies of environmental impacts, particularly when associated with artificial substrates [32].

In 1995, Warwick and Clarke [33] proposed that the phylogenetic structure of a community is clearly important in environmental impact studies. However, the majority of studies investigating aspects of the application and interpretation of results from taxonomic distinction indexes in benthic communities (e.g., References [34–41]) were performed in estuarine areas.

This study evaluates the use of different diversity and taxonomic distinctness indexes and multivariate analyses to test whether assemblages of Harpacticoida colonizing Artificial Substrate Units (ASUs) are an appropriate tool to identify reefs under different levels of anthropogenic pressure. Furthermore, considering that as a result of human activities, most species tend to decline (i.e., losers), and are replaced by a few non-randomly distributed taxa (i.e., winners) [15], we also evaluate if diffused, persistent, anthropogenic impacts associated with large urban centers generate the homogenization and simplification of Harpacticoida assemblages in coral reefs.

2. Materials and Methods

The coastal reefs included in this study were divided into two groups: the Impacted Areas (Rio Doce, Piedade and Paiva), due to their location in the metropolitan region of Recife (which has over four million inhabitants with an average density of over 1000 inhabitants/km², and is the region subjected to the outflow of several highly polluted rivers), and the Control Areas which were distant from large urban centers (Porto de Galinhas, Serrambi and Tamandaré). This allowed for the comparison of environmental conditions and meiofauna compositions between the two groups (Figure 1).

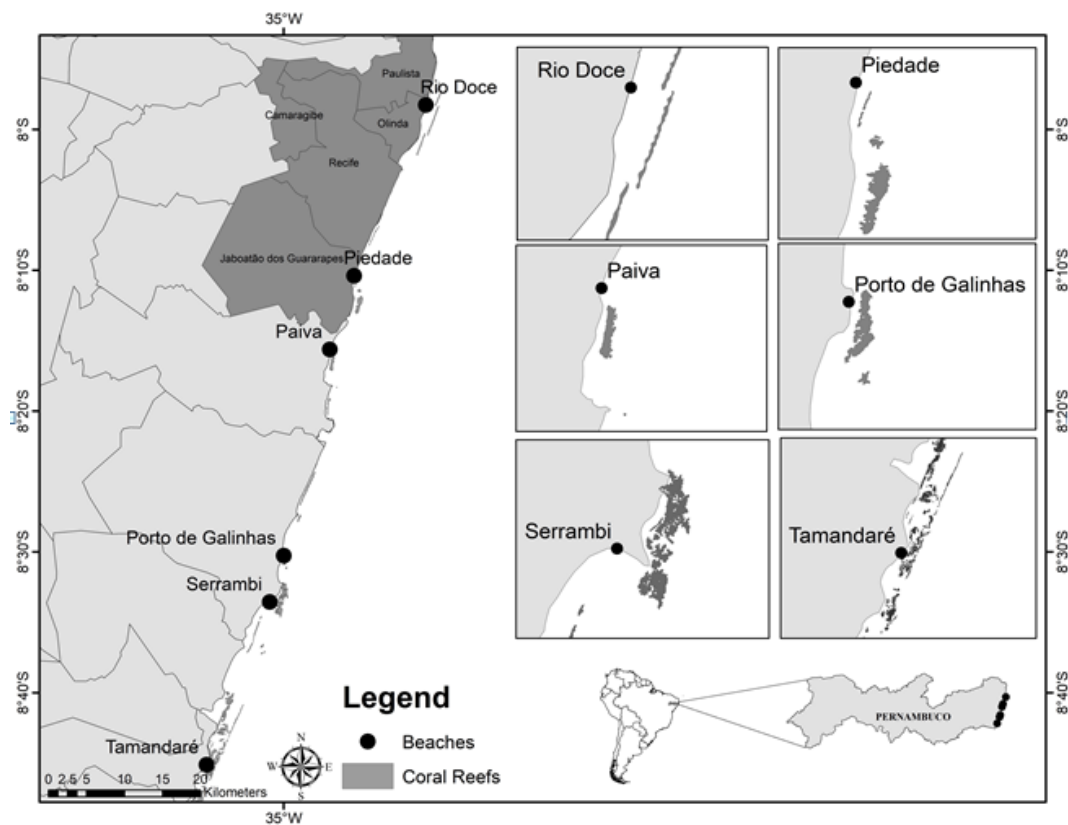


Figure 1. Map of the coastal region of Pernambuco, Brazil, indicating the experimental reef areas. Dark grey areas indicate an average density above 1000 inhabitants/km².

A portion of reef was selected for the experiment implantation in each reef. Each portion selected was the area of reef furthest from the edge of the beach and reasonably well protected from humans bathing, so as to preserve the integrity of the experiment. Since access to these areas depends on tide levels, low tides were selected (0.1 to 0.4 m) as the most adequate times for the implantation procedure and the collection of ASUs, as well as for the analysis of abiotic parameter samples.

During the experiment implantation and removal of the ASUs, temperature and salinity were measured in situ with a thermometer and an optical refractometer (Instrutemp ITREF 10) respectively, and parallel seawater samples were collected using an oceanographic Niskin bottle for chemical analyses at the Chemical Oceanography Laboratory (LOQuim). The dissolved inorganic nutrients were measured by colorimetric methods using a spectrophotometer (Cary 100). Ammonium ($\text{NH}_3 + \text{NH}_4^+$) levels were measured using the blue indophenol method described by Grasshoff et al. [42]. Nitrate (NO_3^-) and nitrite (NO_2^-) levels were measured using the sulfanilamide ($\text{C}_6\text{H}_8\text{N}_2\text{O}_2\text{S}$) method [43]. Dissolved inorganic nitrogen (DIN) was represented by the sum of NO_3^- , NO_2^- , NH_3 and NH_4^+ concentrations.

