

UNIVERSIDADE ESTADUAL PAULISTA "JÚLIO DE MESQUITA FILHO"
INSTITUTO DE BIOCÊNCIAS DE BOTUCATU
PROGRAMA DE PÓS GRADUAÇÃO EM CIÊNCIAS BIOLÓGICAS
ÁREA DE CONCENTRAÇÃO: ZOOLOGIA

DISSERTAÇÃO DE MESTRADO

PREVENDO MUDANÇAS CLIMÁTICAS: RESPOSTA DA MEIOFAUNA A
MUDANÇAS NA DISTRIBUIÇÃO ESPACIAL DE ESPÉCIES-CHAVE

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Botucatu-SP

-2016-

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**Previendo mudanças climáticas: resposta da meiofauna a mudanças na
distribuição espacial de espécies-chave**

Dissertação apresentada ao curso de pós-graduação em Ciências Biológicas: Zoologia, do Instituto de Biociências, Universidade Estadual Paulista (UNESP), Campus de Botucatu, como parte dos requisitos para a obtenção do título de Mestre em Ciências Biológicas – Área de Concentração: Zoologia.

Orientadora: Dra. Tânia Marcia Costa

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**Botucatu – SP
-2016-**

FICHA CATALOGRÁFICA ELABORADA PELA SEÇÃO TÉC. AQUIS. TRATAMENTO DA INFORM. DIVISÃO
TÉCNICA DE BIBLIOTECA E DOCUMENTAÇÃO - CÂMPUS DE BOTUCATU - UNESP BIBLIOTECÁRIA
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Citadin, Monica.

Previendo mudanças climáticas : resposta da meiofauna a mudanças na distribuição espacial de espécies-chave / Monica Citadin. - Botucatu, 2016

Dissertação (mestrado) - Universidade Estadual Paulista "Júlio de Mesquita Filho", Instituto de Biociências de Botucatu

Orientador: Tânia Marcia Costa

Coorientador: Sérgio Antonio Netto

Capes: ECOLOGIA

1. Mudanças climáticas. 2. Distribuição espacial da população. 3. Macroinvertebrados bentônicos. 4. Caranguejo. 5. Biodiversidade - Conservação. 6. Aquecimento global.

Palavras-chave: Deslocamento de espécies; Espécies chave; Meiofauna; Mudanças climáticas; Uca.

Dedico este trabalho à minha família, meu porto seguro: meus pais Moacyr e Ana, meu irmão André e minha cunhada Vanusa e à mais nova integrante da família, minha pequena Valentina.

Agradecimentos

Primeiramente, agradeço a Deus, pelo dom da vida, por ter me dado uma família maravilhosa, pela sua constante presença em minha vida, por iluminar meus caminhos e me dar sabedoria para concluir mais esta etapa.

Agradeço aos meus pais Ana e Moacyr pelo amor incondicional, por terem me mostrado caminhos do bem, pelo apoio constante em todas as minhas decisões e por não terem medido esforços para que eu e meu irmão tivéssemos tudo quanto podiam nos dar.

Ao meu irmão André e minha cunhada Vanusa pelo companheirismo, pelos momentos de descontração, por aturarem as minhas “chatices” nos momentos de estresse, e por terem trazido ao mundo a Valentina, minha alegria dos finais de semana.

Agradeço à minha orientadora Tânia, por aceitar que eu fizesse parte da sua “ganguê” de orientandos, por todos os conselhos e conhecimento transmitidos, pelos momentos de descontração nas festinhas e nas saídas de campo.

Ao meu co-orientador e amigo professor Sérgio, pela motivação a fazer ciência, por me apresentar ao mundo da meiobentologia, por dar aulas “alucinantes” e por ter me auxiliado em todos os momentos desta etapa.

Aos professores Ronaldo Christofolletti e Fabiane Gallucci por aceitarem fazer parte da banca examinadora deste trabalho.

À família Garcez que me adotou e amparou durante o tempo que morei em São Vicente, pelo companheirismo, amor e amizade que construímos.

Aos colegas do Laboratório de Ecologia e Comportamento Animal – LABECOMA da UNESP-São Vicente, pelo companheirismo e amizade, pela ajuda e conhecimentos compartilhados e pela zoeira sem limites.

À equipe do Laboratório de Ciências Marinhas da UNISUL, pela amizade, carinho e companheirismo, pela ajuda e conhecimentos compartilhados, pela zoeira e risadas infinitas e pelas tardes regadas a “cafezões”.

Às queridas Jessica e Jacqueline Kúcera, pela amizade incondicional e por visitarem minha mãe e manterem o clubinho da maldade mesmo durante o tempo que estive fora.

À minha segunda família, Tio Beto, Tia Dorinha, Tuane, Vózinha, Larissas e Gustavo, por todo o amor, carinho e companheirismo.

Aos meus queridos primos Samira e Vinicius, pelo amor e carinho, e pela paciência e compreensão por eu ter sido uma péssima prima mais velha nos últimos tempos.

A todos aqueles que de alguma forma contribuíram para que eu pudesse concluir mais esta etapa.

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1 APRESENTAÇÃO

2 O aumento das emissões de gases do efeito estufa, principalmente o dióxido de
3 carbono, metano e óxido nitroso desde a revolução industrial tem gerado aumentos
4 significativos na temperatura atmosférica. Esse aumento da temperatura está relacionado
5 a outras mudanças climáticas, tais como alterações nos padrões de precipitação e na
6 frequência de extremos climáticos (secas, inundações e tempestades severas, por exemplo),
7 além de aumento de temperatura na superfície dos oceanos, aumento do nível do mar, e
8 derretimento de geleiras (IPCC, 2014).

9 As alterações climáticas e mudanças físicas e geomorfológicas nos oceanos
10 podem afetar os ecossistemas costeiros, uma vez que mudanças nos padrões de
11 pluviosidade e no nível dos oceanos podem provocar variações de salinidade (Lee *et al.*,
12 2011) e aumento de sedimentação em zonas litorâneas (Ralston *et al.*, 2013). Áreas
13 costeiras de média e alta latitude, por exemplo, provavelmente enfrentarão mudanças na
14 biodiversidade devido à entrada de novas espécies oriundas de regiões adjacentes (Chen *et*
15 *al.*, 2011; Molinos *et al.*, 2015).

16 Modelos climáticos têm mostrado mudanças nos padrões de distribuição de
17 espécies de caranguejos do gênero *Uca*. Devido às alterações no clima, espécies de *Uca*
18 tenderão a ampliar sua distribuição espacial migrando em direção aos polos (Nabout *et al.*,
19 2009, tese). Atualmente, existem 11 espécies de *Uca* com ocorrência na costa Atlântica da
20 América do Sul (Bezerra, 2012). O limite máximo de distribuição destas espécies ao Sul é a
21 cidade de Florianópolis, porção central de Santa Catarina, com exceção da espécie *Uca*
22 *uruguayensis* que tem ocorrência até Mar Chiquita, província de Buenos Aires, Argentina e
23 *Uca mordax* que ocorre até Torres no Rio Grande do Sul (Thurman *et al.*, 2013).

