

Can environmental pollution by metals change genetic diversity? *Ucides cordatus* (Linnaeus, 1763) as a study case in Southeastern Brazilian mangroves



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ABSTRACT

Industrial areas on estuarine systems are commonly affected by heavy metals, affecting all local biota. Random Amplified Polymorphic DNA (RAPD) was used to evaluate genetic diversity of *Ucides cordatus* at mangroves in southeastern Brazil (Juréia, J; São Vicente, SV; and Cubatão, C), with distinct pollution levels by metals. The genetic diversity of this species was compared with concentrations of metals (Cd, Pb, Cu, Cr and Hg) in the environment. A pollution gradient was confirmed (SV > C > J), with low levels detected in water, except for mercury in SV. All metals in the sediment samples were below Threshold Effect Level (TEL), without an apparent biological risk to the biota. Genetic distance was very similar between J and C, with SV occurring as an out-group. RAPD was a powerful tool to investigate the effect of metal pollution on genetic diversity of this mangrove crab, and to evaluate the conservation status of the mangrove ecosystem.

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1. Introduction

Mangroves are an ecosystem with intense recycling of nutrients, providing natural nursery grounds for many animal species (e.g. fish, crustaceans and mollusks) (Schaeffer-Novelli, 1995; Spalding, 2012; Lee et al., 2014), and an ideal site for feeding and nesting of birds (Vannucci, 2001; Luther and Greenberg, 2009; Huertas et al., 2016). Biodiversity is relatively low in mangroves, due to physiological adaptations required to support the wide variation of salinity and due to its anoxic/unstable/contaminated sediment (Pinheiro et al., 2008a; Thilagavathi et al., 2013; Remaili et al., 2016).

Recently, mangroves has been purposed to monitor climatic changes (Schaeffer-Novelli et al., 2016), mainly using some species of decapod crustacean that inhabitant this environment (Gilman et al., 2008; Pinheiro & Almeida, 2015; Siddig et al., 2016). Among these crustaceans *Ucides cordatus* (Linnaeus, 1763) is an endemic brachyuran that is widely distributed in tropical mangroves of the western Atlantic (Melo, 1996). This species builds its galleries in the sediment and feeds on senescent

leaves and propagules present on the substrate (Koch and Nordhaus, 2010; Christofolletti et al., 2013), processing 84% of the mangrove litter (Koch & Wolff, 2002). Besides playing a relevant role in carbon cycle by litter processing (Begon et al., 1996; Hogarth, 1999), this crab species is also an important food and subsistence source to coastal communities (Alves & Nishida, 2003), and has been recently used as bioindicator species for evaluating the conservation status of mangrove areas (Pinheiro et al., 2013). *U. cordatus* has also been successfully used as a tool to diagnose and classify the human impacts on environmental quality of mangrove ecosystem based on a multi-level analysis (Duarte et al., 2016).

In Brazil, the southeastern coast is the most disturbed, mainly the central coast of the São Paulo State. From this, the Metropolitan Region of 'Baixada Santista' (MRBS) including nine municipalities with around 1.7 million inhabitants (Pinheiro et al., 2008b) stands out. The ecosystems in this coastal region are under significant pressure due to industrial and port activities, with a historic use that dates back >500 years (Oliveira et al., 2008). Some estuarine areas (e.g. São Vicente municipality) are characterized by the presence of many still houses on the river banks and estuaries (Azevedo et al., 2012), without sanitary condition and correct destination of solid wastes (Cordeiro & Costa, 2010). While in Brazilian mangroves the main threats are the harvesting of mangrove wood, deforestation for aquaculture ponds, and intense

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property speculation (Macintosh & Ashton, 2002; Ellison, 2008), at RMBS industrial activities, such as a port complex, prevails, posing an imminent risk, with toxic waste, which may affect the biota (De Wolf et al., 2004). Duarte et al. (2016) quantified and qualified the main pollutants in São Paulo coast areas, with a high concentration of metals and organic contaminants in Central area (where São Vicente and Cubatão are placed) when compared to those registered in South area (where there are conservation units as Juréia).

Several studies have described the effects of pollution on mangrove vegetation (Silva et al., 1990; Arrivabene et al., 2015; Souza et al., 2015) and crustaceans (Harris & Santos, 2000; Valdovinos & Zúñiga, 2002; Álvaro et al., 2015), but the use of molecular tools for detecting pollution impacts on the genetic level, particularly in estuarine ecosystem, are scarce. The Random Amplified Polymorphic DNA (RAPD), i.e., is one technique applied in ecotoxicology studies (De Wolf et al., 2004; Giantsis et al., 2012; Liu et al., 2012; Salem et al., 2014; Zhang et al., 2016) for quantification of alleles (bands) and analyses of their addition/loss due to mutation, inversion, deletion or chromosomal rearrangement (De Wolf et al., 2004). This genetic fingerprinting tool is a relatively inexpensive and fast method for evaluating pollutants effects on a broad range of DNA damages, thus, improving environmental risk assessment, mainly in developing countries as Brazil, where this study was carried out.

