

Is nematode colonisation in the presence of *Scolecipis* in tropical sandy-beach sediment similar to the colonisation process in temperate sandy beaches?

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Received September 2, 2011 – Accepted March 19, 2012 – Distributed February 28, 2013

(With 6 figures)

Abstract

The role of a dominant macrobenthic polychaete, *Scolecipis squamata*, in the colonisation of defaunated tropical sediments by sandy-beach nematodes was investigated and compared with a previous colonisation experiment carried out on a temperate sandy beach. Experimental cylinders, equipped with lateral windows allowing infaunal colonisation, were filled with defaunated sediment containing two treatments, with and without *S. squamata*. These cylinders were inserted into microcosms containing sediment with indigenous meiofauna collected from the field. The treatments were incubated in the laboratory at ambient temperature and salinity for 7, 14 and 21 days. The nematode assemblages in both treatments did not differ in composition between treatments and from the natural assemblages, suggesting that all the species were equally able to colonise the experimental cores. The presence of the polychaete did not affect the development of the nematode community composition, in contrast to the results from a previous temperate-beach experiment. However, our results did not indicate whether the difference in results was caused by the different behaviour of the polychaete specimens, or by the different composition and response of the present nematode community.

Keywords: meiofauna, *Scolecipis squamata*, Fazenda Beach, microcosm experiment, biological interactions.

A colonização dos sedimentos de praias tropicais por nematódeos na presença de *Scolecipis* é similar ao processo de colonização que ocorre em praias arenosas temperadas?

Resumo

Esse estudo avaliou a influência da espécie *Scolecipis squamata* (Polychaeta) no processo de colonização de sedimentos defaunados obtidos em uma praia arenosa tropical, além de comparar esses resultados com um estudo similar realizado em uma praia temperada. Sedimentos, previamente defaunados, foram colocados em amostradores com aberturas laterais, que permitiram a colonização da meiofauna. Foram definidos dois tratamentos, um com a presença e outro com a ausência da espécie *S. squamata*, além do controle. Os amostradores desses tratamentos foram alocados em unidades experimentais do tipo microcosmo, as quais continham sedimento com a meiofauna residente. Os tratamentos foram incubados em laboratório por 7, 14 e 21 dias, com condições controladas de temperatura e salinidade. Os nematódeos não apresentaram diferenças significativas em termos de composição entre os tratamentos e nem em relação ao controle, sugerindo que todas as espécies desse grupo foram, igualmente, capazes de colonizar as unidades experimentais. Esses resultados indicaram que a presença do poliqueto não afetou a estrutura da comunidade de nematódeos, o que representou um resultado contrário ao obtido para a praia temperada. No entanto, não se pode concluir se essas diferenças entre as praias estariam relacionadas ao comportamento diferencial de *S. squamata* ou pela presença de comunidades de nematódeos distintas nas praias.

Palavras-chave: meiofauna, *Scolecipis squamata*, Praia da Fazenda, experimento de microcosmo, interações biológicas.

1. Introduction

The presence of nematodes as a group in the sediment is independent of the sediment composition (Vanaverbeke et al., 2000), but, in general, nematodes are said to be highly dominant in sand finer than 300 µm (McLachlan and Brown, 2006). Several studies have largely demonstrated the importance of sediment characteristics, such as median grain size, silt content and sorting as key aspects structuring the composition and diversity of free-living nematodes (e. g. Wieser, 1959; Ward, 1973; Heip and Decraemer, 1974; Vincx, 1989; Vincx et al., 1990; Vanreusel, 1990; Vanaverbeke et al., 2002, 2011). Then, despite the geographic area or the tidal regime of a sandy beach, the sediment seems to be more important in structuring the nematode community. However, the three-dimensional sediment of sandy beaches is not exclusively inhabited by nematodes; diverse communities including species of different sizes of organisms, such as micro-, meio- (other than nematodes) and macrofauna are also found in sandy beach sediments. These organism group sizes generally interact with each other by means of trophic interactions (Schratzberger and Warwick, 1999; Tita et al., 2000; Aarnio et al., 2001) or non-trophic interactions (Van Colen et al., 2009; Braeckman et al., 2011; Maria et al., 2011) and, therefore, interactions between macrofauna and nematodes also influence the nematode composition of sandy beach sediments (Maria et al., 2011).