Concentrations of silicate (SiO_2) and dissolved inorganic phosphorus (DIP) were determined based on the production of a blue compound when orthosilicic acid and phosphate ions (PO_4^{3-}), reacted with acidified molybdate [42]. Particulate materials in suspension (PMS) were quantified by a gravimetric method, determined by Strickland and Parsons [42]. Dissolved oxygen (DO) was analyzed following a modified Winkler method [43], and its saturation (DO%) was calculated according to the International Oceanographic Tables [44].

Total Alkalinity (TA) samples were poisoned with mercury chloride (HgCl_2) to prevent biological alterations, and measurements were taken following Dickson et al.'s [45] methodology, represented by its titration with hydrogen chloride (HCl) at 0.1 N. Certified Reference Materials (CRM) provided by Andrew Dickson (Scripps Institutions of Oceanography, San Diego, CA, USA) were used for the calibration and validation of the values. Simultaneously, pH was measured using a Thermo ScientificTM pH meter Star A211.

The ASUs selected for this study were rectangles of 10 × 5 cm synthetic grass that adequately mimic the turf algae environment and have been used in experimental laboratory studies investigating the anthropogenic impacts on meiofauna communities in reef environments [9,46].

The artificial turfs were secured with silk cords using 1.6 mm thick nylon string, minimizing the chances of breaking away and loss of the structures, due to wave action. The cords were tied to the infralittoral zone (1–2 m depth) of the reef with 10 ASU replicates placed in each area. Colonization time was standardized as one month for all reef areas. The decision of experimental duration was based on the elevated temperatures in northeast Brazil (commonly above 22 °C), during which it is expected that the life cycle of meiofaunal organisms allow the occurrence of two to three generations of initial colonizers (see for example, the model for harpacticoid copepods [47]).

Four replicates were used from each area, following the visual evaluation of the ASUs' integrities post-experiment (i.e., the artificial turf was visually, completely covered by colonizers, and nylon strings and silk cords were unbroken). In areas where more replicates were undamaged, four more were randomly selected. The ASUs were added to plastic pots, fixed in 4% formaldehyde solution, and taken to the laboratory where meiofauna were extracted from the substrate and washed successively with running, filtered water over sieves with mesh size 0.5 mm (to remove the macrofauna) and 0.045 mm (to retain the meiofauna).

After the collection of the colonized ASUs and the separation of the meiofauna, the first 80 harpacticoid copepod individuals were removed from each replicate (320 per reef) and were placed in Eppendorf microtubes with 70% alcohol for later analysis. The sample size followed published studies on anthropogenic impacts affecting harpacticoid species colonizing ASUs: Costa et al. [32] (5 replicates per estuary × 30 individuals per replicate); Sarmiento et al. [48] (4 replicates per treatment × 60 individuals per replicate). The individuals were fixed on slides, and the identification of species was accomplished through the observation of the entire animal under an optic microscope, based on taxonomic keys and descriptions [49–51], as well as other publications. Since total counts of harpacticoid copepods were available for all replicate samples, proportions of identified species were adjusted to densities for analysis. A list with harpacticoid copepod species found in reef regions in the state of Pernambuco was created using identification data from Sarmiento and Santos [31], Barreto [52] and Nascimento [53]. This list was considered as a regional list, to which all lists from the reefs included in this study could be compared in order to estimate the taxonomic distinction indexes: taxonomic diversity (Delta), taxonomic distance (Delta*), average taxonomic distance (Delta+), total taxonomic distance (sDelta+), and the variation of taxonomic distance (Lambda+) [33,35,54]. The regional list was organized under Linnean classifications considering five levels of taxonomic hierarchies (species, genus, family, sub-order and order). The Delta and Delta* indexes were calculated based on the untransformed abundance data. The Delta+, sDelta+ and Lambda+ indexes were calculated based on the presence/absence of species in coastal reefs. Shannon-Wiener diversity (H' , using a log of base 2), Margalef's diversity, Pielou's evenness (J'), and species richness indexes were also calculated. The mathematical models followed Clarke and Warwick [33,54] and Salas et al.'s [55] recommendations, and PRIMER[®] software (Plymouth Routines in Multivariate Ecological Researches) packages v6 and PERMANOVA were used for the calculations.

A nested PERMANOVA analysis was performed to evaluate the differences between the two groups of areas (factor: Impact) and between reef areas (factor: Reef) within impact levels for multivariate community structure. To measure the similarity between colonizer fauna samples from the ASUs, the Bray-Curtis index was used. Non-metric multi-dimensional scaling (NMDS) was used to represent the Bray-Curtis matrix graphically in a two-axis space. A similarity percentage (SIMPER) analysis was applied to determine which groups were responsible for the dissimilarities between the two groups of areas (factor: Impact) and PERMDISP was used to test if there were differences in multivariate dispersion. A nested PERMANOVA was also used to evaluate the differences between the two groups of areas for abiotic parameters. To measure the distance between samples considering environmental descriptive parameters, the Euclidean distance was used. Mann-Whitney tests were also used to compare the two groups of areas considering individual abiotic parameters. Student's t-tests were used to compare the two groups of areas in relation to their univariate indexes. In the absence of

the homogeneity of variances, a PERMANOVA was used (which, in this case, is reduced to an ANOVA whose significance is tested by permutations).