Our main objective was to evaluate if the presence of heavy metals in environmental matrices (water and sediment) could affect the genetic diversity and structure of the mangrove crab *U. cordatus*. To this aim, we used Random Amplified Polymorphic DNA (RAPD) technique to

assess the genetic variation of *U. cordatus* from three mangrove areas from Southeastern Brazil (Juréia, São Vicente and Cubatão), characterized by distinct pollution levels by metals (Cd, Pb, Cu, Cr and Hg). A positive association between genetic parameters and concentration of pollutants will allow the use of this tool in studies about environmental conservation, helping government management plans related to the mangrove ecosystem and the target species.

2. Materials and methods

2.1. Study area

Three mangrove areas, which presumably differ in their degree of environmental impacts, were investigated (Fig. 1) from January to March 2009. Juréia is a legally protected area that composes a great 'Mosaic of Conservation Units' near the mouth of the Una River ($24^{\circ}26'0''$ S– $47^{\circ}04'5''$ W), with presence of a traditional community comprising a few people (São Paulo, 2006). São Vicente was represented by a mangrove area near the Branco River ($23^{\circ}56'2''$ S– $46^{\circ}28'1''$ W) in São Vicente Municipality, with an impact of 316,324 inhabitants (IBGE, 2010), 11 sources of industrial pollution (Pinheiro et al., 2013), and also impacted. In turn, Cubatão was comprised by a mangrove close to USIMINAS (an important Brazilian steel Company), and adjacent to the Morrões River ($23^{\circ}52'5''$ S– $46^{\circ}22'2''$ W), in Cubatão Municipality, with 116,010 inhabitants (IBGE, 2010). This is one of the largest industrial centers in Brazil (23 industrial complexes, 111 factories, and >300