Among the macrofauna organisms from medium to very fine sandy beach sediments *Scolecipis squamata* Müller, 1806 (Polychaeta) is often very abundant in the North and South Atlantic, the North Pacific, the Indian Ocean and the Mediterranean Sea (Souza and Borzone, 2000). It can reach high abundances in the upper intertidal zone (Elliot et al., 1997; Degraer et al., 2003), the mid-tide level (Knott et al., 1983; Souza and Borzone, 2000; Janssen and Mulder, 2005) or in the subtidal (Knott et al., 1983; Hartmann-Schröder, 1996; Souza and Borzone, 2000). Recent morphometric studies have shown that this supposedly cosmopolitan species is, rather, a species complex, at least in the South Atlantic, and its members can be distinguished only by a detailed morphological analysis combined with electron microscopy (MacCord and Amaral, 2005; Rocha et al., 2009).

Polychaetes identified as *Scolecipis squamata* can modify the environment through their burrowing and deposit-feeding activities, and by producing pseudofaeces (Dauer, 1983; Pardo and Amaral, 2004; Van Hoey et al., 2004). In experimental treatments with fauna from an ultradissipative sandy beach at De Panne, North Sea, Belgium, *S. squamata* was able to facilitate the early establishment of two non-predatory nematode species by inhibiting the continuous colonisation of initially dominant opportunistic nematode species represented only by *Enoplolaimus litoralis* (Maria et al., 2011). However, the generality of these results is not yet established. While it cannot be excluded that individuals presently attributed to *S. squamata* in different parts of the world are actually different species, it can be assumed that they

do share similar behavioural characteristics. Therefore, by conducting nematode colonisation experiments with *S. squamata* in different regions of the world, it can be investigated whether the positive or negative effects of the presence of a dominant macrofaunal species established in temperate, macrotidal beaches (Maria et al., 2011) can be generalised to other beach types in other worldwide regions. Therefore, a laboratory microcosm experiment was set up to test the effect of *S. squamata* on the colonisation of azoic sediments by free-living nematodes in a tropical macrotidal sandy beach in Brazil. We specifically tested the null hypothesis that the nematode community in newly colonised sediments is unaffected by the presence of the polychaete, and our results were compared with previously conducted experiments on macrotidal sandy beaches in temperate regions (Maria et al., 2011). The results of this study will (1) increase the knowledge of macrofauna-meiofauna interactions in tropical sandy beaches, and (2) allow an assessment of the generality of these patterns across different sandy beaches.

2. Material and Methods

2.1. Study area and sediment sampling

Sediment from the upper 10 cm was collected from the upper intertidal level from Praia da Fazenda, a tropical, dissipative sandy beach (44° 48' W - 44° 52' W and 23° 20' - 23° 22' S) in the Parque Estadual da Serra do Mar, municipality of Ubatuba, São Paulo, Brazil (Figure 1) four weeks prior to the experimental set-up (15th May 2008). The intertidal area is approximately 3.5 km long, with a mangrove located at the north inlet edge.

2.2. Experimental set-up

Sediments previously collected were defaunated and made inorganic by burning to 500 °C for 4 h in a muffle furnace. One day before the experimental set-up (19th June 2008), triplicate field control (FC) samples were collected using Perspex corers (10 cm²) to a depth of 10 cm, in order to collect baseline information on the resident nematode community. Then, large volumes of sand were collected from the same area visited on 15th May. This sediment was homogenised in the field by successive rework and taken to the laboratory to fill the microcosm aquaria of 0.24 m². *Scolecipis squamata* individuals were sampled by sieving sediment from the upper intertidal level of the beach. They were kept alive until the experimental set-up in an aquarium filled with sand and oxygenated sea water.

Six microcosms, each consisting of a plastic aquarium (72 l), were filled with homogenised sediment inhabited by natural meiofaunal and macrofaunal sandy-beach communities to a depth of 12 cm. The sediments were left untouched for one day, to allow the community to stabilise. One corner (96 cm²) of the aquarium was kept free in order to place a water pump, silicone tubes and air stones in a plastic container, to avoid any disturbance of the sediment. Thirty-six experimental cylinders (10 cm²) were allocated to three types of treatment (see below) and were