Multivariate analyses followed Anderson [56], MacArdle and Anderson [57], and Clarke and Warwick's [58] recommendations, whereas parametric univariate analyses followed application suggestions by Zar [59]. Statistical analyses were performed using PRIMER[®] software packages v6 and PERMANOVA, and using the STATISTICA v12 program. A significance level of 5% was used for all analyses.

3. Results

The results of the physical-chemical characterization of the environment at the time of the experiment implantation and collection of the colonized ASUs can be found in Table 1. Considering the abiotic variables, the control and impacted areas were compared using a nested PERMANOVA analysis, and results indicated a significant difference (Pseudo-F = 1.999; $p < 0.001$). The reefs nested within each area were not significantly different from each other (Pseudo-F = 1.450; $p = 0.227$).

In the control areas, SiO₂ was represented by a mean concentration of 6.67 $\mu\text{mol L}^{-1}$ and, in the impacted areas, 25.15 $\mu\text{mol L}^{-1}$. The dissolved inorganic nutrients (DIN) presented a mean of 2.26 $\mu\text{mol L}^{-1}$ and 8.41 $\mu\text{mol L}^{-1}$, and dissolved inorganic phosphorus (DIP) presented mean values of 0.19 $\mu\text{mol L}^{-1}$ and 0.57 $\mu\text{mol L}^{-1}$ in the control and impacted areas, respectively.

The SiO₂, DIN and DIP were represented by significantly higher concentrations in impacted areas when compared to the control areas ($p = 0.008$; $p = 0.003$; $p = 0.04$, respectively). DO%, salinity, PMS, pH and TA were not significantly different between the impacted and control areas. Significant differences ($p = 0.01$) between these groups were found for temperature ($^{\circ}\text{C}$), with a mean of 28 $^{\circ}\text{C}$ for impacted, and of 29.86 $^{\circ}\text{C}$ for control areas.

Table 1. Mean values (\pm standard deviation) of the environmental variables according to the study areas: PMS (mg L^{-1}), salinity, temperature ($^{\circ}\text{C}$), DO (%), pH, TA ($\mu\text{mol L}^{-1}$), SiO₂ ($\mu\text{mol L}^{-1}$), DIN ($\mu\text{mol L}^{-1}$), and DIP ($\mu\text{mol L}^{-1}$).

Reef Areas	PMS	Salinity	Temperature	DO	pH	TA	SiO ₂	DIN	DIP
Tamandaré	45.1 (± 3.9)	40.5 (± 2.1)	29.5 (± 0.7)	133.4 (± 7.8)	8.20 (± 0.07)	2340.9 (± 23.4)	9.40 (± 4.30)	1.63 (± 0.21)	0.15 (± 0.04)
Serrambi	33.1 (± 6.2)	38.5 (± 0.7)	29.95 (± 1.3)	153.8 (± 33.5)	8.31 (± 0.11)	2315.7 (± 51.1)	5.38 (± 1.57)	2.80 (± 1.56)	0.22 (± 0.08)
P. Galinhas	38 (± 1.4)	37 (± 1.41)	30.15 (± 1.6)	119.4 (± 2.6)	8.22 (± 0.03)	2352.1 (± 0.6)	5.24 (± 1.38)	2.36 (± 1.36)	0.19 (± 0.02)
Paiva	46.6 (± 13.3)	38.5 (± 0.7)	29 (± 1.4)	147.5 (± 13.2)	8.40 (± 0.10)	2313.5 (± 29.6)	16.17 (± 11.98)	8.95 (± 11.23)	0.30 (± 0.10)
R. Doce	61.2 (± 37.3)	38 (± 1.4)	28 (± 1.4)	140.6 (± 24.1)	8.22 (± 0.05)	2317.0 (± 56.3)	39.39 (± 11.54)	11.68 (± 2.77)	1.03 (± 0.12)
Piedade	40.9 (± 4.9)	36 (± 1.4)	27 (± 1.4)	120.5 (± 18.7)	8.30 (± 0.03)	2331.9 (± 7.9)	19.91 (± 19.08)	4.61 (± 3.05)	0.40 (± 0.08)

In this study, 14 families, 35 genera and 63 species of harpacticoid copepods were registered in the six areas of the study's coastal reefs (Table 2). Of these, only 23 species occurred in both the control and impacted areas. The results indicate that *Ectinosoma* sp1 was the species with the highest abundance and was found to be present in all collection areas. In the control areas, we observed a greater number of total (55) and exclusive (32) species when compared to impacted areas, where only eight species were found to be exclusive out of 31 total species. Densities (average \pm standard deviation) were very similar for both areas (impacted: 1078 \pm 410 individuals/ASU; control: 1144 \pm 656 individuals/ASU).

In the control areas, 12 species were found to have representative relative abundances (relative abundance $\geq 3\%$ of the total individuals in the area), with *Stenhelia* sp. (10.75%), *Parastenhelia spinosa* (5.25%) and *Amphiascoides* sp1 (5%) having the highest values. Whereas, in impacted areas, only six species were considered to be representative, with *Ectinosoma* sp1 (29%), *Ameira* sp2 (17%) and *Scutellidium* sp. (7.25%) having the highest values.

Table 2. List of harpacticoid copepod species identified on the artificial substrates obtained in the control areas (Porto de Galinhas, Serrambi and Tamandaré) and impacted areas (Rio Doce, Piedade, Paiva) in Pernambuco, Brazil, together with percentage contribution of species in each area.