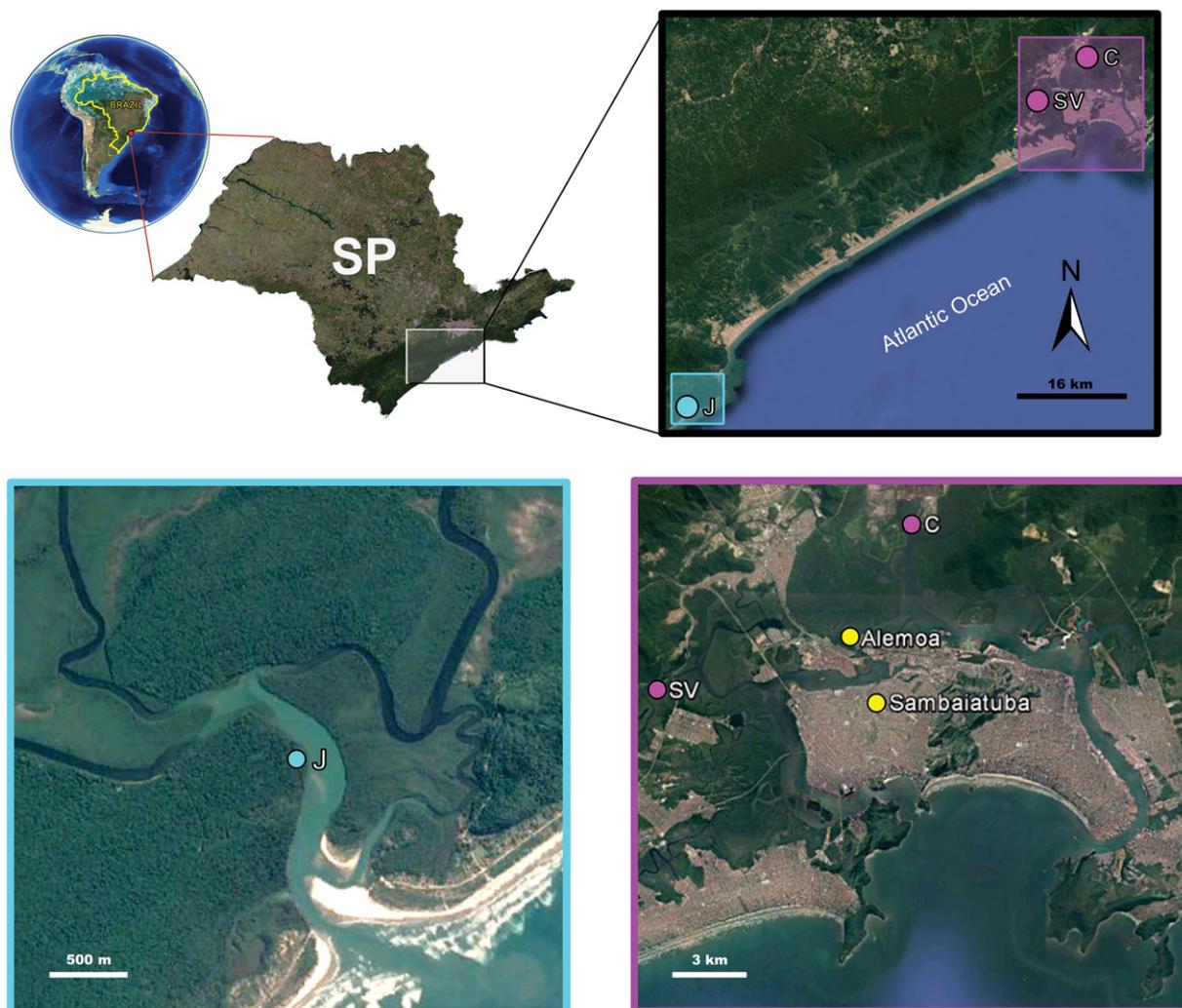


Fig. 1. Map of São Paulo state coast (Brazil), showing the three mangrove areas studied (C, Cubatão; J, Juréia; and SV, São Vicente), and the locations of Sambaibatuba and Alemoa dumps.

pollution sources, according Pinheiro et al., 2012), that produce fertilizers, steel and petrochemical compounds. In the 1980s, the high pollution at Cubatão was the subject of major global environmental discussions because the incidence of several diseases, most of them linked to genetic mutations (e.g. anencephaly in newborns) (Monteleone-Neto & Castilla, 1994). Other diffuses sources of pollutants that act in estuarine areas of Central coast of São Paulo state are Sambaiatuba dump (23°55'18"S–46°23'50"W) and Alemoa dump (23°56'40"S–46°23'11"W).

2.2. Metals

In each mangrove area, we collected three non-adjacent sediment samples at different depths (surface, 15, 30 and 45 cm), and three non-contiguous water samples along of a perpendicular transect from the riverbank toward the upper salt flat ("apicum") of the mangroves. The concentrations of five metals (Cd, Pb, Cu, Cr and Hg) from these environmental samples were measured at CEATOX (IB/UNESP Botucatu) by wet mineralization with 65% HNO₃ (Basset et al., 1981) and read by an atomic absorption spectrophotometer (GBC - 932 AA) (Athanasopoulos, 1993).

The concentrations of metals in water were compared to reference values for brackish water 'class 1' (minimum human impact), according to National Council of the Environment (CONAMA n° 357/2005, see Table 1). This Brazilian law, however, does not provide the reference values for estuarine sediments (Oliveira et al., 2007). Therefore, these results were compared to guidelines provided by Environment Canada (1999), comprising two categories: 1) Threshold Effect Level (TEL), the concentration below which adverse biological effects are rarely observed (<10%); and 2) Probable Effect Level (PEL), the concentration above which adverse biological effects are frequently observed (Table 1). We assume that Cd, Cr, Cu, Hg and Pb are transferred from the environment (water and sediment) to *U. cordatus* (tissues) because there is evidence that the mangrove crab and its main food source may accumulate these heavy metals (Pinheiro et al., 2012). Additionally, *U. cordatus* individuals from Juréia presented lower frequency of micronucleated cells (Pinheiro et al., 2013; Duarte et al., 2016) and higher neutral red retention time (Duarte et al., 2016) when compared to those sampled in Cubatão and São Vicente.

Data were entered in spreadsheets, and the statistical analyses were conducted using 'R' Version 3.2.3 (R Core Team, 2016). The metal concentrations were first tested for normality (Shapiro-Wilk) and homoscedasticity of the variances (Bartlett's test), which are prerequisites for the application of one- or two-way ANOVA or Student's *t*-test. When the data did not satisfy these prerequisites, the nonparametric Kruskal-Wallis test was applied (Zar, 1999). A statistical level of 5% was used to evaluate all analyses.

2.3. Capture of crabs and acquisition of genetic material

In each mangrove area, 15 specimens of *U. cordatus* were manually collected. All crabs had a cephalothoracic width (CW) > 60 mm, in

Table 1

Reference values of threshold effect level (TEL) and probable effect level (PEL) for concentrations of metals in sediment and brackish-water 'class 1' (with minimum human impact).

Metals	Sediment ^a (µg/g = mg/kg)		Brackish water ^b (µg/mL = mg/L)
	TEL	PEL	
Cu	19.0	110.0	0.005
Cd	0.67	4.20	0.005
Cr	52.0	160.0	0.050
Pb	30.0	110.0	0.010
Hg	0.13	0.70	0.0002

^a Environment Canada (1999).

^b National Council of the Environment (CONAMA law n° 357/2005).

accordance with IBAMA n° 52/2003 (Ibama, 2003). In the laboratory, about 1 cm³ of muscle tissue from the cephalothorax was removed from each animal and placed in Eppendorf tubes (2 mL) containing absolute ethanol. The tissues were then frozen at –20 °C for later molecular analyses.