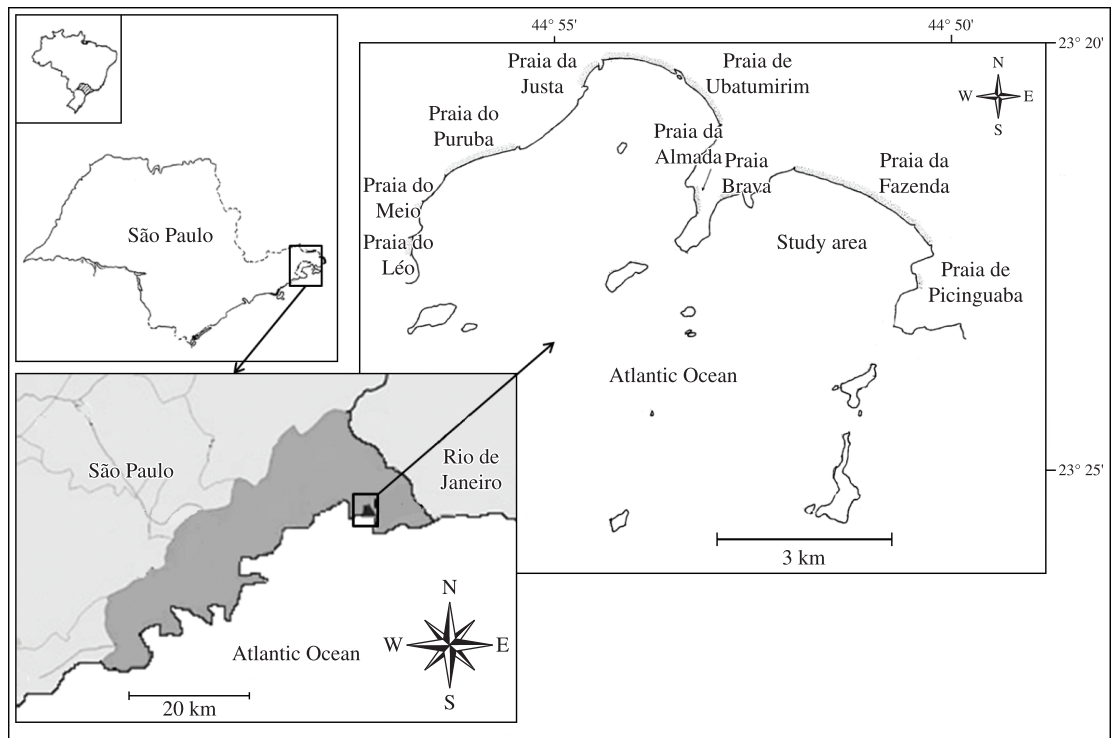


Figure 1. Study area localized in the Parque Estadual da Reserva do Mar at the southeast coast of Brazil.

randomly pushed into the sediment of different aquaria at a regular distance of 10 cm (Figure 2a). Before the addition of the experimental corers, a similar-sized corer removed the corresponding amount of sediment from the aquarium, in order to reduce disturbance. The experimental corers had two opposite lateral windows, each 5.3 cm × 2.6 cm, covered by gauze with a mesh size of 300 µm (Figure 2b). These lateral windows allowed meiofaunal migration from the adjacent sediment. The upper 2 cm of the gauze was in contact with the water, in order to allow mimicking of tides inside these corers; however, the water level never overtopped the edge of the corers.

Experimental cylinders with three different treatments were randomly distributed in the 6 aquaria. They consisted of:

1. Indigenous control (IC): natural sediment with indigenous community, collected simultaneously with the sediment used to fill the aquarium. This type of control was used to check for a possible effect of the use of the corer;
2. Azoic treatment (AT): defaunated sediment; and
3. *Scolecipis* treatment (ST): defaunated sediment + 5 specimens of *S. squamata*, which corresponded to the size of the polychaetes used in the North Sea experiment.

Immediately after the set-up of the experiment, the sediment in the aquarium was covered with 3.5 cm filtered seawater of natural salinity (35).

The experiment ran in a temperature-controlled room (25 °C) in a day and night light regime of 12: 12. Tides

were simulated twice a day, to a maximum water depth of 3.5 cm above the sediment layer. The water entered into the experimental cylinders through the upper 2 cm of the gauze, covering the lateral windows of the cylinder. The sediment was submerged for 2 hours and exposed to the air for 10 hours. Changes in salinity of the seawater were monitored daily, and increases due to evaporation were avoided by adding deionised water to the water reservoir, thereby maintaining the natural salinity.

Three replicates of each treatment were removed from different aquaria and transferred into a plastic container at 7, 14, 21 days post-placement during simulated low tide. At the same time, control samples (AQ) were randomly collected using a 10-cm² cylinder. Immediately after the removal of the experimental cylinders, the holes were filled with similar-sized empty corers to prevent the surrounding sediment from collapsing. All samples were preserved in a 10% formaldehyde solution until sample processing.

2.3. Sample processing in the laboratory

After the experiment ended, nematodes were extracted from the sediment by centrifugation with Ludox (Heip et al., 1985). Macrofauna was excluded by means of a 1-mm sieve. All organisms retained on a 38-µm sieve were counted and enumerated under a dissecting microscope. A sub-sample of 100 random nematodes were transferred to De Grisse solution (De Grisse, 1969) and mounted on slides for further identification to genus and species. Cylinders from the *Scolecipis* treatment were checked to assess if the organisms were still alive on the day of the sampling.