Sub-Order	Family	Genus	Species	Control	Impact
Polyarthra	Longipediidae	<i>Longipedia</i>	<i>Longipedia</i> sp.	1.9	1.3
Oligoarthra	Ameiridae	<i>Ameira</i>	<i>Ameira parvula</i>	3.4	0.0
			<i>Ameira</i> sp1	5.5	0.0
			<i>Ameira</i> sp2	2.2	15.7
			<i>Ameira</i> sp3	0.5	2.9
			<i>Ameira</i> sp4	0.4	0.0
		<i>Sarsameira</i>	<i>Sarsameira knorri</i>	0.9	0.0
			<i>Sarsameira</i> sp1	0.0	0.4
			<i>Sarsameira</i> sp2	0.7	0.0
	Canthocamptidae	<i>Mesochra</i>	<i>Mesochra</i> sp1	0.9	0.0
			<i>Mesochra</i> sp2	0.2	1.5
			<i>Mesochra</i> sp3	0.2	0.0
		<i>Nannomesochra</i>	<i>Nannomesochra</i> sp.	0.8	1.0
	Dactylopusiidae	<i>Dactylopusia</i>	<i>Dactylopusia</i> sp1	0.8	0.0
			<i>Dactylopusia</i> sp2	1.9	1.9
			<i>Dactylopusia tisburyoides</i>	3.2	5.8
		<i>Diarthrodes</i>	<i>Diarthrodes</i> sp1	0.4	0.0
			<i>Diarthrodes</i> sp2	0.2	0.0
		<i>Paradactylopodia</i>	<i>Paradactylopodia brevicornis</i>	0.5	0.3
	Ectinosomatidae	<i>Ectinosoma</i>	<i>Ectinosoma</i> sp1	3.4	33.6
			<i>Ectinosoma</i> sp2	1.8	1.9
			<i>Ectinosoma</i> sp3	0.4	0.0
		<i>Pseudobradya</i>	<i>Pseudobradya</i> sp.	1.1	0.0
	Hamondiidae	<i>Ambunguipes</i>	<i>Ambunguipes</i> sp.	0.5	0.0
	Harpacticidae	<i>Harpacticus</i>	<i>Harpacticus obscurus</i>	5.6	0.0
			<i>Harpacticus</i> sp.	0.0	1.4
	Laophontidae	<i>Applanola</i>	<i>Applanola hirsuta</i>	0.0	0.6
		<i>Echinolaophonte</i>	<i>Echinolaophonte</i> sp.	0.0	0.2
		<i>Laophonte</i>	<i>Laophonte cornuta</i>	0.5	0.0
			<i>Laophonte parvula</i>	3.9	0.0
			<i>Laophonte</i> sp1	0.2	0.0
			<i>Laophonte</i> sp2	0.0	0.2
		<i>Paralaophonte</i>	<i>Paralaophonte brevis</i>	0.0	0.7
			<i>Paralaophonte congenera</i>	1.3	0.0
			<i>Paralaophonte</i> sp.	0.0	3.9
	Miraciidae	<i>Amonardia</i>	<i>Amonardia</i> sp.	0.4	0.0
		<i>Amphiascoides</i>	<i>Amphiascoides</i> sp1	5.6	0.4
			<i>Amphiascoides</i> sp2	0.7	1.5
			<i>Amphiascoides</i> sp3	0.2	0.0
		<i>Amphiascopsis</i>	<i>Amphiascopsis cinctus</i>	1.8	1.5
		<i>Amphiascus</i>	<i>Amphiascus (Minutus)</i> sp.	0.6	0.3
			<i>Amphiascus (Varians)</i> sp.	0.4	0.0
			<i>Amphiascus</i> sp.	5.3	0.0
		<i>Dactylopodamphiascopsis</i>	<i>Dactylopodamphiascopsis</i> sp.	0.5	0.0
		<i>Delavalia</i>	<i>Delavalia</i> sp.	0.5	0.0
		<i>Diosaccus</i>	<i>Diosaccus</i> sp.	0.5	0.0
		<i>Haloschizopera</i>	<i>Haloschizopera</i> sp.	0.0	0.4
		<i>Melima</i>	<i>Melima</i> sp1	3.2	0.8
			<i>Melima</i> sp2	2.9	0.0
			<i>Melima</i> sp3	0.2	0.0
		<i>Paramphiascella</i>	<i>Paramphiascella</i> sp.	4.4	0.8
		<i>Robertgurneya</i>	<i>Robertgurneya</i> sp.	1.4	6.2
		<i>Robertsonia</i>	<i>Robertsonia knoxi</i>	3.1	0.0
			<i>Robertsonia mourei</i>	1.0	0.0
			<i>Robertsonia</i> sp1	0.7	0.5
			<i>Robertsonia</i> sp2	1.4	0.9
		<i>Stenhelia</i>	<i>Stenhelia</i> sp.	1.3	0.0
	Parastenheliidae	<i>Parastenhelia</i>	<i>Parastenhelia spinosa</i>	13.1	0.5
	Peltidiidae	<i>Alteutha</i>	<i>Alteutha</i> sp.	3.4	0.0
		<i>Eupelte</i>	<i>Eupelte</i> sp.	0.5	2.4
	Pseudotachidiidae	<i>Xouthous</i>	<i>Xouthous</i> sp.	1.6	2.0
	Thalestridae	<i>Parathalestris</i>	<i>Parathalestris</i> sp.	0.4	0.0
	Tisbidae	<i>Scutellidium</i>	<i>Scutellidium</i> sp.	1.4	8.6

In the comparison between the control and impacted areas, significant differences were found for richness (Pseudo-F = 29.807; $p < 0.001$), Margalef diversity (Pseudo-F = 34.268; $p < 0.001$), evenness (Pseudo-F = 21.088; $p = 0.003$) and Shannon-Wiener diversity (Pseudo-F = 31.486; $p < 0.001$). For all of these indexes, the control areas demonstrated higher average values when compared to impacted areas (Figure 2).