24 Caranguejos do gênero *Uca*, conhecidos como caranguejos chama-maré, são
25 organismos típicos de áreas estuarinas e ocupam diferentes porções da zona entremarés

1 em regiões de clima tropical e subtropical (Crane, 1975). Caranguejos chama-maré são
2 considerados espécies chave uma vez que atuam como engenheiros de ecossistema
3 alterando as propriedades físicas e químicas do ambiente durante suas atividades de
4 escavação e alimentação (Payton *et al.*, 2002). Além disso, estudos recentes sugerem que
5 caranguejos *Uca* desempenham um papel fundamental controlando as comunidades
6 meiobênticas estuarinas (Ólafsson, 2003; Citadin *et al.*, 2016, no prelo).

7 Assim, no trabalho apresentado a seguir, buscamos avaliar os efeitos causados na
8 estrutura e funcionamento meiofauna devido à entrada de uma nova espécie chave na região
9 sul da América do Sul, como resposta às variações climáticas. Para isto, foram usadas duas
10 espécies de caranguejos chama-maré (*Uca uruguayensis* e *Uca leptodactyla*) em experimentos
11 em laboratório usando mesocosmos nos quais a densidade e a diversidade de *Uca* foram
12 manipuladas. Ao final dos experimentos, amostras de sedimento foram coletadas para análise
13 da meiofauna. Nossos resultados mostraram efeitos significativos na meiofauna devido ao
14 aumento de densidade e diversidade de caranguejos do gênero *Uca*.

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O trabalho a seguir será submetido à revista “Diversity and Distributions – A Journal of Conservation Biogeography” e, portanto, está formatado conforme as normas disponíveis no website da revista: [http://onlinelibrary.wiley.com/journal/10.1111/\(ISSN\)1472-4642](http://onlinelibrary.wiley.com/journal/10.1111/(ISSN)1472-4642).

1 **Projecting climate changes: response of meiofauna communities to shifts in spatial**
2 **distribution of keystone species**

3

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1 **ABSTRACT**

2 **Aim** Current trend of climate changes may affect biodiversity by shifting species distributions
3 patterns. *Uca* crabs, keystone species of intertidal estuarine sediments, are supposed to shift
4 their spatial distribution by poleward displacement as a response of global warming. We develop
5 and test models assessing whether the entrance of a new fiddler crab species, which directly
6 affects the diversity and abundance of keystone species, may affect the structure and
7 functioning of meiobenthic communities in intertidal areas.

8 **Location** The southern Atlantic coastal zone of South America.

9 **Methods** We used replacement series and partial additive design in indoor mesocosms
10 experiments, manipulating the diversity and density of keystone species. Experiments lasted for
11 20 days and then sediment samples were taken for meiofaunal analysis. The effect of keystone
12 species was quantified by structural and functional univariate descriptors and structure of
13 multivariate meiofaunal communities. We utilized permutational analysis of variance
14 (PERMANOVA) to assess differences in nematodes assemblages among treatments.

15 **Results** Nematodes were the most abundant meiofaunal group at both experiments. At the
16 replacement series experiment, the dominant genera of nematodes were *Microlaimus*,
17 *Terschellingia*, and *Anoplostoma*. Increasing keystone species diversity did not affect the
18 structure of nematodes assemblage, however it affected the functioning by reducing
19 predators/omnivores nematodes. *Microlaimus* was the dominant genus at the partial additive
20 design experiment. Increasing keystone species diversity and density affected the univariate
21 descriptors of structure and functioning of nematodes assemblage as well as the multivariate
22 structure. The total density of nematodes and density of *Microlaimus* were reduced with the
23 increase of *Uca* density. Increasing fiddler crabs density also led to changes in the multivariate
24 structure of nematodes assemblage. The increase of diversity of keystone species reduced non-
25 selective deposit feeders nematodes in the high level of *Uca* density.

1 **Main conclusions** Our findings suggested that the entrance of new species of *Uca* crabs in
2 the Southern of South America as a response of global warming may affect meiobenthic
3 communities. Increasing density and diversity of *Uca* may affect the structure and the
4 functioning of meiofauna due changes in top-down control.

5 **Keywords**

6 Global warming, nematodes assemblage, poleward displacement, *Uca*.

1 INTRODUCTION

2 The increase of anthropogenic emissions of greenhouse gases since the beginning
3 of industrial era has raised the atmospheric concentrations of carbon dioxide, methane and
4 nitrous oxide resulting in higher air temperatures. Each of the last three decades has been
5 successively warmer than any preceding decade since 1850. The period from 1983 to 2012
6 was likely the warmest 30-year period of the last 1400 years (IPCC, 2014).

7 Climate variability had altered marine environments. Among other effects general
8 effects, warming temperatures in the atmosphere led to a decrease in permafrost and sea ice
9 extent (Lawrence *et al.*, 2008), increase in sea surface temperature (Levitus *et al.*, 2008) and rise
10 in sea level (Church & White, 2006). Regional effects also have been described, such as the
11 current increase in rainfall along southwest Atlantic marine ecoregions (Bernardino *et al.*, 2015),
12 change in salinity along estuaries (Lee *et al.*, 2011) and increase of sedimentation in coastal areas
13 (Ralston *et al.*, 2013). Against such stressful conditions, species are adapting to the new
14 scenarios by shifting their phenology or changing their spatial distribution (Parmesan & Yohe,
15 2003).

16 Investigating changes in spatial distribution of species and interactions are key
17 issues to predict the possible consequences of climate change (Van der Putten *et al.*, 2010).
18 Climate changes have increased the occurrence of invasive species, since changes in abiotic
19 factors (e.g. temperature) increase the likelihood of species migrate to areas where the
20 conditions are favorable, which could, in turn, affect local species and ecosystems (Walther *et*
21 *al.*, 2009; Burgiel & Muir, 2010). The entrance of a new species in an ecosystem may significantly
22 affect native species. Non-native species may compete with native species (Mooney & Cleland
23 2001) and alter the availability of resources, causing suppression or enhancement of the relative
24 abundance of native species (Didham *et al.*, 2005) and thus affect species evenness.

25 Coastal areas of medium and high latitudes will probably face changes in
26 biodiversity caused by the entrance of new species migrating from adjacent regions due to

1 climate warming (Chen *et al.*, 2011; Molinos *et al.*, 2015). This is the case of *Uca* crabs, an
2 Ocypodidae keystone species, that is predicted to change their range of distributions next years,
3 migrating poleward (J.C. Nabout *et al.*, unpublished data). Keystone species are those whose
4 effects are disproportionately large relative to their abundances (Power *et al.*, 1996). *Uca* crabs,
5 known as fiddler crabs, are keystone species as they modify the physical and chemical
6 environment during their burrowing and feeding activities (Payton *et al.*, 2002). These crabs also
7 play a key role in controlling estuarine meiofauna (Ólafsson, 2003; Citadin *et al.*, 2016 *in press*).
8 Meiofauna is a group of small metazoans between 45 and 500 μm in size and constitutes the
9 most abundant and diverse group of benthic marine invertebrates. Meiofaunal organisms
10 contribute to the transfer of matter and energy by feeding on detritus, bacteria, microalgae and
11 other microorganisms, and being a food source for higher trophic levels (Giere 2009; Schüffel
12 *et al.* 2013).

