



Perkinsus beihaiensis (Perkinsozoa) in oysters of Bahia State, Brazil

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Abstract

This study reports the pathogen *Perkinsus beihaiensis* in oysters of the genus *Crassostrea* on the coast of the State of Bahia (Brazil), its prevalence, infection intensity and correlation with salinity. Oysters (n = 240) were collected between October and December 2014 at eight sampling stations between latitudes 13°55'S and 15°42'S. The laboratory procedures included macroscopic analysis, histology, culture in Ray's fluid thioglycollate medium (RFTM), Polymerase Chain Reaction (PCR) and DNA sequencing. PCR and sequencing have been used for the genetic identification of oysters as well. Two species of oysters have been identified: *Crassostrea rhizophorae* and *C. brasiliana*. In both oyster species *P. beihaiensis* was the only *Perkinsus* species detected. In *C. rhizophorae*, the average prevalence was 82.8% by histology and 65.2% by RFTM. In *C. brasiliana*, the prevalences were 70.5% and 35.7%, respectively. The higher prevalence of *P. beihaiensis* in *C. rhizophorae* was probably influenced by salinity, with which was positively correlated ($r > 0.8$). In both oysters, *P. beihaiensis* was located mainly in the gastric epithelium. The infection was generally mild or moderate, without apparent harm to the hosts, but in cases of severe infection, there was hemocytical reaction and tissue disorganization. The generally high prevalence in the region suggests that oysters should be monitored with respect to this pathogen, especially in growing areas.

Keywords: bivalves, diseases, pathology, salinity.

Perkinsus beihaiensis (Perkinsozoa) em ostras do Estado da Bahia

Resumo

Este estudo relata o patógeno *Perkinsus beihaiensis* em ostras do gênero *Crassostrea* no litoral do Estado da Bahia (Brasil), sua prevalência, intensidade de infecção e correlação com a salinidade. As ostras (n = 240) foram coletadas entre outubro e dezembro de 2014 em oito estações amostrais entre as latitudes 13°55'S e 15°42'S. Os procedimentos laboratoriais incluíram análise macroscópica, histologia, cultivo em meio de tioglicolato de Ray (RFTM), reação em cadeia da polimerase (PCR) e sequenciamento de DNA. PCR e sequenciamento foram também utilizados para a identificação genética das ostras. Foram identificadas duas espécies de ostras: *Crassostrea rhizophorae* e *C. brasiliana*. Em ambas as espécies, *P. beihaiensis* foi a única espécie de *Perkinsus* detectada. Em *C. rhizophorae*, a prevalência média foi de 82,8% por histologia e de 65,2% por RFTM. Em *C. brasiliana*, as prevalências foram de 70,5% e 35,7%, respectivamente. A maior prevalência de *P. beihaiensis* em *C. rhizophorae* foi provavelmente influenciada pela salinidade, com a qual este apresentou correlação positiva ($r > 0,8$). Em ambas as espécies, *P. beihaiensis* esteve localizada principalmente no epitélio gástrico. A infecção foi geralmente leve ou moderada, sem danos aparentes aos hospedeiros, mas em casos de infecção severa, houve reação hemocitária e desorganização de tecidos. As prevalências geralmente altas na região sugerem que as ostras devam ser monitoradas com relação a este patógeno, principalmente em áreas de cultivo.

Palavras-chave: bivalves, doenças, patologia, salinidade.

1. Introduction

It is known that for the success of mollusks farming, management and monitoring of diseases are essential. According to review by Boehs et al. (2012), in the last decade have been recorded various pathogens in bivalve mollusks from the Brazilian coast, including representatives of bacteria, protozoa, fungi and metazoans. Nevertheless,

in general there is still few information on diseases in bivalves of economic interest, both in Brazil and throughout South America, particularly with regard to the life cycle of pathogens and the interaction of these with their hosts.

Among the pathogens already reported at Brazilian coast, is the genus *Perkinsus* Levine, 1978 (Perkinsozoa),

which affects mollusks in various parts of the world, including some bivalves of great economic interest, such as oysters. Seven species are described worldwide, two of which (*P. marinus* and *P. olseni*) are notifiable to the World Organization for Animal Health (OIE, 2012). In Brazil, four species were recorded in the past five years, all in the Northeast: *P. beihaiensis*, *P. marinus*, *P. olseni* and *P. chesapeakei* (Dantas-Neto, 2015).