2.4. DNA extraction and RAPD technique

For the extraction of genomic DNA, we followed the protocol described by Sambrook et al. (1989), with adaptations. Muscle tissue (20 mg) from each animal was macerated in liquid nitrogen, under aseptic conditions, and transferred to a 15 mL tube. A lysis solution was added to each tube, containing 80 µL of Proteinase K buffer, 40 µL of Proteinase K, 20 µL of 20% SDS and 240 µL of ultrapure water. Each tube was then inverted ten times to homogenize the solution with the macerated tissue. The tubes were left in a 60 °C bath overnight. Next day, the tubes were removed from the bath, and once they reached room temperature, 100 µL of 6 M NaCl was added while the tube was gently shaken. The tubes were then centrifuged at 13,000 rpm at room temperature, for 10 min. The supernatant was recovered and centrifuged once more. The supernatant was recovered again and 1 V of ice-cold isopropanol was added. Each tube was inverted for 2 min, and left for 8 min at room temperature, then centrifuged again at 13,000 rpm at 4 °C for 15 min. The supernatant was discarded and 1 mL of ice-cold 70% ethanol was added. The tubes were centrifuged again at 13,000 rpm, 4 °C for 5 min. The supernatant was discarded and the pellet was dried at room temperature. The pellet was then resuspended in 100 µL of sterilized ultrapure water, and refrigerated (4 °C) for at least 5 h, before being used. The quality of the extracted DNA was evaluated in agarose gel (0.8%) stained with ethidium bromide (0.1 10⁻³ V).

For the RAPD technique, 21 decamer primers of the Eurofins MWG Operon RAPD 10mer Kits were tested for amplification quality, seven of which (OPE12, OPE14, OPE15, OPE18, OPG5, OPI7 and OPJ9; see Table 2) resulted in reliable amplification.

The 20 µL polymerase chain reactions (PCR) had the final concentrations: 0.25 mM of dNTP, 1 × reaction buffer, 4 mM of MgCl₂, 0.1 U/µL of Taq DNA polymerase, 0.4 µM of primer, and 0.4 ng/µL of template DNA (Oliveira-Neto et al., 2007a, b). The thermocycler (MJ-Research, ICN-PTC-100-Programmable Thermal Controller) conditions involved initial denaturation for 3 min at 95 °C, followed by 35 cycles of 15 s. denaturation at 94 °C, 30 s. annealing at 35 °C, 1 s. extension at 72 °C, and one 4-min cycle at 72 °C for final extension. We used negative and positive controls in all reactions and each reaction was carried out in duplicate to minimize non-reproducibility issues.

The amplification products were observed in a Gel Doc 1000 gel documentation system (Bio-Rad, Hercules, California, USA), with the software Quantity OneR (Bio-Rad). Based on this gel analysis, binary numerical matrices were constructed (0, absence; 1, presence) for comparing the bands amplified with each primer. The distance migrated by the fragments (mm) was measured and converted to size in base pairs utilizing the *Fragment Size Calculator* (available at <http://www.basic.northwestern.edu/biotools/SizeCalc.html>).

We quantified the genetic diversity using *GenAlEx 6.5* (Peakall & Smouse, 2012) to calculate the following descriptive statistics: number

Table 2

Oligonucleotides used, and their sequences.

Oligonucleotide	Sequence
OPE12	5'-TTATCGCCCC-3'
OPE14	5'-TGCGGCTGAG-3'
OPE15	5'-ACGCACAACC-3'
OPE18	5'-GGACTGCAGA-3'
OPG5	5'-CTGAGACGGA-3'
OPI7	5'-CAGCGACAAG-3'
OPJ9	5'-TGAGCCTCAC-3'

of bands, number of private bands, percentage of polymorphic loci and unbiased expected heterozygosity (H_E) assuming Hardy-Weinberg model for each sampling locality. To describe how the genetic variation is structured, we carried out the Analysis of Molecular Variance (AMOVA; Excoffier et al. 1992) and estimated F_{ST} analogue (Φ_{ST}) also using *GenAlEx 6.5* (Peakall & Smouse, 2012) with 9999 permutations for testing the hypothesis that there is no difference among groups. Furthermore, using *poppr 2.2.1* (Kamvar et al., 2014), we calculated Nei's genetic distance (1978) among individuals and used Neighbor-Joining (NJ) clustering method (Saitou & Nei, 1987) with 10,000 bootstrap pseudo-replication to measure branching supports. Finally, we performed a Discriminant Analysis of Principal Components (DAPC; Jombart et al., 2010) to find the number of genetically related individuals that best explain our data and describe the clusters with Bayesian Information Criterion. This method is implemented with *adegenet 2.0.1 R* package (Jombart, 2008), and it relies on data transformation using Principal Component Analysis (PCA) prior to a Discriminant Analysis (DA) that partitions the genetic variation into between group and within group, maximizing the former (Jombart et al., 2010). To avoid overfitting, we used the function *optim.a.score*.