13 Here we focus our research on two abundant species of fiddler crabs, *Uca*
14 *uruguayensis* and *Uca leptodactyla*, to investigate if changes in distributions of keystone species
15 due climate changes may affect interactions in intertidal areas, by means of meiofauna. Fiddler
16 crabs are conspicuous intertidal organisms from tropical and subtropical regions. Along the
17 Atlantic coast of South America, *U. leptodactyla* ranges from Turiamo, Aragua State, Venezuela
18 to Florianopolis, Santa Catarina State, Brazil meanwhile *U. uruguayensis* range of distribution is
19 from Cabo Frio, Rio de Janeiro State, Brazil to Mar Chiquita, Buenos Aires Province (Fig. 1)
20 (Thurman *et al.*, 2013). The region of the Cape of Santa Marta at Laguna city, State of Santa
21 Catarina, South Brazil, is a biogeographic transition zone. This region is the southern limit of
22 mangroves along American continent (Schaeffer-Novelli *et al.*, 2000). The State of Santa Catarina
23 is also the southern limit of distribution of many species of fiddler crabs, with *U. rapaz*, *U.*
24 *leptodactyla*, *U. burgersi* and *U. thayeri* occurring up to the central portion of the State.
25 Southwards, only *U. uruguayensis* and *U. mordax* occurs (Thurman, 2013).



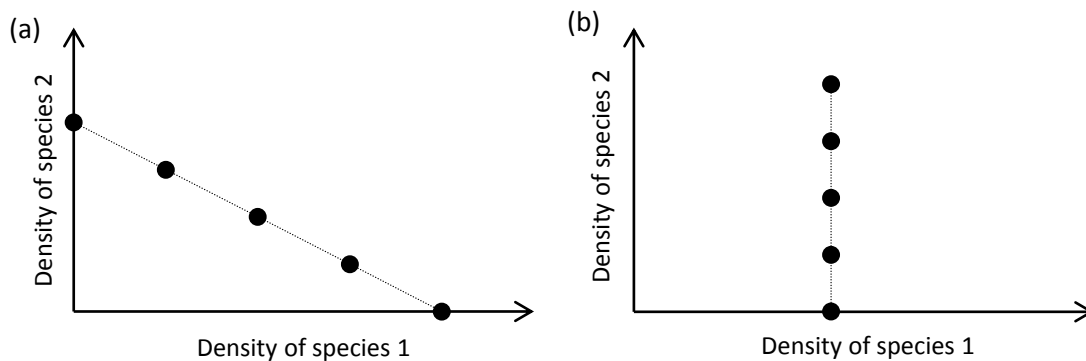
1
2 Figure 1 South America distribution of the fiddler crabs *Uca uruguayensis* and *Uca leptodactyla* according
3 to Thurman *et al.* (2013).

4
5 Considering the putative trend of *Uca* species to expand southwards, and their
6 strong top-down regulation on meiofauna (e.g. M.A. Fortuna *et al.*, unpublished data; Citadin *et*
7 *al.*, 2016 *in press*), we develop and test models assessing whether the entrance of a new fiddler
8 crab species, which directly affects the diversity and abundance of local keystone species, may
9 affect the structure and functioning of meiobenthic communities in intertidal areas.

10 11 **METHODS**

12 We established two mesocosm setups to mimic the putative effects of the arrival
13 of a new keystone species on meiofaunal communities. The experimental designs were based
14 on the replacement series from De Wit (1960) and the partial additive design from Harper (1977)
15 (Fig. 2). These designs differ in the way species are manipulated, in the replacement series the
16 total density of species is held constant in single and multiple-species treatments while the
17 proportion of each species is changed. In the partial additive design, density of one species is

1 kept constant while other varies. The arrival of a new keystone species leads to increases in
 2 species density and diversity. Thus, firstly, to test the hypothesis of increase in keystone species
 3 diversity due shifts in distribution, as a result of climate change, may affect meiofaunal
 4 communities, we used the replacement series. Then, we tested the effects on meiofauna due
 5 changes in keystone species density together with diversity in a partial additive design.



6
 7 Figure 2 Experimental models used in this study: (a) in the replacement series, total density is constant
 8 and the proportion of each species changes; (b) in the partial additive design, the total density varies as
 9 density of one species is constant and other varies.

10

11 **Set-up Experiment 1 – Replacement series**

12 The replacement series experiment was performed between the 27th of October
 13 and the 16th of November 2014. Total density of keystone specimens was fixed in 8
 14 inds/mesocosm (natural field densities may reach 240 crabs/m² (Masunari, 2006), *i.e.* 8,15
 15 inds/mesocosm). The experiment included four treatments - one with a single keystone species
 16 and three with two keystone species. The single-species treatment was composed by 8 *U.*
 17 *uruguayensis* (8U), and two-species treatments comprised: 6 *U. uruguayensis* + 2 *U. leptodactyla*
 18 (6U+2L); 4 *U. uruguayensis* + 4 *U. leptodactyla* (4U+4L) and 2 *U. uruguayensis* + 6 *U. leptodactyla*
 19 (2U+6L) (Fig. 3). Each treatment was replicated four times.

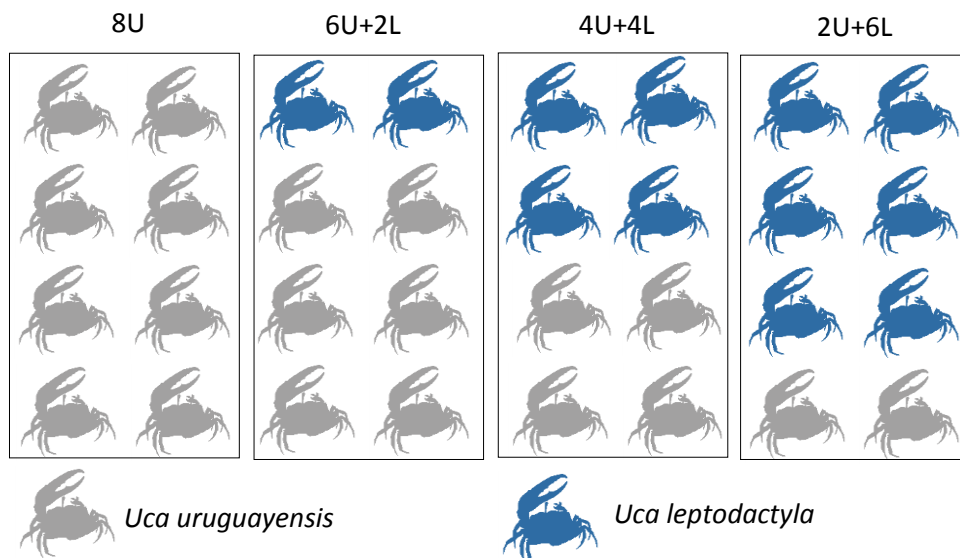
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21 **Set-up Experiment 2 – Partial additive design**

22 The partial additive design was performed between the 26th of January and the 15th
 23 of February of 2015. In this experiment, both keystone species density and richness were

1 manipulated. First, densities of *U. uruguayensis* were kept constant (two individuals) and *U.*
 2 *leptodactyla* were added two by two through the treatments to mimic the entrance of this
 3 fiddler crab in temperate zones due to climate change. So that, these treatments were
 4 composed by: 2 *U. uruguayensis* + 2 *U. leptodactyla* (2U+2L); 2 *U. uruguayensis* + 4 *U.*
 5 *leptodactyla* (2U+4L) and 2 *U. uruguayensis* + 6 *U. leptodactyla* (2U+6L). Besides, single-species
 6 treatments composed only by *U. uruguayensis* were also included with the same levels of total
 7 density than two-species treatment (4U; 6U and 8U) (Fig. 4) to test if the potential effects on
 8 meiofauna were due by the increase of richness or density of the keystone species.