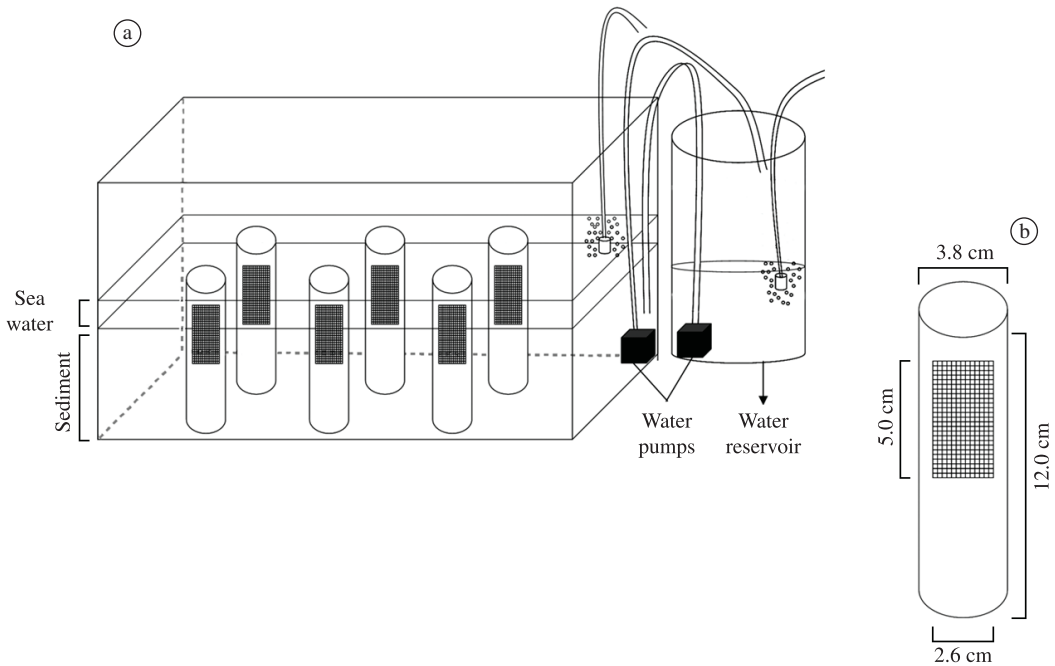


Figure 2. Schematic drawing of the experimental set-up. A: aerated microcosms (57 × 37 × 31 cm), B: syringes with lateral window of 5 cm × 2.6 cm which were filled with the different types of treatments.

2.4. Data analyses

Nematode assemblages from all the treatments and sampling dates were analysed using univariate and multivariate techniques. Total densities per 10 cm², species richness (S) and diversity (Shannon diversity index – H' log_e) were calculated for each treatment.

Differences between nematode densities of FC and AQ, and between FC and IC were analysed by t-tests. Differences in nematode densities among sampling times in AQ and IC were analysed by one-way ANOVA after checking that the necessary assumptions were met.

Experimental effects on total nematode density per 10 cm², species richness (S) and diversity (H') were tested by two-way analysis of variance (two-way ANOVA). When significant differences were detected, Tukey HSD tests were applied for testing for pairwise differences. Cochran's test was applied to check the homogeneity of the variances. Differences in nematode community structure were analysed by non-metric Multi-Dimensional Scaling (MDS) using the Bray-Curtis Similarity on non-transformed data for each sample. A one-way PERMANOVA was applied to analyse differences in the nematode community structure among FC, AQ and IC; and a two-way design was applied to evaluate differences in the community structure among treatments (AQ, IC, AT and ST) and over time (Anderson et al., 2008). Since a PERMANOVA test can show significant differences between groups, but does not distinguish between a difference due to factor effects or dispersion (variance), homogeneity of multivariate dispersion was tested with PERMDISP, using distances

among centroids calculated in the treatment × time group. The PERMDISP test was never significant, indicating equally dispersed distances to centroids. In case of a significant result in the PERMANOVA design, pairwise tests for the significant term were performed. In cases of restricted number of possible permutations in pairwise tests, p-values were obtained from Monte Carlo samplings (Anderson and Robinson, 2003). The species contributing most to within-group similarity were identified by the two-way crossed SIMPER analysis. All the multivariate analyses, and the calculation of S and H' were performed using the PRIMER v6 with PERMANOVA + add on software package (Clark and Gorley, 2006; Anderson et al., 2008), and the t-test and ANOVA were done using STATISTICA 7.0.