3. Results

3.1. Analysis of metals

Detectable Pb concentrations in water samples were registered in all mangrove areas with concentrations following a normal tendency (Shapiro-Wilk, $P > 0.05$), but without difference among mean concentrations when these areas were compared ($t = 0.73$; $P = 0.51$) (Table 3). At Juréia, the Pb concentration in water was $<0.05 \mu\text{g/mL}$ (estuary and 10 m from the riverbank), but higher in water samples obtained more internally in mangrove forest (0.12 and $0.28 \mu\text{g/mL}$, with 20 and 30 m from the riverbank, respectively); in Cubatão Pb concentrations in water were more homogeneous ($0.12\text{--}0.19 \mu\text{g/mL}$). All the other metals had concentrations below detection limits, both in Juréia and Cubatão (Cu and Cr: <0.05 ; Cd: <0.01 ; and Hg: $<0.0001 \mu\text{g/mL}$), except for Hg in São Vicente ($0.44 \mu\text{g/mL}$).

Concentration of Hg in mangrove sediment may be considered absent in all mangrove areas and its strata ($<0.10 \text{ng/g}$). The other registered metals did not have a normal distribution (Shapiro-Wilk, $P < 0.05$), without a significant difference among sediment strata ($0.40 < KW < 1.64$; $P > 0.41$) in all mangrove areas (Table 4). Concentrations of Cu, Cd, Cr and Pb were higher at São Vicente and Cubatão but were similar between them, both contrasting from Juréia (Kruskal-Wallis, $5.81 < KW < 16.34$; $P < 0.015$).

Table 3

Concentrations of metals in sediment (mean \pm standard error) at four depths in mangroves from Juréia, Cubatão and São Vicente, São Paulo state (Brazil). Means of concentration in a same metal followed by an equal letter did not differ ($P > 0.05$).

Mangrove	Depth	Metal concentrations ($\mu\text{g}\cdot\text{g}^{-1}$ * and $\text{ng}\cdot\text{g}^{-1}$ **)				
Areas	(cm)	Cu*	Cd*	Cr*	Pb*	Hg**
Juréia	Surface	1.92 \pm 0.12	0.053 \pm 0.003	3.17 \pm 0.17	4.56 \pm 0.14	<0.10
	15	1.30 \pm 0.17	0.046 \pm 0.003	3.07 \pm 0.61	3.18 \pm 0.25	<0.10
	30	1.06 \pm 0.12	0.043 \pm 0.003	2.14 \pm 0.17	2.78 \pm 0.53	<0.10
	45	1.72 \pm 0.45	0.056 \pm 0.007	4.37 \pm 1.33	3.97 \pm 0.85	<0.10
	Total	1.39 \pm 0.30a	0.046 \pm 0.005a	3.01 \pm 0.79a	3.30 \pm 0.61a	<0.10
Cubatão	Surface	3.82 \pm 0.37	0.064 \pm 0.009	6.39 \pm 0.74	5.33 \pm 0.53	<0.10
	15	4.05 \pm 0.22	0.077 \pm 0.013	6.91 \pm 0.40	6.18 \pm 0.53	<0.10
	30	3.78 \pm 0.21	0.063 \pm 0.009	7.47 \pm 1.25	5.41 \pm 0.21	<0.10
	45	2.81 \pm 0.49	0.057 \pm 0.009	5.24 \pm 0.97	3.86 \pm 0.95	<0.10
	Total	3.30 \pm 0.40b	0.060 \pm 0.010ab	6.05 \pm 0.91b	4.75 \pm 0.73ab	<0.10
São Vicente	Surface	5.81 \pm 1.28	0.100 \pm 0.023	7.07 \pm 0.45	10.33 \pm 0.56	<0.10
	15	5.35 \pm 0.34	0.063 \pm 0.003	6.00 \pm 0.80	5.93 \pm 2.98	<0.10
	30	4.32 \pm 0.18	0.087 \pm 0.007	5.26 \pm 0.69	10.04 \pm 2.11	<0.10
	45	5.29 \pm 0.69	0.080 \pm 0.010	7.72 \pm 1.73	10.61 \pm 0.30	<0.10
	Total	4.81 \pm 0.72b	0.077 \pm 0.014b	6.07 \pm 1.05b	8.31 \pm 1.96b	<0.10

Table 4

Concentration of metals in water (mean \pm standard error; or <'value', minimum concentration detected), in mangroves from Juréia, Cubatão and São Vicente, state of São Paulo, Brazil. Means of concentration of Pb followed by an equal letter did not differ ($P > 0.05$).