9



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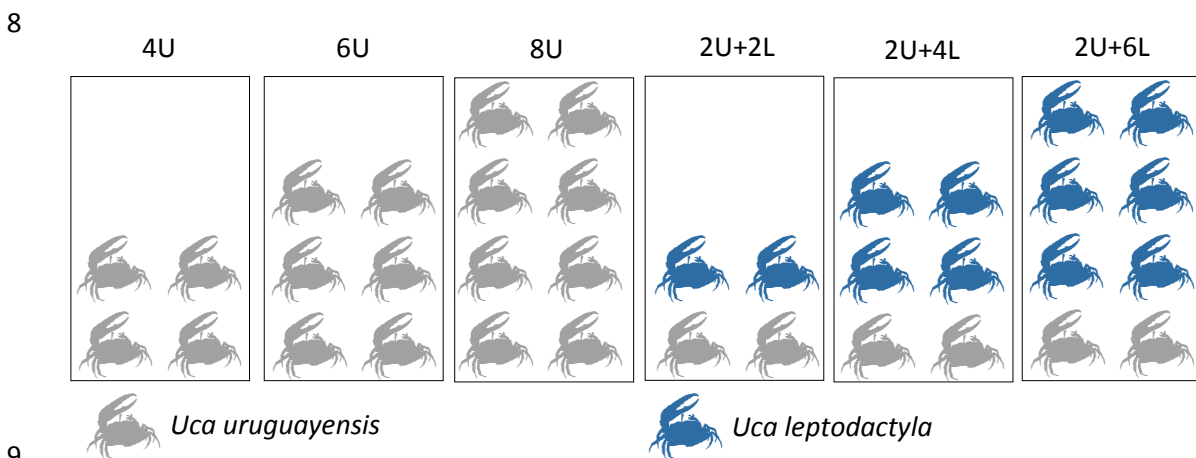
11 Figure 3 Set-up experiment 1 – replacement series. U: *Uca uruguayensis*; L: *Uca leptodactyla*.

12

13 **Experimental conditions**

14 In both experiments, *Uca* crabs were maintained in indoor mesocosms consisting
 15 in glass aquariums (29.5x11.5x20 cm, length x width x height). Sediments (upper than 5 cm) and
 16 fiddler crabs were collected on an unvegetated sandflat along the Una estuary, south-eastern
 17 Brazil (24°24'50"S, 47°04'14"W), at the Juréia-Itatins Reserve, part of a network protected areas
 18 of Atlantic rainforest (Marques & Duleba 2004), and transported to the laboratory inside plastic
 19 boxes. Only adult males of the two crabs species (from 8 to 10 mm of carapace width) were used
 20 in both experiments. Sediment were composed by very fine sand with an average total organic

1 matter of 1.52% (L.F. Natálio *et al.*, unpublished data), and chlorophyll *a* content ranging from 0
 2 to 11.98 $\mu\text{g g}^{-1}$ (Citadin *et al.*, 2016 *in press*). In laboratory, sediments were gently homogenized,
 3 and subsamples for the mesocosms were taken up to complete a layer of 5 cm of sediment.
 4 Fiddler crabs were included in mesocosms only 48 h after sediments to permit meiofauna
 5 stratification and crabs acclimation at laboratory. The experiments ran with constant
 6 temperature (28° C). By the end of the experiments, crabs were counted and sediment samples
 7 were collected from each mesocosm to evaluate the changes in meiofauna.



9
 10 Figure 4 Set-up experiment 2 – Partial additive design. U: *Uca uruguayensis*; L: *Uca leptodactyla*.

11

12 Sampling and sample processing

13 A total area of 19.23 cm² of sediment was taken from each mesocosm from the
 14 experiment 1, (two PVC corers of 3.5 cm in diameter to a depth 1 cm), and 28.84 cm² for the
 15 experiment 2 (three PVC cores of 3.5 cm x 1 cm). Differences in volume of sampled sediment
 16 were due to the difference in density of meiofauna inside the mesocosms, lower in experiment
 17 2. All sediment samples were fixed in 4% formalin and processed following Somerfield *et al.*
 18 (2005). Sediment was washed in fresh water, sieved through 500 and 63 μm mesh openings, and
 19 fauna on the smaller mesh was extracted by flotation in Ludox TM-50 (specific gravity of 1.15).
 20 The extracted fauna was placed in embryo dishes with glycerol (65% water, 30% alcohol and 5%
 21 glycerin) and left to evaporate for about 10 h, then permanent slides were made.

22

1 **Data analysis**

2 Two replicates, one from 2U+4L treatment and other from 2U+6L treatment of
3 experiment 2 had crabs died by the end of the experiment, and were excluded from statistical
4 analysis. For data analysis, only nematodes were used as they accounted for 97% of the total
5 meiofauna. Both univariate and multivariate statistical methods were used to test the effects of
6 fiddler crabs density and/or diversity on nematodes community. Permutational analysis of
7 variance (PERMANOVA; Anderson *et al.*, 2008) were used to assess differences in nematodes
8 univariate descriptors and multivariate structure among treatments. PERMANOVA was chosen
9 because it allows for mixed designs, an interaction term, and does not assume a normal
10 distribution of errors.

11 In univariate analysis both structural and functional nematodes descriptors were
12 used. Structural descriptors included total density, number of genera, Shannon-Wiener diversity
13 (\log_2) and the density of numerically dominant genera. Yet as functional attributes of the
14 nematodes, we used the relative abundance of each feeding type (Wieser, 1953; 1960), the
15 index of trophic diversity (Heip, 1985) and the maturity index (Bongers *et al.*, 1991).

16 For the experiment 1, univariate one-way PERMANOVA tests were run on Euclidean
17 distance matrices with 999 permutations and with unrestricted permutation of raw data.
18 Multivariate one-way PERMANOVA tests were run on Bray-Curtis similarity matrices. For both
19 analysis, when significant differences were detected ($p < 0.05$), pairwise tests P-value based on
20 Monte Carlo (MC) were applied. Ordinations of the multivariate faunal data were represented
21 by non-metric multidimensional scaling (nMDS).

22 For the second experiment two-way PERMANOVA were used to assess differences
23 in univariate and multivariate faunal structure among keystone species' density (fixed factor
24 with three levels: 4, 6 and 8 crabs) and richness (fixed factor with two levels: *U. uruguayensis*
25 and *U. uruguayensis* + *U. leptodactyla*, nested in keystone species density). Univariate
26 PERMANOVA tests were run on Euclidean distance matrices with 999 permutations, and the

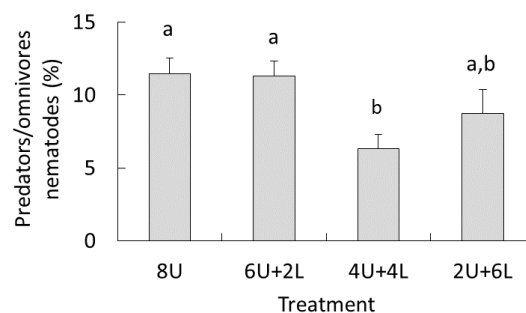
1 residuals were permuted under a reduced model. PERMANOVA tests for differences in the
 2 multivariate community structure were carried out on Bray-Curtis similarities matrices. When
 3 significant differences were detected ($p < 0.05$), pairwise tests P-value based on Monte Carlo
 4 (MC) were applied. Ordinations of the multivariate faunal data were represented by non-metric
 5 multidimensional scaling (nMDS).