3. Results

3.1. Effect on density and diversity of the nematode communities

The nematode densities recorded in AQ samples were not significantly different from the values recorded from the field samples (time zero) (Figure 3; t-test, $t = -0.49$, $p = 0.65$) and did not change significantly over the course of the experiment (Figure 4, one-way ANOVA, $F_{2,6} = 0.94$, $p = 0.44$). Considering the indigenous controls, the densities were also not statistically different from the field samples (Figure 3; t-test, $t = 0.81$, $p = 0.46$). There was no significant change over time in the densities of IC over the course of the experiment (Figure 4, one-way ANOVA, $F_{2,6} = 0.63$, $p = 0.57$).

Nematode densities were not affected by treatment or by time, whereas species richness was significantly affected by the time \times treatment interaction term (Table 1, Figure 5). Tukey HSD (Table 1) indicated that these differences were caused by the higher species richness found in AQ at day-7 and in IC at day-14 and day-21. H' diversity was significantly affected by treatment only, since a higher diversity was found at IC (Table 1, Figure 5).

3.2. Effect on the nematode community composition

A mean of 16.983 individuals were counted in this study, belonging to 21 species and 2 unidentified genera. *Daptonema* sp. A, *Theristus* sp. A, *Theristus* sp. C, *Neochromadora* sp., and *Nudora besnardi* (Gerlach, 1956) were the dominant species in both the field control and experimental treatments (Appendix). The first three species are classified as non-selective deposit feeders and the latter two as epistrate feeders, according to Wieser (1953). No significant differences in nematode community composition were observed among FC, AQ, and IC treatments at the beginning of the experiment (one-way PERMANOVA: Pseudo- $F_{2,6} = 1.92$, $p = 0.057$). In addition, nematode communities from both AQ and IC did not change significantly over time (one-way PERMANOVA;

and Pseudo- $F_{2,6} = 1.30$, $p = 0.253$ and Pseudo- $F_{2,6} = 1.586$, $p = 0.198$, respectively).

Nematode community composition was significantly affected by treatment and time, but not affected by the interaction term (Table 2). The pairwise test showed that the nematode communities from AQ were different from AT, and the nematode communities present at day-7 were significantly different from the communities encountered at day-14 (Figure 6, Table 2).

The species contributing to the similarity within each treatment and time indicated by two-way crossed SIMPER are listed in Table 3. Within-group similarity in AQ and IC was mainly determined by *Daptonema* sp. A and *Nudora besnardi*, whereas *Daptonema* sp. A and *Theristus* sp. A were much more important in AT. The difference between day-7 and day-14 mainly occurred by the replacement of the dominant *Theristus* sp. C by *Theristus* sp. A on the latter sampling day (Table 3).

4. Discussion

4.1. Experimental set-up

Initial changes in nematode densities, diversity and community composition caused by manipulation of the

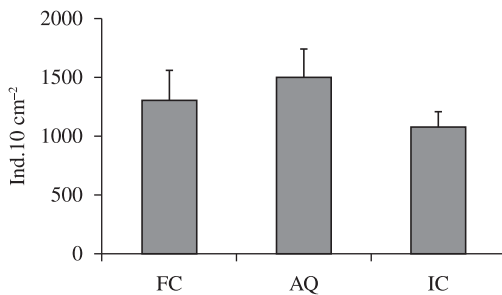


Figure 3. Mean nematode density in the field control ($t = 0$) and in the initial stages of the experiments ($t = 7$). FC: field control, AQ: aquarium control, IC: indigenous control. Error bar represents \pm SE ($n = 3$).

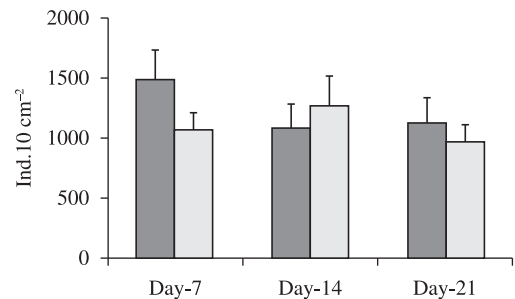


Figure 4. Mean nematode density in the controls over the 21 days of the experiment. Black bars represent AQ and grey bars represent IC. Error bar shows \pm SE ($n = 3$).

Table 1. Results from two-way ANOVA for the treatment and time effects on nematodes univariate measurements and overview of the Tukey HSD-test for number of species and Shannon diversity. Field controls are not included in this analysis.