Perkinsus beihaiensis Moss, Xiao, Dungan and Reece, 2008 was first described in southern China (Moss et al., 2008) in *Crassostrea ariakensis* (Fujita, 1913) and in *C. hongkongensis* Lam and Morton, 2003. Sanil et al. (2012) reported this species in *C. madrasensis* (Preston, 1916) in India. In Brazil, *P. beihaiensis* was recorded in species of *Crassostrea* in northeastern Brazil, in the states of Ceará (Sabry et al., 2009, 2013), Paraíba (Queiroga et al., 2015) and Bahia (Luz and Boehs, 2016), as well as in *Anomalocardia brasiliiana* (Gmelin, 1791) (Veneridae) (Ferreira et al., 2015) from Ceará. Overall, there is still few information on this species in terms of their ecology, potential hosts, and risks for the production of mollusks.

The oysters *Crassostrea rhizophorae* (Guilding, 1828) and *C. brasiliiana* (Lamarck, 1819) are important resources for consumption and sale of extractive communities on the Brazilian coast. The cultivation of these mollusks is already practiced in several locations. This study reports the presence of *P. beihaiensis* in these species at Bahia State, with information about pathology, infection intensity, prevalence, and correlation with salinity.

2. Material and Methods

2.1. Collections

Sampling was done at eight sampling stations (St1-St8) located between latitudes 13°55'S and 15°42'S, in a stretch of about 200 km. Stations St1-St4 were located in the municipality of Camamu, St5 and St6 in Ilhéus and St7 and St8 in Canavieiras (Figure 1, Table 1). Samples were collected between October and December 2014, under authorization number 20912-3, granted by the Chico Mendes Institute for Biodiversity Conservation – ICMBio, Brazil.

The specimens were removed with the help of a knife, from the trunks or roots of red mangrove *Rhizophorae mangle* L. (St1-St5 and St8) or from rocks and concrete structures (St6 and St7), always during low tide periods. At each sampling station were collected 30 oysters. These were brought in buckets containing a small portion of sea water and processed in a period of 6 hours after their collection. Water temperature and salinity were recorded in each day of sampling using respectively a standard mercury thermometer and an optical refractometer Atago S/Mill.

2.2. Laboratory procedures

The oysters were measured for their long axis (height = dorsal-ventral axis), according Galtsoff (1964). After opened, the specimens were analyzed macroscopically for the presence of clinical signs of *Perkinsus* sp., such as weight loss, pale appearance, shrinkage of mantle and

presence of pustules, as previously reported by Bower et al. (1994) and by Bondad-Reantaso et al. (2001).

Cross section of about 5 mm was made on each specimen, which was fixed in Davidson solution (Shaw and Battle, 1957) for 24-30 hours. The tissues were processed by classical histological technique, including dehydration in a series of increasing alcohol concentrations and embedding in paraffin. Sections of thickness 5 µm were cut using a microtome and were stained with Harris haematoxylin and eosin (H&E) (Howard et al., 2004). The histological sections were analyzed under Olympus light microscope at 40 and 100 ×. After analysis, the slides were deposited in the Marine Mollusks Laboratory (LMM) at the State University of Santa Cruz, under the care of the first and last authors.

Gills and rectum of each oyster (n = 240) were removed and incubated for culturing in Ray's fluid thioglycollate medium - RFTM, developed by Ray (1966), following international protocol (OIE, 2012) with addition of antibiotics (Penicillin and Streptomycin) and antifungal (Nystatin), to inhibit the proliferation of microorganisms. The culture medium was kept in the dark for 7 days at room temperature (20-25 °C). The tissues were macerated and stained in 3% iodine solution for visualization under a light microscope.

Samples with positive result in RFTM were subjected to PCR. Total DNA extraction was performed from subsamples of the digestive gland and gills, which were preserved in 95% ethanol. For this step, it was used DNazol (*Invitrogen*®) following the manufacturer's protocol and phenol-chloroform protocol proposed by OIE (2012). For the PCR reactions was used the primer pair PerKITS 85/750, exclusive to members of the genus *Perkinsus* (except for *P. qugwadi*) (Casas et al., 2002). The PCR reactions were performed in volumes of 12.5µL, containing 100-200ng of genomic DNA, 1x PCR buffer concentrate, 1.5 mM of MgCl₂, 2.5 mM of dNTP, 10 picomol of each primer and 1 U Taq DNA polymerase (*Invitrogen*®). The DNA of *P. beihaiensis* identified previously by Luz and Boehs (2016) was used as positive control. The protocol included: DNA denaturation at 94 °C for 5 minutes; 35 cycles of amplification at 94 °C for 40 seconds, 60 °C for 40 seconds and 72 °C for 1 minute followed by a final extension at 72 °C for 7 minutes. The PCR products were separated on 1.5% agarose gels, stained with *Syber Safe*® and visualized using UV light. Twenty positive samples were randomly selected for DNA sequencing. The samples were sequenced by ACTGene Molecular Analysis (Porto Alegre, Brazil). The obtained sequence was deposited in GenBank.