7 RESULTS

8 Experiment 1 - Replacement series

9 Nematodes accounted 97% of the total individuals, so that other meiofaunal groups
 10 were excluded from statistical analyses. Nematodes densities ranged from 36 to 126 inds/10
 11 cm^2 , distributed in 25 genera belonging to 14 families (Appendix 1). The genera *Anoplostoma*
 12 (*Anoplostomatidae*), *Microloaimus* (*Microloaimidae*) and *Terschellingia* (*Linhomoeidae*) were the
 13 numerically dominant genera contributing for 25,1%, 19,9% and 18% (respectively) of the total
 14 fauna.

15 The structural and functional descriptors of nematodes did not vary significantly
 16 with the increase in keystone species diversity (Table 1). The exception was the relative
 17 abundance of predators/omnivores nematodes ($p(\text{MC}) = 0,037$). The pairwise PERMANOVA test
 18 showed difference between the groups of treatments 6U+2L and 4U+4L; and 8U and 4U+4L
 19 (results of pairwise tests are on Appendix 2), predators/omnivores nematodes were lower in the
 20 4U+4L treatment (Fig. 5).



21
 22 Figure 5 Mean values (\pm SE) of predators/omnivores nematodes on treatments. U: *U. uruguayensis*; and
 23 L: *U. leptodacyla*. Different letters mean differences found on pairwise tests.

1 Non-metric multidimensional scaling showed that the multivariate structure of
 2 nematodes did not differ between the treatments since no clear pattern of aggregation of the
 3 groups can be observed (Fig. 6). Multivariate PERMANOVA also showed no difference between
 4 treatments ($p(\text{MC})=0,263$; Table 1).

5

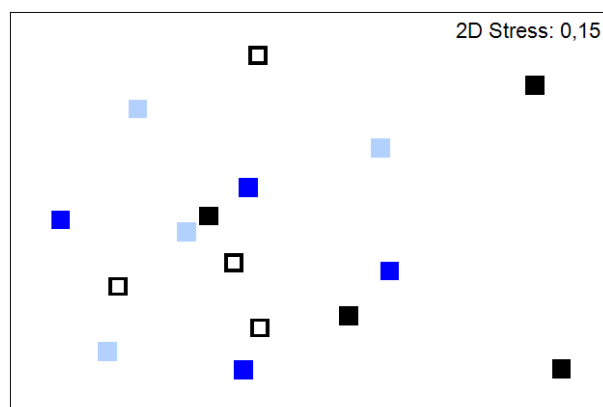
6 Table 1 Results of PERMANOVA evaluating the increase in keystone species diversity on multivariate
 7 structure and on structural and functional univariate descriptors of nematodes. Bold values indicate
 8 $p \leq 0.05$. Pairwise tests are available on Appendix 2. df: degrees of freedom; SS: sum of squares; MS: mean
 9 squares; $p(\text{MC})$: p-value obtained with Monte Carlo permutation test.

Source of variation	df	SS	MS	F	$p(\text{MC})$
Multivariate structure					
Treatment	3	1152,8	384,26	1,2007	0,263
Residual	12	3840,4	320,04		
Structural descriptor					
Density					
Treatment	3	3303,7	1101,2	1,5694	0,253
Residual	12	8420,1	701,68		
Number of genera					
Treatment	3	10,5	3,5	0,7433	0,568
Residual	12	56,5	4,7083		
Diversity (H')					
Treatment	3	0,0151	0,0050	0,8040	0,526
Residual	12	0,0753	0,0062		
<i>Anoplostoma</i> density					
Treatment	3	155,83	51,943	0,7309	0,575
Residual	12	852,77	71,065		
<i>Microlaimus</i> density					
Treatment	3	278,87	92,957	1,5802	0,243
Residual	12	705,94	58,828		
<i>Terschellingia</i> density					
Treatment	3	301,93	100,64	2,3724	0,128
Residual	12	509,07	42,422		
Functional descriptor					
Selective deposit feeders					
Treatment	3	279,56	93,187	1,544	0,236
Residual	12	724,23	60,353		

1 Table 1 Results of PERMANOVA evaluating the increase in keystone species diversity on multivariate
 2 structure and on structural and functional univariate descriptors of nematodes. Bold values indicate
 3 $p \leq 0.05$. Pairwise tests are available on Appendix 2. df: degrees of freedom; SS: sum of squares; MS: mean
 4 squares; p(MC): p-value obtained with Monte Carlo permutation test.

Source of variation	df	SS	MS	F	p(MC)
Non-selective deposit feeders					
Treatment	3	116,44	38,814	1,1044	0,379
Residual	12	421,72	35,144		
Epistratum feeders					
Treatment	3	134,49	44,829	0,8939	0,496
Residual	12	601,78	50,148		
Predators/Omnivores					
Treatment	3	71,569	23,856	4,0687	0,037
Residual	12	70,36	5,8633		
Index of Trophic Diversity					
Treatment	3	0,0030	0,0010	2,0308	0,158
Residual	12	0,0059	0,0004		
Maturity Index					
Treatment	3	0,0069	0,0023	0,2092	0,872
Residual	12	0,1321	0,0110		

5



6

7 Figure 6 MDS ordination for $\log(x+1)$ transformed nematodes multivariate structure. Black squares 8U;
 8 dark blue: 6U+2L; light blue: 4U+4L and open squares 2U+6L. U: *U. uruguayensis*; and L: *U. leptodactyla*

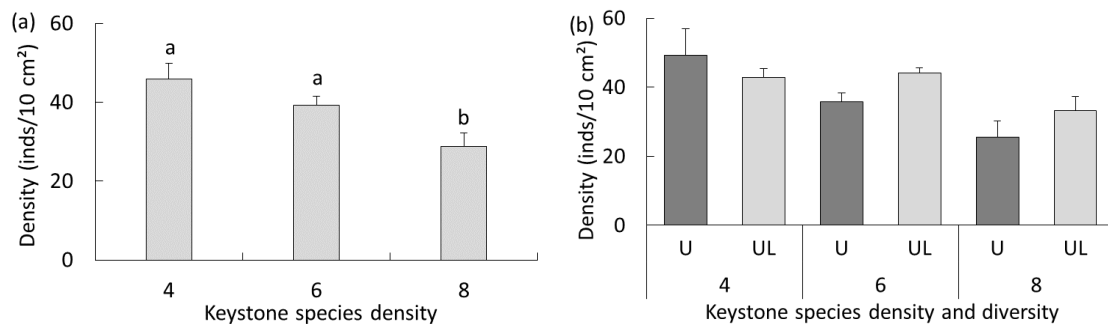
9

10 Experiment 2 – Partial additive design

11 As in the replacement series experiment, nematodes were the dominant
 12 meiofaunal group and account more than 95% of the total fauna, so that, only nematodes were
 13 used on statistical analysis. Nematodes densities ranged from 19 to 64 inds/10², distributed in

1 18 genera belonging to 14 families. (Appendix 3). The genus *Microlaimus* (Microlaimidae) was
 2 the numerically dominant genera contributing for 82,1%, of the total fauna.