ANOVA	Treatment		Time		Treatment \times Time	
	F _(3,24)	p	F _(2,24)	p	F _(6,24)	p
Density	0.79	0.507	2.68	0.089	2.06	0.096
Species richness (S)	23.18	<0.001*	0.25	0.805	3.18	0.019*
Diversity (H')	11.36	<0.001*	2.54	0.100	0.99	0.448
Groups compared		Result for S		Result for H'		
AQ \times IC		no difference		AQ < IC		
AQ \times AT		no difference		no difference		
AQ \times ST		day 7 (ST < AQ)		no difference		
IC \times AT		day 14, 21 (IC > AT)		IC > AT		
IC \times ST		day 14, 21 (IC > ST)		IC > ST		
AT \times ST		no difference		no difference		

AQ: aquarium control, IC: indigenous control, AT: azoic treatment, ST: *Scolecipis* treatment. *: significant values

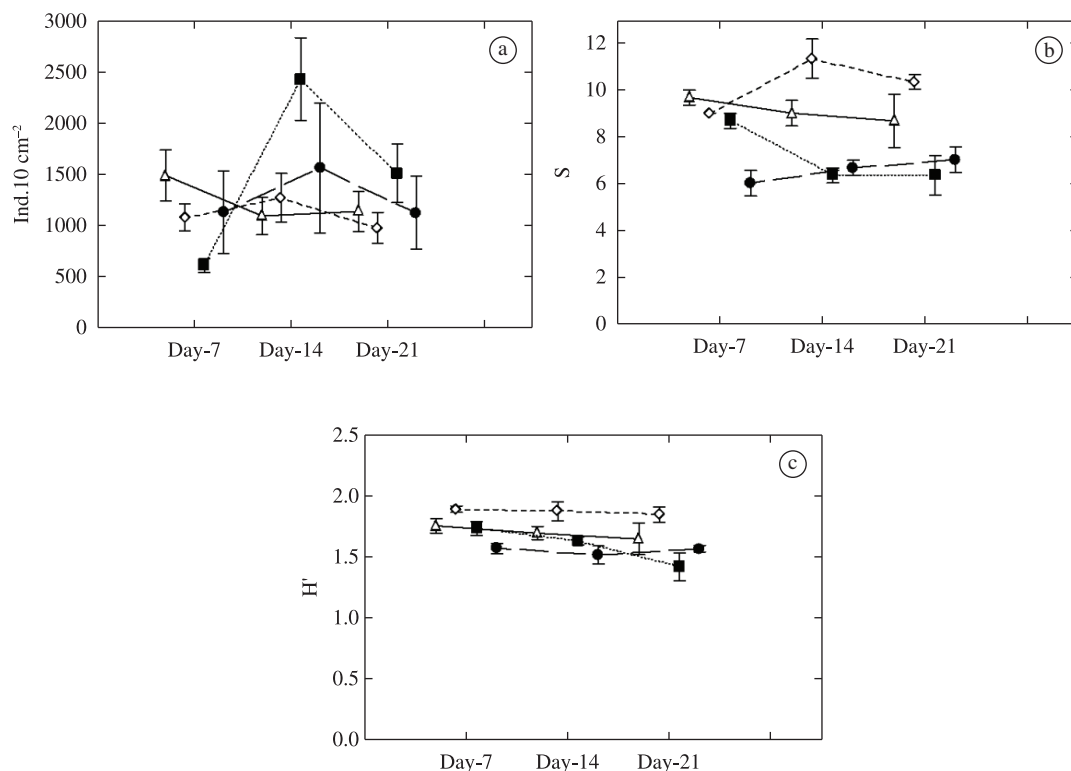


Figure 5. Univariate indices for nematode assemblages over the 21 days of the experiment. Treatment results (black symbols) were plotted against data obtained for the laboratory controls – AQ and IC – (open symbols) that served as potential species pool for colonisation of the defaunated sediment. a) mean total nematode density, b) species richness (S), c) Shannon diversity index (H'). Error bar represents $\pm SE$ ($n = 3$). AQ: triangles, IC: diamonds, AT: squares, ST: circles.

Table 2. Results from two-way PERMANOVA using Bray-Curtis similarity on non-transformed data showing the effect of treatment and time on nematode community and results from pair-wise tests using Bray-Curtis similarity on non-transformed, showing the treatment and time effect on nematode communities. Abbreviations as used in Table 1.