The similarity between the replicates in the control and impacted areas can be seen in the NMDS representation (Figure 3). Table 3 shows the results of the SIMPER analysis. *Parastenhelia spinosa*, *Laophonte parvula* and *Amphiascoides* sp1 were the species that contributed most to the control reef similarity, whereas *Ectinosoma* sp1, *Ameira* sp2 and *Scutellidium* sp. were the species that contributed most to impacted reef similarity. The SIMPER average values of similarity between replicates of impacted areas (49.76) were found to be almost twice as large as those between replicates of the control areas (24.44), resulting in significantly different values (Student's *t*-test = 11.49; $p < 0.001$). Significant differences were found using a nested PERMANOVA for the comparison between the control and impacted areas, and among reefs within the control and impacted areas (Table 4). The PERMDISP analysis showed significant differences between the reefs included in this study ($F = 5.719$; $p = 0.047$) regarding homogeneity of dispersion, with control reefs showing greater dispersion. This pattern can be seen in the NMDS.

Table 3. Cumulative percentage (Cum. %) of species contribution to average similarity for the control and impacted areas and dissimilarity between the control and impacted area replicates.

Control		Impact & Control	
Average similarity: 24.44		Average dissimilarity = 86.52	
Species	Cum. %	Species	Cum. %
<i>Parastenhelia spinosa</i>	24.38	<i>Ectinosoma</i> sp1	17.47
<i>Laophonte parvula</i>	31.79	<i>Ameira</i> sp2	26.68
<i>Amphiascoides</i> sp1	38.84	<i>Parastenhelia spinosa</i>	33.59
<i>Harpacticus obscurus</i>	45.66	<i>Scutellidium</i> sp.	38.16
<i>Paramphiascella</i> sp.	51.53	<i>Robertgurneya</i> sp.	41.66
<i>Robertsonia knoxi</i>	56.79	<i>Dactylopusia tisboides</i>	44.89
<i>Dactylopusia tisboides</i>	61.13	<i>Harpacticus obscurus</i>	47.94
<i>Melima</i> sp1	65.33	<i>Amphiascoides</i> sp1	51
<i>Ectinosoma</i> sp1	69.4	<i>Ameira</i> sp1	53.99
<i>Amphiascus</i> sp.	73.37	<i>Amphiascus</i> sp.	56.95
<i>Alteutha</i> sp.	77.09	<i>Laophonte parvula</i>	59.55
<i>Ameira</i> sp1	79.39	<i>Alteutha</i> sp.	61.97
<i>Melima</i> sp2	81.61	<i>Paramphiascella</i> sp.	64.34
<i>Longipedia</i> sp.	83.66	<i>Paralaophonte</i> sp.	66.36
<i>Ameira parvula</i>	85.7	<i>Ameira parvula</i>	68.34
<i>Amphiascopsis cinctus</i>	87.49	<i>Melima</i> sp1	70.26
<i>Xouthous</i> sp.	89.18	<i>Robertsonia knoxi</i>	72.08
<i>Ameira</i> sp2	90.71	<i>Ectinosoma</i> sp2	73.75
		<i>Ameira</i> sp3	75.43
		<i>Dactylopusia</i> sp2	77.08
Impact		<i>Amphiascopsis cinctus</i>	78.53
Average similarity: 49.76		<i>Melima</i> sp2	79.98
Species	Cum. %	<i>Xouthous</i> sp.	81.39
<i>Ectinosoma</i> sp1	46.21	<i>Eupelte</i> sp.	82.76
<i>Ameira</i> sp2	69.18	<i>Longipedia</i> sp.	84.07
<i>Scutellidium</i> sp.	77.84	<i>Robertsonia</i> sp2	85.22
<i>Dactylopusia tisboides</i>	86.2	<i>Amphiascoides</i> sp2	86.26
<i>Robertgurneya</i> sp.	90.46	<i>Nannomesochra</i> sp.	87.25
		<i>Mesochra</i> sp2	88.23
		<i>Paralaophonte congenera</i>	89.11
		<i>Harpacticus</i> sp.	89.84
		<i>Stenhelia</i> sp.	90.53