2.3. Data processing

The prevalence of the pathogen was calculated as the number of infected oysters divided by the total number of analyzed oysters (Bush et al., 1997). The infection intensity (analyzed from RFTM), was calculated according to the scale developed by Ray (1954) and modified by Sabry et al. (2009), at the following levels (L): Nil infection: zero hyphospores on the whole slide (100 ×) (= L0); Very

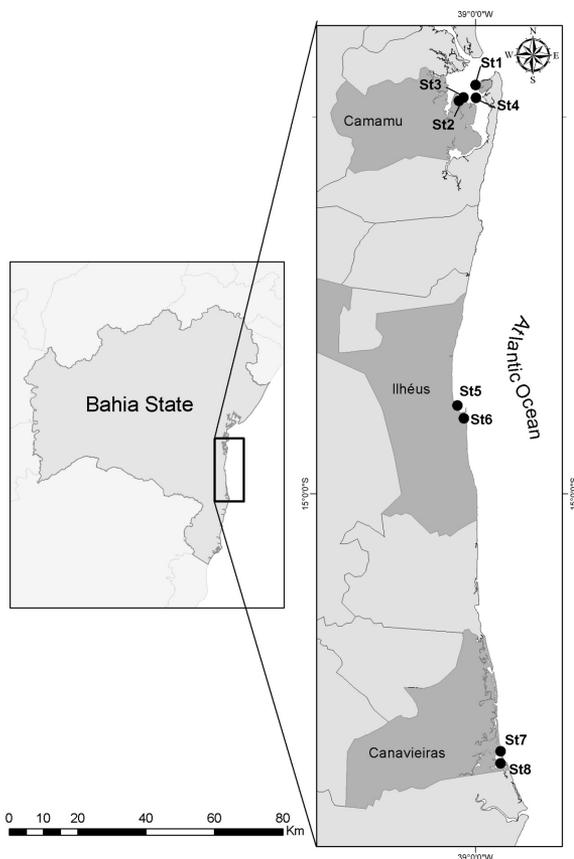


Figure 1. Map of the study area, indicating the collection stations (St1-St8).

Table 1. Information on sampling stations (coordinates, location and specific temperature and salinity data) and results of prevalence (P) of *Perkinsus beihaiensis* in two species of oysters in Bahia, Brazil, for 240 samples collected between October and December 2014 and analyzed by histology and by Ray’s fluid thioglycollate medium (RFTM), as follows: - Absence of the species or data; Cb = *Crassostrea brasiliana*; Cr = *Crassostrea rhizophorae*; S = salinity; T = temperature.

Sampling stations	Coordinates	Localities	T (°C)	S (‰)	P (%) – Histology		P (%) - RFTM	
					Cr	Cb	Cr	Cb
St1	13°55'. 818"S, 39°. 00.963"W	Camamu	29	36	100	-	93.3	-
St2	13°56'. 848"S, 39°. 00.867"W	Camamu	28	36	96.6	-	86.6	-
St3	13°57'. 779"S, 39°. 02'. 948"W	Camamu	27	36	86.6	-	83.3	-
St4	13°57'. 962"S, 39°. 00.022'W	Camamu	27	36	86.6	-	86.6	-
St5	14°46'. 113"S, 39°3'. 116"W	Ilhéus	26	18	60.7	-	25.0	-
St6	14°48'10.21"S, 39°2'20.79"W	Ilhéus	28	30	66.6	58.3	16.6	4.16
St7	15°41'. 075"S, 38°56'38.06"W	Canavieiras	29	20	-	86.6	-	76.6
St8	15°42'41.41"S, 38°55'56.17"W	Canavieiras	29	25	-	66.6	-	26.6
Mean ± Standard Deviation	-	-	27.8 ± 1.12	29.6 ± 7.67	82.8 ± 15.9	70.5 ± 14.5	65.2 ± 34.6	35.7 ± 37.0

mild infection: up to 10 hyphospores on the whole slide (100 ×) (= L1); Mild infection: 11-100 hyphospores on the whole slide (100 ×) (= L2); Moderate infection: at least 40 hyphospores observed in 10 different fields (400 ×) (= L3); Severe infection: more than 40 hyphospores observed in 10 different fields (400 ×) (= L4).