Factors	MS	Pseudo-F	p
Treatment _(3,24)	1382	2.06	0.022*
Time _(2,24)	2066	3.07	0.007*
treatment \times time _(6,24)	1109	1.65	0.067
Groups of treatments	T	p	
AQ \times IC	0.97	0.437	
AQ \times AT	1.84	0.014*	
AQ \times ST	1.44	0.100	
IC \times AT	1.92	0.020*	
IC \times ST	1.43	0.123	
AT \times ST	0.99	0.396	
Groups of days	T	p	
7 \times 14	1.91	0.026*	
7 \times 21	1.68	0.037*	
14 \times 21	1.65	0.055	

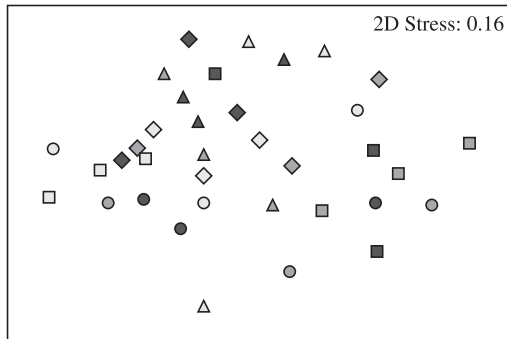
sediment could not be detected by our experimental design. In addition, no temporal changes were observed over the course of the experiment in AQ, again suggesting that using relatively large microcosms in meiofaunal experimental work avoids experimental artefacts (Maria et al., 2011). The absence of changes in density and nematode composition in the IC shows that cage effects did not occur in our experiment.

4.2. Colonisation pattern

A similar colonisation pattern was observed between azoic and *Scolecipis* treatments in terms of density, diversity and community structure. The nematode colonisation was a very rapid process, and all species had the same ability to colonise the newly available sediment since many of the species found in the source community (AQ) were found in the colonising cores (AT and ST). Previous colonisation experiments have shown that the success of a certain nematode genus, such as *Sabatieria* (Schratzberger et al., 2004), *Leptolaimus* (Gallucci et al., 2008) and *Enoplolaimus* (Maria et al., 2011), in colonising abundantly a new area is attributed to their relatively large body size. Our source community (AQ) lacked those large nematodes, mainly

Table 3. Output of two-way crossed SIMPER analysis showing the top 50% typical species for each treatment.

Species	Treatments				Days		
	Aquarium	Indigenous	Azoic	<i>Scolecopsis</i>	7	14	21
<i>Daptonema</i> sp. A	32	24	28	-	20	23	36
<i>Nudora besnardi</i>	23	23	-	25	22	26	-
<i>Theristus</i> sp. A	-	15	24	29	-	24	21
<i>Theristus</i> sp. C	-	-	-	-	21	-	-
Overall similarities	66	70	73	56	68	62	66

**Figure 6.** Non-parametric multi-dimensional scaling ordination based non-transformed species density using Bray-Curtis similarity comparing nematode community among lab controls (AQ: triangles, IC: diamonds) and treatments (AT: squares, ST: circles) over time (day-7: light grey, day-14: dark grey; day-21: black).

reflecting the granulometric characteristics of the studied beach since thin nematodes are more prone to occur in very fine sand (Fleege et al., 2011).

The dominance of Xyalidae and the low number of nematode species also reflect the sediment grain size of Fazenda beach. A high abundance of Xyalidae in sandy beaches composed of fine-grained sediments was already evidenced by Gheskiere et al. (2004), Hourston et al. (2005) and Gourbault and Warwick (1994). In addition, fine sediment has a low diversity of nematodes and is generally inhabited by non-selective deposit feeders (Vanaverbeke et al., 2011), which includes members of Xyalidae.

Although density, diversity and community structure were not significantly different, differences were found in terms of species richness. At day 7, a significant difference was found between ST and AQ, which may indicate that the chemical unattractiveness of the sediment is a barrier for the survival of successful rapid-colonising species (Maria et al., 2011). After day 7, there was a slight but significant rise in species richness in ST, indicating that environmental conditions had improved. Similar observations were made in temperate beaches (Maria et al., 2011) and were attributed to the recovery of microbial communities after defaunation (Stocum and Plante, 2006).

No significant differences in terms of density, diversity and nematode community composition were found between the azoic and the *Scolecopsis* treatment, in contrast to the findings obtained for a temperate, macrotidal sandy beach (Maria et al., 2011). In the latter study, a significant difference in the diversity between AT and ST was observed, attributable to the ability of *S. squamata* to inhibit the initially high colonisation rates of an opportunistic, large-sized predatory nematode, *Enoplolaimus litoralis* Schulz, 1936. A similarly prominent, predatory nematode was not present in the experimental and natural communities of the tropical Brazilian beach. Although *Nudora besnardi* and *Neochromadora* sp. both have an armed buccal cavity with sclerotised teeth and/or denticles, their buccal cavity is mainly adapted to pierce diatoms and/or scrape sand particles (Moens and Vincx, 1997). *Nudora besnardi* can also be relatively large in size (0.9–1 mm), but will never attain the large size of *E. litoralis* (1.4–2.5 mm), to which its high mobility and high colonisation rate in the beginning of the experiment was attributed (Maria et al., 2011). Therefore, the discrepancy between the results obtained here and the previous colonisation experiment may be a consequence of different nematode communities in both geographic areas which reflect directly the sediment composition of both areas.