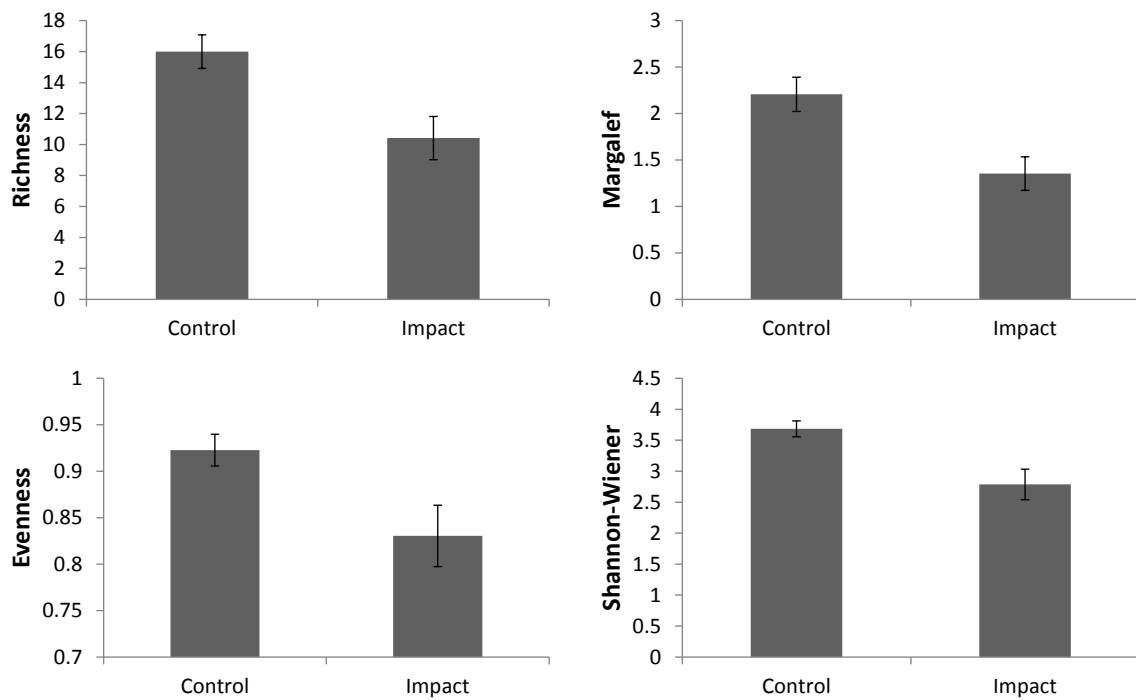


Figure 2. Average values of univariate bioindicators: species richness (S); Margalef diversity index (d); Pielou’s evenness index; and Shannon-Wiener diversity index (H’) in the control and impacted areas, based on the data from the harpacticoid copepods. Bars define confidence intervals of 95%.

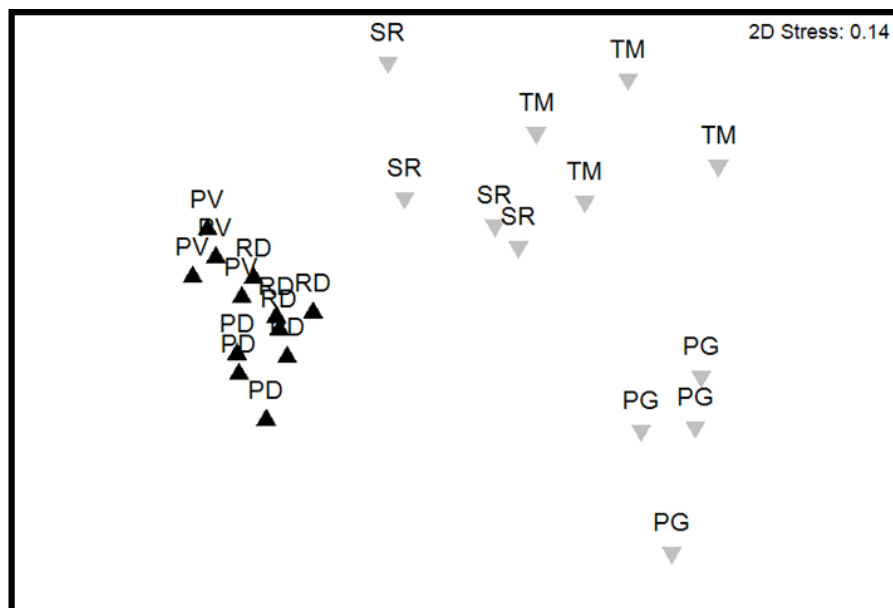


Figure 3. Non-metric multidimensional scaling (NMDS) ordination of *Harpacticoida* community structure. The control areas are represented by the blue symbols and impacted areas are represented by the green areas. Control areas (SR = Serrambi, PG = Porto de Galinhas and TM = Tamandaré) and impacted areas (PV = Paiva, PD = Piedade and RD = Rio Doce).

Table 4. The results from the multifactorial PERMANOVA analysis for the structure of the harpacticoid copepod community for factors Impact and Reef (within IMPACT).

Source	Df	SS	MS	Pseudo-F	P (MC)
Impact	1	21,768	21,768	3.6876	0.0097
Reef (Im)	4	23,612	5903	4.5396	0.0001
Res	18	23,406	1300.3		
Total	23	68,786			

The average values of the taxonomic distance indexes for each area can be found in Figure 4. In the comparison of taxonomic distance indexes using a Student’s t-test, only the Delta values showed significant differences between the control and impacted areas ($p = 0.0291$). Figure 5 shows the values of Delta+ for each of the sample reefs that were positioned within or very close to the 95% confidence interval funnel, with no deviations related to anthropogenic impact.