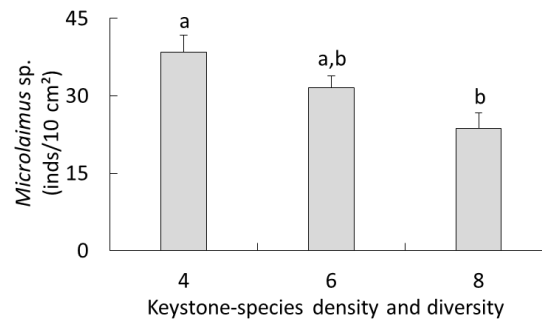
3 The total density of nematodes and the density of *Microlaimus* sp. were affected
 4 by the increase of keystone species density (both $p(\text{MC})= 0,017$) but not by the increase of
 5 keystone species richness ($p(\text{MC})=0,331$ and $0,0333$, respectively) (Table2). For total density of
 6 nematodes, the pairwise test showed this descriptor varied between the levels of crabs' density
 7 4 and 8, and 6 and 8 but not between 4 and 6 (Figure 7a). Although fiddler crabs diversity did
 8 not affect nematodes density, it can be seen in the Figure 7b that two species treatments with
 9 higher levels of density, the densities of nematodes were higher than in one-species treatment.



10 Figure 7 Mean values (\pm SE) of density of nematodes at (a) the densities of 4, 6 and 8 keystone species
 11 and at (b) the densities of 4, 6 and 8 keystone species divided by single and two-species treatments. U:
 12 *Uca uruguayensis*; UL: *Uca uruguayensis* and *Uca leptodactyla*. Different letters mean differences found
 13 on pairwise tests.
 14

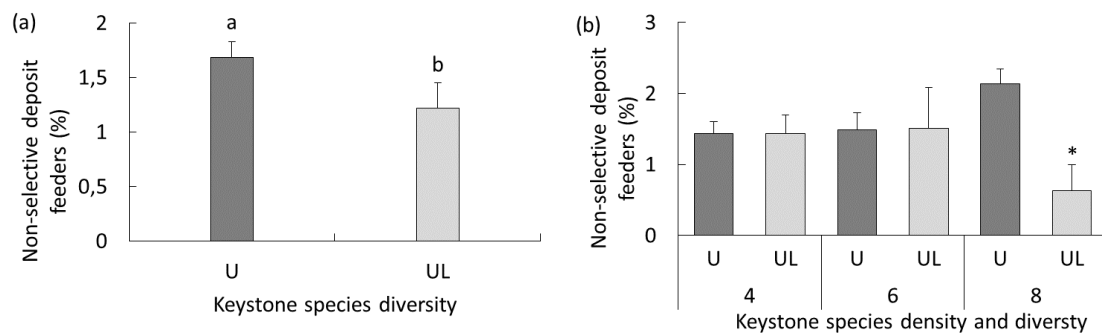
15

16 As total density, densities of *Microlaimus* sp. also decreased with higher density of
 17 fiddler crabs (Fig. 8). *Microlaimus* sp. density varied among the levels 4 and 8 crabs, but not
 18 between 4 and 6, and 6 and 8 (results of pairwise tests are available on Appendix 4). All the other
 19 descriptors of nematodes structure was not affected by the increase of *Uca* density nor richness
 20 ($p(\text{MC})>0,05$; Table2).



1
2 Figure 8 Mean values (\pm SE) of *Microlaimus* sp. density at the densities of 4, 6 and 8 keystone species.
3 Different letters mean differences found on pairwise tests.

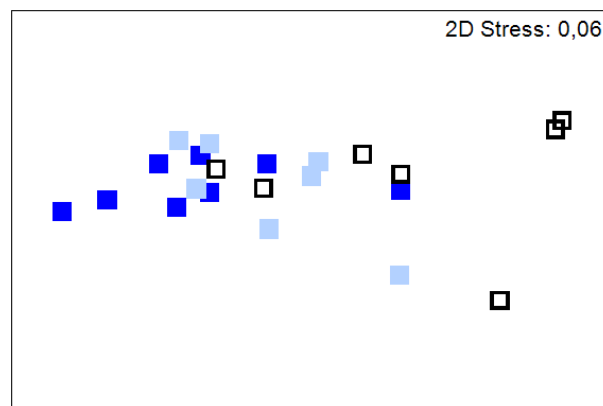
4
5 The relative abundance of non-selective deposit feeders nematodes did not vary
6 significantly between keystone species density ($p(\text{MC})=0,924$), but between keystone species
7 richness it did ($p(\text{MC})=0,031$; Table 2). Non-selective deposit feeders nematodes abundance was
8 low in the presence of the two keystone species (Fig. 9a). This effect was observed at the high
9 level of fiddler crab density (Appendix 4; Fig. 9b).



10
11 Figure 9 Log(x+1) transformed mean values (\pm SE) of non-selective deposit feeders nematodes on (a)
12 single and two-species treatments and on (b) single and two-species treatments at the densities of 4, 6
13 and 8 keystone species. U: *Uca uruguayensis*; UL: *Uca uruguayensis* and *Uca leptodactyla*. Different letters
14 and asterisk (*) mean differences found on pairwise tests.

15
16 The non-metric multidimensional scaling showed a clear separation between the
17 groups. The treatments with 4 crabs are placed mostly on the left of the graph, while treatments
18 with 8 crabs are mostly on the right, 6 crabs treatments are at the center, being intermediate
19 between 4 and 8. Multivariate PERMANOVA showed that fauna varied with keystone species
20 density ($p(\text{MC})=0,008$; Table 2), but with keystone species richness ($p(\text{MC})=0,239$). Pairwise tests

- 1 showed difference between the groups of 4 and 8 crabs, but not between 4 and 6, and 6 and 8
 2 (Appendix 4).



3
 4 Figure 10 MDS ordination for nematodes multivariate structure at densities of 4 (dark blue); 6 (light blue)
 5 and 8 (open squares) crabs.

6

7 Table 2 Results of two-way PERMANOVA evaluating the increase in keystone species diversity and density
 8 on multivariate structure and structural and functional univariate descriptors of nematodes. Data were
 9 log(x+1) transformed when it was necessary. Bold values indicate $p \leq 0.05$. Pairwise tests are available on
 10 Appendix 4. df: degrees of freedom; SS: sum of squares; MS: mean squares; p(MC): p-value obtained with
 11 Monte Carlo permutation test.