The absence of interactions between nematodes and the polychaete could also be due to the species of polychaete used in our experiment. The species used here belongs to a species complex that includes *S. squamata*, *S. chilensis* and *S. goodbodyi* (Rocha et al., 2009). These species differ among each other in small morphological details, such as the shape of the notopodial lamellae and their fusion with gills (Rocha et al., 2009). Few studies have focused on the biology and ecology of these species (Hernandez et al., 2008; MacCord and Amaral, 2005, 2007), but our results might indicate that species that are so closely related can show different behaviours in the sediment. However, further testing is necessary to resolve the question of whether the divergent results between this experiment and the previous experiment with North Sea fauna (Maria et al., 2011) might be related to differences in the nematode community composition or to differences in the behaviour of, perhaps, different members of the *S. squamata*-group in the colonisation process.

Acknowledgements – We are thankful to Adriane Wandeness, Claudio Bloise and Márcio Macedo for their help during the

sampling and transport of the sediment, as well as Sérgio Bonecker and Ana Bonecker for providing the experimental room facilities of their laboratory. We thank Dr. Janet W. Reid, JWR Associates, for her critical revision of the English. The reviewer is also acknowledged for her comments in an earlier version of the manuscript. The first author was sponsored by a Ph.D. scholarship from Vlaamse Interuniversitaire Raad – VLIR-UOS. This paper contributes to FWO project G.0041.08N.

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Appendix

Appendix 1. Mean total density of the nematode species per 10 cm² within each treatment at each specific sampling time.

Species	Day-7					Day-14				Day-21			
	FC	AQ	IC	AT	ST	AQ	IC	AT	ST	AQ	IC	AT	ST
<i>Ascolaimus</i> sp.	0	0	0	0	0	0	13	0	0	0	0	0	0
<i>Bolbolaimus</i> sp.	0	0	0	0	0	0	5	0	0	0	0	0	0
<i>Cobbia</i> sp.	0	24	0	5	0	13	11	0	9	6	20	0	6
<i>Daptonema</i> sp.A	440	431	252	123	166	253	346	637	236	316	334	494	271
<i>Daptonema</i> sp.B	86	79	65	2	0	32	32	0	0	46	47	13	3
<i>Daptonema</i> sp.C	6	0	0	4	0	0	5	58	0	0	0	13	0
<i>Dichromadora</i> sp.	3	0	0	0	0	0	0	0	0	0	0	0	0
<i>Maryllynia</i> sp.	222	77	17	14	10	17	46	0	2	8	3	0	0
<i>Neochromadora</i> sp.	50	70	187	53	198	85	101	465	176	93	92	304	187
<i>Nudora besnardi</i>	243	250	199	156	409	286	471	389	314	392	197	89	149
<i>Odontophora</i> sp.	11	8	6	0	0	0	11	0	0	3	2	0	0
<i>Paracyatholaimus</i> sp.	3	0	0	0	0	7	0	0	23	3	39	0	5
<i>Pseudosteineria</i> sp.	0	3	8	8	2	0	13	0	0	0	0	6	6
<i>Pselionema</i> sp.	6	6	0	0	0	0	0	0	0	0	6	0	0
<i>Sabatieria</i> sp.	35	19	31	0	0	6	51	14	0	27	20	6	12
<i>Scaptrella</i> sp.	0	6	0	0	0	0	0	0	0	0	0	0	0
<i>Thalassironus</i> sp.	8	0	0	0	0	0	0	0	4	0	0	0	3
<i>Theristus</i> sp.A	141	179	158	108	192	225	308	552	571	143	105	434	372
<i>Theristus</i> sp.B	19	0	0	0	0	5	0	0	0	0	0	0	0
<i>Theristus</i> sp.C	31	329	149	127	148	151	193	310	219	94	103	147	106
<i>Theristus</i> sp.D	0	0	0	0	0	3	0	0	0	0	0	0	0
Non-identified genera	3	0	5	0	0	0	0	0	0	0	0	0	0

AQ: aquarium control, IC: indigenous control, AT: azoic treatment, ST: Scolelepis treatment.