Mangrove area	Metal concentrations ($\text{mg}\cdot\text{L}^{-1}$)				
	Cu	Cd	Cr	Pb	Hg
Juréia	<0.05	<0.01	<0.05	0.10 \pm 0.07a	<0.10
Cubatão	<0.05	<0.01	<0.05	0.15 \pm 0.02a	<0.10
São Vicente	<0.05	<0.01	<0.05	0.18 \pm 0.07a	0.44 \pm 0.00

3.2. Analysis of molecular genetic markers

All seven primers we used reliably amplified a range of 3–8 loci/individual, resulting in an average of seven loci/individual. Overall, we observed 83.78% of polymorphic bands (considering the entire sampling and all seven primers: SE = 0.0413) and a low expected heterozygosity ($H_E = 0.318$, SE = 0.018). Regarding each population, although we observed that there was no substantial difference on the levels of genetic diversity between the three locations, unexpectedly, samples from Juréia, our reference environment, presented the lowest values for every descriptive statistic we used (Table 5).

We observed the largest amount of the genetic variation within each population (88%). Despite the low genetic variation between the evaluated populations (12%), the genetic differentiation between them was significantly higher than expected by chance ($\Phi_{ST} = 0.117$, $P < 0.001$). The genetic relation between individuals based on Nei's genetic distance (Nei, 1978) and NJ clustering revealed a complex scenario where individuals from the three sampling locations do not compose well defined (nor statistically supported by the bootstrapping procedure) clades (Fig. 2A). The genetic structure was better described by using DAPC (Jombart et al., 2010). The number of groups (k) that best explained our data was three. The pattern of genetic structure we observed based on two Principal Components that we retained after performing the *optim.a.score* function suggested that each sampling location is composed of a mixture of genetically distinct individuals (Fig. 2B). Interestingly, most individuals from São Vicente composed one inferred group, and only a few individuals from this location were assigned to the other genetic clusters. These clusters were majoritarily constituted by a mixture of individuals from both Cubatão and Juréia (Fig. 2B).

4. Discussion

Mangrove sediments are considered an important geochemical barrier to immobilize pollutants as metals (Neinavaz et al., 2012) due

Table 5

Genetic diversity of *Ucides cordatus* measured by DNA fingerprinting based on RAPD, with samples from 15 individuals. Where: H_E, heterozygosity; SE, standard error.

Population	Juréia	Cubatão	São Vicente
Number of Bands	32	36	34
Number of Private Bands	0	0	1
H _E (±SE)	0.278 ± 0.033	0.314 ± 0.030	0.363 ± 0.029
Polymorphic Loci (%)	75.7	89.2	86.5

to its high organic matter, cation exchange capacity (CEC), and finer fractions (Faridah-Hanum et al., 2014). The mobility and bioavailability of metals in the environment can be modulated by sorption/desorption and precipitation/dissolution (Skrbic and Djurisc-Mladenovic, 2010). Therefore, in acidic environments occur the weakening of these chemical bonds, a reduction of CEC and release of metals to ground water (Sidi et al., 2015; Ramos et al., 2016), which is specially verified in estuarine areas with a wider range of the salinity (Bryan & Langston, 1992; Gonçalves & Carvalho, 2006). Despite the foregoing, in the present study, metal concentrations registered in mangrove sediments were below the Threshold Effect Level (TEL, according Environment Canada, 1999), and are not expected to cause observable adverse effects to the biota. Even though some crustacean species may accumulate metals in a higher concentration in its tissues than in the sediment (Fratini et al., 2008), these damages can be reduced due to metal tolerance, the existence of an efficient detoxification system or by tolerant strains

(Ortega et al., 2016). According to Ahearn et al. (2004), a higher content of metals can be immobilized in hepatopancreas (detoxification organ) of the decapod crustaceans, also occurring in a dissolved form (Wang & Rainbow, 2005, 2008) or as granules (Corrêa et al., 2002; Corrêa-Junior et al., 2003). However, metal concentrations in this digestive gland may exceed its detoxifying action, so that metals can be registered in other tissues, as in gills, gonads, and musculature (Barrento et al., 2009; Pinheiro et al., 2012).

In São Paulo state (Brazil), Pinheiro et al. (2013) previously verified higher metal concentrations in environmental matrices (water and sediment) in the mangroves of Cubatão when compared to those of Juréia, confirming the better conservation status of this latter and its relevance as ecological station. Moreover, there is a communication between Cubatão and São Vicente at an inner area of the Estuarine System of Santos-São Vicente, sharing the industrial effects and sanitary/waste disposal problems in these municipalities.