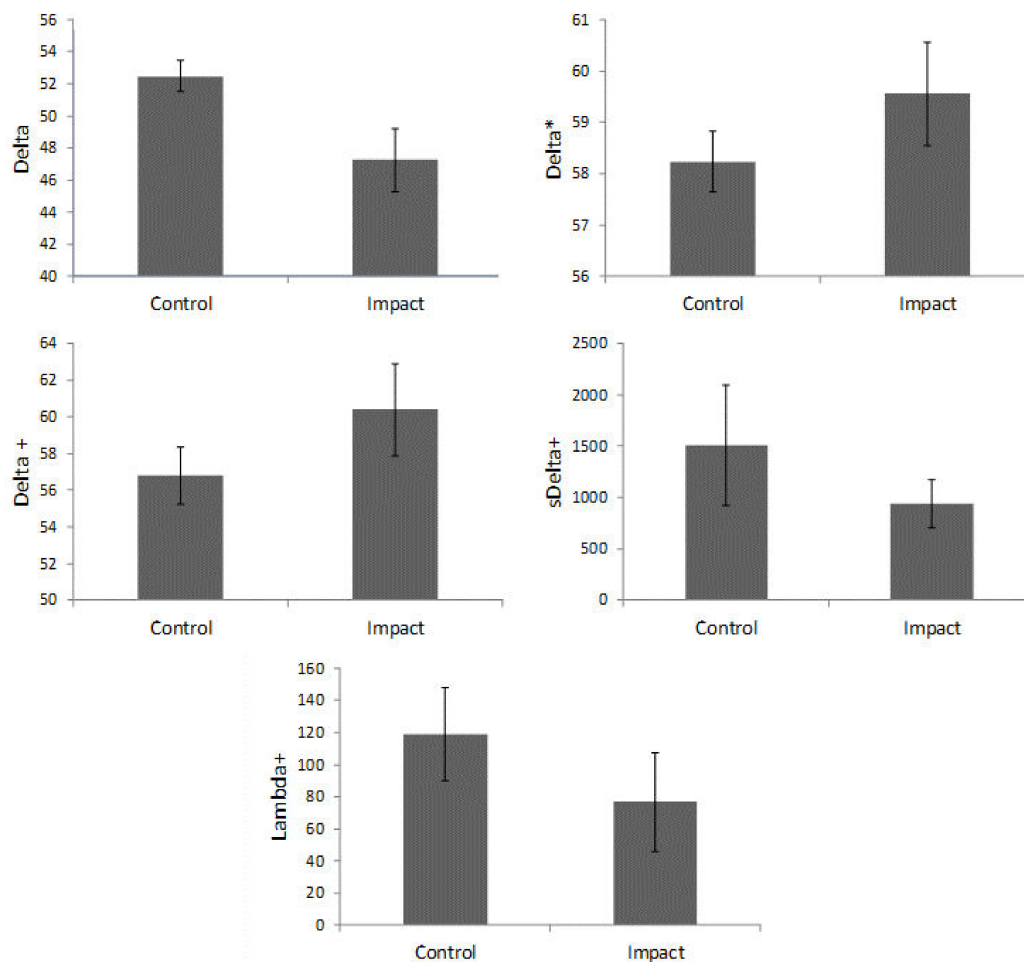


Figure 4. Average values of taxonomic diversity (Delta), taxonomic distance (Delta*), average taxonomic distance (Delta+), variation of taxonomic distance (sDelta+) and total taxonomic distance (Lambda+) for the control and impacted areas. Bars define 95% confidence intervals.

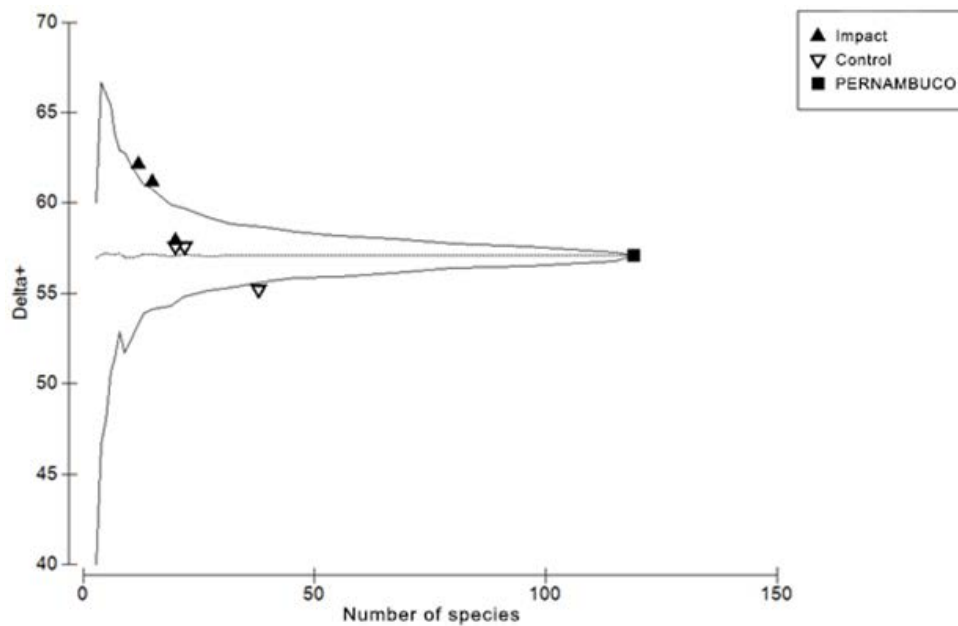


Figure 5. Average taxonomic distance values (Delta+) of six reefs in Pernambuco plotted against the number of species in each area with 95% confidence limits for random samples of subsets from a species in the regional list.

4. Discussion

Environmental variables are fundamental in the comparison and interpretation of meiofauna distribution patterns in areas exposed to different levels of anthropogenic impact [2]. In coral reef ecosystems, the water is generally considered oligotrophic [60,61], supersaturated with oxygen, and the nutrient concentrations are mainly influenced by atmospheric deposition, organic matter remineralization processes, sediment resuspension, or by riverine/estuarine input. The DIN, DIP and SiO₂ values registered for the control areas indicated minimal concentrations, which is expected for a non-impacted reef and allows higher levels of diversity [62–65].

However, for the a priori considered impacted areas, the urbanization levels and the consequent run-off of domestic and industrial wastewater are probably the main reasons for the higher nutrient concentrations [63]. The PERMANOVA analysis for nutrients demonstrated relevant differences between the control and impacted areas, supporting the a priori classification of the areas based on their geographical location and/or urbanization.