Source of variation	df	SS	MS	F	p(MC)
Multivariate structure					
<i>Uca</i> density	2	1940,2	970,1	4,0555	0,008
<i>Uca</i> richness (<i>Uca</i> density)	3	990,52	330,17	1,3803	0,239
Residual	16	3827,3	239,2		
Nematodes structure					
Density					
<i>Uca</i> density	2	1025,7	512,83	6,4585	0,017
<i>Uca</i> richness (<i>Uca</i> density)	3	300,02	100,01	1,2595	0,331
Residual	16	1270,5	79,403		
Number of genera					
<i>Uca</i> density	2	13,452	6,7259	1,9245	0,194
<i>Uca</i> richness (<i>Uca</i> density)	3	2,1548	0,7182	0,2055	0,907
Residual	16	55,917	3,4948		
Diversity (H')					
<i>Uca</i> density	2	0,0292	0,0146	0,5067	0,628
<i>Uca</i> richness (<i>Uca</i> density)	3	0,0271	0,0090	0,3137	0,824
Residual	16	0,4610	0,0288		

1 Table 2 Results of two-way PERMANOVA evaluating the increase in keystone species diversity and density
 2 on multivariate structure and structural and functional univariate descriptors of nematodes. Data were
 3 log(x+1) transformed when it was necessary. Bold values indicate $p \leq 0.05$. Pairwise tests are available on
 4 Appendix 4. df: degrees of freedom; SS: sum of squares; MS: mean squares; p(MC): p-value obtained with
 5 Monte Carlo permutation test.

Source of variation	df	SS	MS	F	p(MC)
<i>Microlaimus</i> density					
<i>Uca</i> density	2	750,82	375,41	5,8627	0,017
<i>Uca</i> richness (<i>Uca</i> density)	3	236,74	78,914	1,2324	0,333
Residual	16	1024,5	64,033		
Nematodes functioning					
Selective deposit feeders					
<i>Uca</i> density	2	60,901	30,451	1,0745	0,358
<i>Uca</i> richness (<i>Uca</i> density)	3	5,402	1,8007	0,0635	0,978
Residual	16	453,41	28,338		
Non-selective deposit feeders					
<i>Uca</i> density	2	0,0474	0,0237	0,0739	0,924
<i>Uca</i> richness (<i>Uca</i> density)	3	3,8954	1,2985	4,0454	0,031
Residual	16	5,1356	0,3209		
Epistratum feeders					
<i>Uca</i> density	2	89,319	44,659	1,327	0,291
<i>Uca</i> richness (<i>Uca</i> density)	3	47,47	15,823	0,4701	0,709
Residual	16	538,47	33,654		
Predators/omnivores					
<i>Uca</i> density	2	0,9453	0,4726	2,1736	0,146
<i>Uca</i> richness (<i>Uca</i> density)	3	1,4923	0,4974	2,2875	0,109
Residual	16	3,4794	0,2174		
Index of Trophic diversity					
<i>Uca</i> density	2	0,0158	0,0079	1,2154	0,313
<i>Uca</i> richness (<i>Uca</i> density)	3	0,0118	0,0039	0,6036	0,63
Residual	16	0,1044	0,0065		
Maturity index					
<i>Uca</i> density	2	0,0017	0,0008	0,1780	0,856
<i>Uca</i> richness (<i>Uca</i> density)	3	0,0130	0,0043	0,8694	0,476
Residual	16	0,0799	0,0049		

6

7 DISCUSSION

8 As we hypothesized, the poleward displacement of keystone species as a response
 9 of climate change may affect the structure and functioning of the meiofauna communities of

1 intertidal flats. In our experiments, changes in fiddler crabs diversity affected nematodes
2 functioning, by altering the relative abundance of predators/omnivores nematodes and non-
3 selective deposit feeders. Moreover, the increase in density of fiddler crabs affected negatively
4 the nematodes assemblage decreasing their densities and the density of the most abundant
5 genera. The entrance of a new invertebrate species in marine environments may affect the
6 native community by altering food web dynamics and modifying physical environment (Crooks,
7 1998; Mistri, 2003; Greene *et al.*, 2011; Chan & Bendell, 2013).

8 In our replacement series experiment, high fiddler crab diversity did not lead to
9 changes in structure of meiofauna. This result may be due to *U. uruguayensis* and *U. leptodactyla*
10 are similar (Crane, 1975) and may explore similar niches by reworking similar volumes of
11 sediments at almost identical depths (Machado *et al.*, 2013). O'Connor & Bruno (2009) also did
12 not found changes in the structure of diverse prey communities such as our study with
13 meiofauna, with shifting predator richness. According to the authors, studies reporting
14 significant effects on prey composition with the increase of predator richness, (e.g. Siddon &
15 Witman, 2004; Finke & Denno, 2005; Byrnes *et al.*, 2006; Douglass *et al.*, 2008; Naddafi &
16 Rudstam, 2013), used a small number of prey species (one to five species). The effects of
17 predator richness therefore, seem to be more important under conditions of low prey diversity.
18 If this is true, future experiments should be done including different nematodes communities
19 with high and low levels of nematodes diversity.

20 While the increase of keystone species diversity did not affect the structure
21 nematode assemblages, it did affect their trophic functioning. The relative abundance of
22 predators/omnivores nematodes declined when more *U. leptodactyla* were present. The exact
23 nature of this change in the trophic structure of the nematode assemblages is not easily to
24 explain. The most abundant predators/omnivores nematodes found in this work belong to the
25 genus *Oncholaimus* (Oncholaimidae), nematodes that usually have large body and occupy the
26 surface sediment layers (Moens *et al.*, 2013). Considering *Oncholaimus* size, or the superficial

1 layer of sediment that these nematodes occupy, the genera could be more easily ingested by
2 crabs or disturbed in their presence. However, although fiddler crabs may feed actively on
3 meiofauna, they probably do not select species as they feed by flotation feeding (Quinn, 1980;
4 Dittmann, 1993), so the size of the organism may be important during this process.
5 Unfortunately, we did not measure the nematodes biomass to evaluate if there is a relationship
6 between the size of nematodes and *Uca* species. Future studies should include analysis of
7 nematodes biomass to explore this question.

8 When we evaluated the effects of keystone species diversity together with density
9 in the partial additive design experiment, no effects of *Uca* diversity on nematodes assemblage
10 structure were found, confirming our results from the replacement series. Increasing fiddler
11 crabs density, however, affected significantly the structure of nematodes. The total density of
12 nematodes and the density of the most abundant genus (*Microlaimus*) declined with increasing
13 fiddler crabs density independently of the species used. Similarly, in a field enclosure experiment
14 Guidetti (2007) observed a decrease in the density of sea urchin, caused by an enhancement of
15 predation, when he increased predator density. Our results confirm the important top-down
16 regulation of the meiofauna by fiddler crabs by means of sediment disturbance and predation
17 observed by Ólafsson (2003) and Citadin *et al.* (2016 *in press*). The increase of keystone species
18 therefore, may reduce nematodes density directly via predation, or indirectly via modification
19 of the sediment environment.

20 Although *Uca* diversity caused no significant effects in nematodes structure, the
21 mean density of nematodes was higher in the two-species treatments at high and mid levels of
22 keystone species density. These results may suggest a reduction of the top-down control when
23 both species of crabs are together in high densities. The reduction of predation is not unusual
24 when two species forage together, Siddon & Witman (2004), for instance, reported a significant
25 risk reduction for urchins when two predators were combined.