It is partially surprising an increasing gradient in pollution levels following the hierarchical order São Vicente (SV) > Cubatão (C) > Juréia (J), since it contrasts with our initial hypothesis that Cubatão was the most polluted mangrove area. All the mangrove areas studied show an extremely low level of metals in water, except for Hg in São Vicente, which had a concentration 2200 times higher than established by legal Brazilian limit. This fact may be explained by an emission source of Hg near this sampling site. High concentration of some more heavy metallic pollutants dissolved in water (e.g. Pb and Hg), may be increased nearby from domestic sewage and garbage dumps leachate

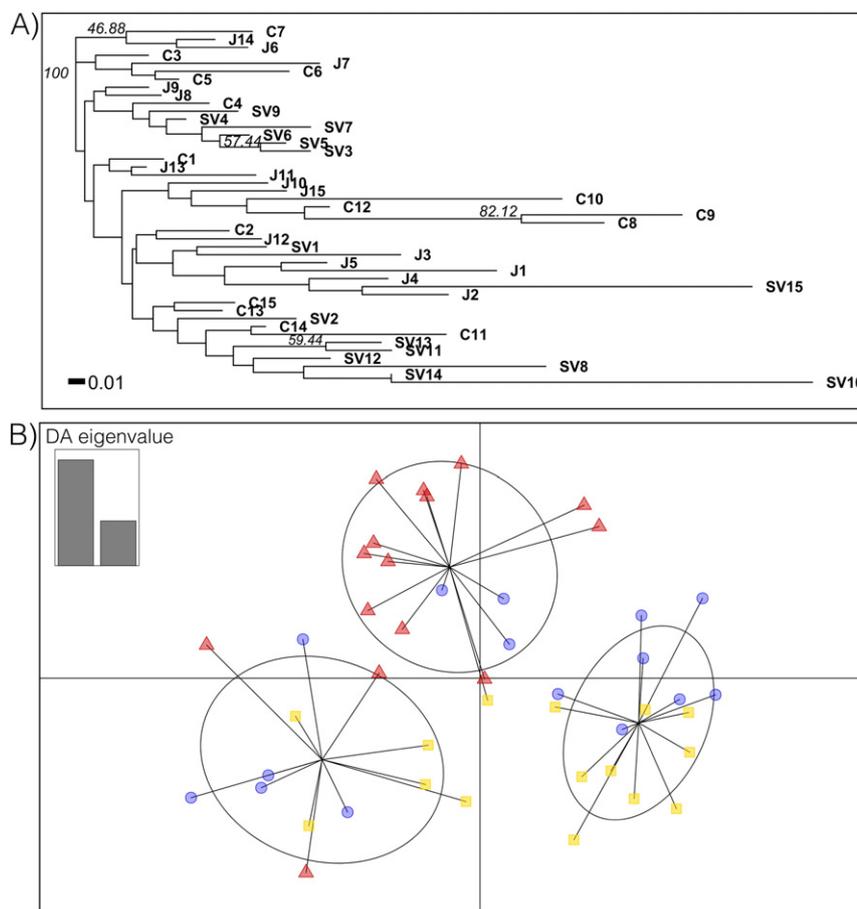


Fig. 2. Genetic structure of *Ucides cordatus* based on DNA fingerprinting from three locations (C, Cubatão; J, Juréia; and SV, São Vicente) with distinct pollution levels by metals. A) Dendrogram constructed based on Nei's genetic distance (1978) using Neighbor-Joining algorithm. Each number indicates the bootstrapping procedure support for each node (only values >40%). B) A scatterplot of the first two principal components of the Discriminant Analysis of Principal Components (DAPC) considering the number of groups that best explained our data as three. Each inferred groups is shown by the inertia ellipses and connected symbols (blue circles, Cubatão; yellow squares, Juréia; and red triangles, São Vicente).

(Sambaiatuba and Alemoa dumps – Fig. 1). According to Christensen et al. (1994) landfill leachate is characterized by some water-based solution of pollutants as heavy metals, with reports confirming higher concentrations near those contaminated sources even after 25 years of interruption of the waste storage activity (CETESB, 2001; Hortellani et al., 2005). To maximize this scenario the region of São Vicente is impacted by carbon processing from metallurgical slag handling (e.g. Branco river), a serious pentachlorophenol record caused by an improper disposal in environment, as well as cyanide, phenolic compounds and other xenobiotics (CETESB, 2001). Otherwise, the valley of Cubatão city was negatively marked by high levels of contamination, and known as ‘death valley’ in the 60s, a fact that seems to be reversing since 1984 after the pollution-control program establishment by São Paulo State Environmental Protection Agency – CETESB (Gonçalves et al., 2012).

The concentrations of all the metals measured in this study were below the reference levels, except for Hg in São Vicente. However, some crustacean species may have higher metal concentrations in their tissues than those recorded in their surrounding environment, a fact that highlights the importance of bioaccumulation in these organisms (Ugolini et al., 2004; Remaili et al., 2016). Metal concentrations in tissues and organs of *U. cordatus* from São Vicente were consistently lower than the legal levels according data informed by CETESB (2001), as well as conducted by Pinheiro et al. (2012). Nonetheless, these legal levels of each metal concentration enables the recommended consumption of this fishery resource at Estuarine System of Santos-São Vicente in that moment, a fact not confirmed more recently by Pinheiro et al. (2012). According to these authors in crabs captured in Cubatão mangroves chromium levels in muscle and hepatopancreas were three and six times higher, respectively, than permitted by Brazilian legislation. Manganese concentrations were also high, although a legal contamination limit has not been legally established.