Despite the high nutrient concentrations in the impacted areas, the oxygen supersaturation due to the intense wave action upon the reefs, together with the high pH and TA values, indicate that the environments in this study do not suffer from hypoxia and/or coastal water acidification [66].

The synthetic grass ASU used in this study was recently tested in coastal reef environments along the coastline of northeast Brazil [52,53]. The results indicated that ASUs made of synthetic grass were an effective representation of the structure of meiofaunal communities in phytal environments in coral reef areas. Across the world, ASUs have been used to monitor the in situ change in biodiversity as a result of different sources of anthropogenic pollution [67], and have been found to be potentially useful as a method of biomonitoring coastal environments [23], a result that is herein supported.

In 1995, Warwick and Clarke [33] found a continuous decrease in the taxonomic distance of a marine community along a gradient of increasing estuarine contamination where species diversity remained constant, indicating a phylogenetic signal of the impact. However, Hall and Greenstreet [36] demonstrated that in demersal fish communities, TDIs (taxonomic distance indexes) exhibited the same behaviors as conventional diversity indexes when comparing disturbance levels in this environment. Somerfield et al. [68] also concluded that it was not possible to consistently observe

the decreasing pattern of TDI values for macrofauna, and found an increased impact in deeper regions. Thus, Salas et al. [55] tested the robustness of taxonomic distinction measures. They applied them to different scenarios of estuarine disturbance in Portugal and Spain and found that in the majority of case studies, among the various TDIs used, only the total index of taxonomic distinction was relatively satisfactory in discriminating situations of disturbance and contrast with the used indexes (Shannon-Wiener, Margalef and eco-exergy). In the present study, the classic indexes of ecological diversity clearly characterize the differences between the control and impacted areas. However, the taxonomic distinction measures, with the exception of the taxonomic diversity (Delta) index, did not show clear signs of impact differentiation, a finding which has been previously found in several of the case studies cited above (e.g., References [36,55,68]). These results suggest that the strong loss of harpacticoid copepod species in coastal reefs under the impact of multistressors does not present a significant phylogenetic signal.

The diversity indexes calculated for coral reef harpacticoid copepods (Margalef, evenness and Shannon-Wiener diversity) were found to be significantly greater in the control areas when compared to impacted areas. Furthermore, the community structure of harpacticoid copepods was also found to be highly sensitive to differences between the control and impacted areas based on PERMANOVA results. Mirto and Danovaro [23] found greater taxon richness in a meiofaunal community in control areas compared to impacted areas when investigating the dynamic colonization of meiofauna at the level of large taxonomic groups on artificial substrates made of nylon brushes in the port of Ancona, in the north of the Adriatic Sea. Other studies, using classic indexes, also found that anthropogenic stressors, such as fishing activities [69,70], mechanical disturbances [71], sedimentation increase [72], species invasions [73], temperature increase [74] and water pollution [75] had negative effects on reef environments. These stressors led to a decrease in richness and/or diversity of coral communities. Anthropogenic disturbances may create spatial and temporal variability in community structures in terrestrial, aquatic, and marine ecosystems [76,77]. Currently, the rate at which environmental conditions are changing introduces the concern of ecosystems becoming simplified, with less spatial variability [78], especially in extremely diverse tropical ecosystems, such as coastal coral reefs [79]. Our results indicate that changes in environments caused by anthropogenic multistressors cause the simplification of coral reef harpacticoid copepod communities, resulting in greater homogeneity within and between impacted areas and control areas. Similar results were found for benthic communities under the impact of seawater acidification [80,81].

In several previously studied areas, the simplification of communities driven by anthropogenic impacts has raised questions on how the diversity of an ecosystem influences its function. Recent studies affirmed that a greater number of species can lead to greater temporal stability of an ecosystem [82]. More than 50% of species identified in this study were exclusively present in the control areas, suggesting that this notable difference may be due to the reduction in harpacticoid copepod richness and could potentially affect the function of this ecosystem in impacted areas, a theme that deserves further attention.

In the control areas, the species with the highest densities were *Stenhelia* sp., *Parastenhelia spinosa* and *Amphiascoides* sp. The species that were found exclusively in the control area reefs were *Parastenhelia spinosa*, *Amphiascus* sp., *Laophonte parvula*, *Harpacticus obscurus*, *Ameira* sp. and *Ameira parvula*. In a study on the direct impact of human trampling in a reef area on the coast of Pernambuco, Sarmiento and Santos [31] also verified that *Parastenhelia spinosa* and *Ameira parvula* were some of the most abundant species in a protected area [8]. Furthermore, *Amphiascus* sp. (the same species as in the present study) was found to be one of the species found exclusively in the protected areas [31]. In this way, we conclude that environmental changes caused by anthropogenic multistressors cause the simplification of meiofaunal communities and the homogenization of ecosystems in impacted areas through decreases in richness. Furthermore, we can conclude that indicators based on taxonomic distinction are less efficient than classic indicators in the detection of anthropogenic impacts on harpacticoid copepod associations in coastal reef environments, since these impacts did not present a strong phylogenetic signal.

Author Contributions: P.J.P.S. conceived and outlined the study. M.S.B., B.J.d.S. and P.J.P.S. conducted the field experiment implantation and removal. M.S.B., B.J.d.S. and M.J.F.M. conducted chemical and biological analysis. M.S.B. and P.J.P.S. were responsible for data analysis and interpretation. M.S.B., B.J.d.S. and P.J.P.S. wrote the article.

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