1 The abundances of non-selective deposit feeders nematodes decreased in the
2 higher levels of keystone species diversity and density. This low abundance in this functional
3 feeding group of nematodes may also be due the activities of bioturbation exerted by fiddler
4 crabs. During burrowing activities, fiddler crabs manipulate large quantities of sediment
5 increasing habitat heterogeneity both horizontally and vertically (Hoffman *et al.*, 1984; Botto *et*
6 *al.*, 2008; Sayão-Aguiar *et al.*, 2012). The maximum depth burrowed by *U. leptodactyla* is
7 generally higher than *U. uruguayensis* (Machado *et al.*, 2013), with high abundance of *U.*
8 *leptodactyla* in mesocosms it is possible that more non-selective deposit feeders nematodes are
9 being affected by *Uca*'s disturbance.

10 Our predictions about the effects on nematodes assemblages due climate changes
11 refer exclusively to those expected as a response to the increase of keystone species. Besides
12 changes in patters of keystone species distributions, other points may be considered to evaluate
13 the effects of climate variability on nematodes assemblage. For instance, rising temperatures
14 may affect keystone-species behavior by increasing foraging activities, as observed by Sanford
15 (1999) affecting even further the meiofauna. Increasing temperature may also affect
16 nematodes structure, as seen by Gingold *et al.* (2013), resulting in loss of density, richness,
17 diversity, abundance and community biomass.

18 The increase of keystone species diversity could cause fewer effects in lower
19 latitudes where there is a bigger number of species (Fischer, 1961; Hillebrand, 2004) than in high
20 latitudes. The high biodiversity found in low latitudes increases the possibility of several species
21 perform the same function resulting in ecological redundancy, which is not observed at high
22 latitudes. However, these patterns of species distributions are been modified by climate
23 changes. Tropicalization of temperate zones due to species poleward displacements could lead
24 to increase species richness on temperate zones and poles (Molinos *et al.*, 2015).

25 In summary, our results suggest that global climate changes may indirectly affect
26 meiofauna community via displacement of keystone species. Global warming and climate

1 changes will continue unless anthropogenic emissions of greenhouse gases stop. More studies
 2 evaluating the direct and indirect impacts of climate changes on ecosystems should be done to
 3 predict future changes and explore ways to mitigate them.

4

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- 1 **Appendix 1.** Total abundance of nematodes genera in replacement series experiment on
 2 treatments: 8U; 6U+2L; 4U+4L and 2U+6L. U: *Uca uruguayensis* and L: *Uca leptodactyla*.

Genera	8U	6U+2L	4U+4L	2U+6L
<i>Anoplostoma</i>	104	147	171	140
<i>Chromadorella</i>	4	13	12	15
Chromadoridae indet	0	0	1	1
<i>Dichromadora</i>	60	83	102	90
<i>Hypodontolaimus</i>	0	1	0	0
<i>Neochromadora</i>	3	0	0	1
<i>Ptychollaimelus</i>	0	0	0	1
<i>Paracanthonchus</i>	7	9	14	11
<i>Paracyatholaimus</i>	0	0	1	7
<i>Pomponema</i>	1	0	0	0
Dorylaiminae	2	5	7	5
<i>Ethmolaimus</i>	1	3	4	5
<i>Haliplectus</i>	2	5	2	1
<i>Leptolaimus</i>	3	2	2	0
Linhomoeidae	0	3	2	1
<i>Paralinhomoeus</i>	0	0	2	0
<i>Terschellingia</i>	73	65	146	120
<i>Microlaimus</i>	57	120	131	138
<i>Diplolaimella</i>	0	1	0	0
<i>Diplolaimelloides</i>	2	2	0	1
<i>Thalassomonhystera</i>	5	7	1	5
Monhysteridae	0	1	0	1
<i>Oncholaimus</i>	39	59	40	49
<i>Viscosia</i>	1	1	2	1
Oncholaimidae	4	46	4	1
<i>Halalaimus</i>	4	8	10	9
<i>Oxystomina</i>	2	2	3	1
<i>Subsphaerolaimus</i>	1	1	1	1
<i>Rhabdocoma</i>	5	1	8	0
<i>Trefusia</i>	1	1	1	0

- 1 **Appendix 2.** Pairwise PERMANOVA tests from replacement series experiment for effects of
 2 keystone species treatments (8U; 6U+2L; 4U+4L and 2U+6L, U: *Uca uruguayensis* and L: *Uca*
 3 *leptodactyla*.) on predators/omnivores nematodes. t: pseudo-t; p(MC): p-value obtained with
 4 Monte Carlo permutation test.

Groups	t	p(MC)
8U, 6U+2L	0,083303	0,933
8U, 4U+4L	3,4642	0,012
8U, 2U+6L	1,3783	0,205
6U+2L, 4U+4L	3,5994	0,011
6U+2L, 2U+6L	1,3616	0,236
4U+4L, 2U+6L	1,2574	0,246

- 1 **Appendix 3.** Total abundance of nematodes genera in partial additive design experiment on two
 2 and single keystone species treatments and among levels of density.

Genera	<i>U. leptodactyla</i> + <i>U. uruguayensis</i>			<i>U. uruguayensis</i>		
	4	6	8	4	6	8
<i>Anoplostoma</i>	0	0	1	2	1	2
<i>Axonolaimus</i>	0	0	0	0	1	2
<i>Chromadorella</i>	0	1	0	0	1	1
<i>Desmodora</i>	1	0	1	0	1	0
<i>Dichromadora</i>	0	1	0	0	0	0
<i>Diplolaimelloides</i>	17	19	2	18	12	19
Dorylaiminae	0	1	0	0	0	0
<i>Ethmolaimus</i>	3	3	3	8	1	3
<i>Halalaimus</i>	0	0	1	1	0	1
<i>Haliplectus</i>	12	13	8	15	15	3
<i>Linhomoeus</i>	0	1	0	0	0	0
<i>Microlaimus</i>	410	303	243	477	333	234
Oncholaimidae	0	0	0	1	0	0
<i>Oncholaimus</i>	1	0	0	3	1	0
<i>Paracanthochus</i>	1	2	0	0	0	1
<i>Paracyatholaimus</i>	1	0	0	0	0	0
<i>Siphonolaimus</i>	0	0	0	1	1	0
<i>Terschellingia</i>	31	34	26	32	35	27
<i>Thalassomonhystera</i>	1	0	0	0	1	0
<i>Viscosia</i>	0	0	0	0	1	0

- 1 **Appendix 4.** Pairwise PERMANOVA tests from partial additive design experiment for effects of
 2 keystone species density (4: four crabs, 6: six crabs and 8: eight crabs) and keystone species
 3 richness (U: *U. uruguayensis* and UL: *U. uruguayensis* + *U. leptodactyla*; nested in keystone
 4 species density) on nematodes multivariate structure and univariate descriptors. t: pseudo-t;
 5 p(MC): p-value obtained with Monte Carlo permutation test.

Groups (keystone species density)	Multivariate		Total Density		<i>Microlaimus</i> density	
	t	p(MC)	t	p(MC)	t	p(MC)
4,6	1,1293	0,279	1,2827	0,234	1,5337	0,163
4,8	2,6127	0,013	3,0847	0,01	3,109	0,011
6,8	1,8347	0,056	2,8677	0,02	2,1072	0,056
Non-selective deposit feeders						
Groups (Keystone species richness nested in keystone species density)			t	p(MC)		
U,UL (4)			0,013826	0,994		
U,UL (6)			0,043613	0,963		
U,UL (8)			3,8136	0,011		