In this study, the genetic diversity descriptive statistics revealed a similar level of genetic variation between the three populations. Contrary to our expectations, Juréia population presented a slightly lower genetic diversity regardless of the statistic analysis considered (Table 5). We arbitrarily chose this sampling location to be a reference environment and it indeed presented lower levels of heavy metals. Therefore we expected a higher genetic diversity in the Juréia populations because continuous exposure of animals to xenobiotics may cause a reduction in its genetic variability (Nadig et al., 1998; Ma et al., 2000; Ross et al., 2002; Colin et al., 2016), a process known as “genetic erosion” (Van Straalen & Timmermans, 2002; Ribeiro & Lopes, 2013). One process that may explain our result is the increase of the mutation rate by toxic compounds leading to a higher genetic variation in the locations with higher quantities of pollutants, namely São Vicente and Cubatão. The causal relationship between the metals we quantified and an increase in the genetic diversity is a hypothesis worthy of further investigation specifically designed for addressing this issue.

The description of the genetic structure patterns also provided surprising results. We expected to observe one of two contrasting patterns: absence of clear genetic structure, or that individuals from Cubatão and São Vicente (geographically closer locations) would be more related to each other than to those from Juréia. The former was justified by the lack of genetic differentiation of *U. cordatus* at local (within estuary) and regional geographic scales (including populations 3000 km apart; Oliveira-Neto et al., 2007a, b). The evidence obtained by RAPD and Restriction Fragment Length Polymorphism (Oliveira-Neto et al., 2007a) or by control region of the mitochondrial genome sequence (Oliveira-Neto et al., 2007b), genetic markers with different mutation rates and information content (dominant, co-dominant and haplotypes markers), consistently showed that *U. cordatus* populations are genetically homogeneous across a wide range of geographic scales. Conversely, our second expectation followed the rationale of a simple model of isolation-by-distance (Wright, 1943) in which due to dispersal limitation,

populations could be genetically distinct in a continuously distributed species. Thus, it was also reasonable to expect that geographically closer populations (Cubatão and São Vicente) would be more genetically related when compared to Juréia.

Contrarily to both expectations, we observed a genetic structure pattern in which only individuals from São Vicente composed a relatively delimited group according to our DAPC results (Fig. 2B). This result showed that gene flow indeed is a major biological process for *U. cordatus*, as expected according to previous studies (Oliveira-Neto et al., 2007a, b), since individuals assigned to different genetic clusters are from localities farther than 80 km apart (Cubatão and Juréia). Additionally, it provided evidence that, at fine geographic scale, there is population differentiation mainly regarding individuals from São Vicente. Considering our data, we are not able to identify specifically the biological processes (fine scale gene flow, natural selection and local adaptation, mutation rate differences – see Gama-Maia and Torres, 2016) that have generated this pattern. For instance, an unforeseen pattern of estuarine water flow, for instance, could lead to the differentiation of São Vicente population and this hypothesis could be tested in future studies using a waterscape genetics approach (Selkoe et al., 2015). Alternatively, a higher mutation rate in São Vicente could explain both the differentiation of the individuals that inhabit this area and the presence of the private band only observed in this location. Thus, one particularly promising hypothesis we consider worthy of being tested is the heavy metals genotoxic effects on *U. cordatus* to evaluate whether the levels of pollutants we observed could lead to higher mutation rates.

The lack of studies concerning metals' accumulation in mangroves biota makes it difficult to elucidate the behavior of these compounds in this estuarine ecosystem, and some of them reveal that mitochondrial DNA and microsatellite techniques can be so informative (Kim et al., 2003; Fratini et al., 2008). According to Pinheiro (unpublished data) high Hg levels were previously registered in 67% of the nine mangrove sediment samples from São Vicente, with an average higher than Brazilian reference limits. It may be linked to the accumulation of this metal in hepatopancreas and musculature of the mangrove crab *U. cordatus* and its main food source, the red mangrove (Pinheiro et al., 2012). Register of this metal in water highlights the need for careful monitoring to identify and eliminate pollutant sources.

Moreover our findings also identified an unexpected genetic structure pattern of *U. cordatus* populations, suggesting a complex evolutionary history. We argue that using this crab species as an environment quality indicator considering genetic, physiologic and chemical biomarkers (Pinheiro et al., 2013; Duarte et al., 2016) are a reasonable approach for monitoring. We further confirmed the suitability of genetic markers to investigate the effect of metal pollution on genetic diversity of this mangrove crab and to evaluate environmental remediation measures (Ma et al., 2000; Ross et al., 2002). Finally, we confirmed DNA fingerprinting using RAPD as a particularly interesting method for environmental risk assessment and monitoring, mainly in developing countries as Brazil, due to its relative low cost and fast results.

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