Pearson correlation analysis was used to relate the prevalence data and infection intensity with salinity at the sampling stations and species and a t test was associated to show significant differences. We used the ASSISTAT, version 7.7 beta. A multivariate ordination (Principal Component Analysis - PCA), generated by MVSP software was used to show graphically the correlations. The confidence level was 95%. To ensure high reliability of the salinity values of each place, for these analyzes were used the average of the values obtained in this study with records made over a year in each region, as follows: Lenz and Boehs (2011) and Luz and Boehs (2015) at Camamu, Boehs et al. (2010) at Ilhéus and Zeidan et al. (2012) at Canavieiras.

3. Results

The temperature was between 26 and 29 °C (Mean 27.8 °C ± 1.12) and the salinity between 18 and 36‰ (Mean 29.6‰ ± 7.67) (Table 1).

In molecular analysis to confirm the pathogen, the amplicons showed 700 bp, which is the expected size for protozoa of the genus *Perkinsus*. Sequencing demonstrated that this was *P. beihaiensis*, with 98 to 100% similarity with the sequences available at NCBI, as deposited by Sanil et al. (2012) and Sabry et al. (2013), to India and the Northeast Brazil, respectively. The obtained sequence was deposited in GenBank with number KX923545.

In an infected oyster collected in station St7, in the municipality of Canavieiras, small pustules measuring approximately 1 mm in diameter, yellow-white color, were observed in the mantle and gills. In the other specimens (n = 239) there was no macroscopic signs of the pathogen.

Trophozoites, typically with a large vacuole and peripheral nucleus, like schizonts (trophozoites dividing) were observed in both species. Trophozoites had sizes between 3 and 10 µm (Mean 6.8 µm ± 2.03; n = 240) and were seen mainly in the digestive tract, preferably in the stomach epithelium, but these were observed also in the mantle and adductor muscle, in low-intensity. In the digestive tract, in cases of severe infection, there was moderate hemocytic infiltration and tissue disorganization.

Considering all the sampling stations, the overall mean prevalence in the histology was 82.8% ± 15.9 in *C. rhizophorae* and 70.5% ± 14.5 in *C. brasiliiana* (Table 1).

The results obtained by RFTM showed hyphospores of 2 to 75 µm, being that the lowest values were observed in cultures from station St3 (2-4 µm) and station St7 (2-8 µm). The mean prevalence by RFTM was 65.2% ± 34.6 in *C. rhizophorae* and 35.7% ± 37.0 in *C. brasiliiana* (Table 1). The infection was moderate (L3) to severe (L4) in stations St1-St4 and in station St7. In the other stations dominated

the null level of infection (L0) or oysters had very mild (L1) or mild infection (L2) (Figure 2 and Figure 3).

Statistical analyzes demonstrated a positive correlation between the prevalence of *P. beihaiensis* and salinity ($r = 0.8798$; $p < 0.01$). The salinity was also positively correlated with the infection levels: L2 ($r = 0.7437$; $p < 0.05$), L3 ($r = 0.7261$; $p < 0.05$) and L4 ($r = 0.8841$; $p < 0.01$) and negatively correlated with the level L0 (absence of infection) ($r = -0.7526$; $p < 0.05$). The localities with high prevalence were also strongly correlated with severe levels of infection ($r = 0.8803$; $p < 0.01$).

At the stations with high salinity (St1-St4 where only occurred *C. rhizophorae* and St7 where only occurred *C. brasiliiana*), were observed both high prevalence of *P. beihaiensis* as well high levels of infection. The components 1 and 2 of the Principal Component Analysis - PCA explained 86.5% of the total variability of the data (Figure 3).

4. Discussion

In both species of oysters, *Perkinsus beihaiensis* was detected as the single species of *Perkinsus* present, with preferential localization in the digestive epithelium, especially in the stomach, and in low numbers, it was also seen in the mantle and adductor muscle. The preferred location of *P. beihaiensis* in the digestive epithelium has been reported in previous studies on the genus *Crassostrea* (Moss et al., 2008; Sanil et al., 2012; Sabry et al., 2013; Queiroga et al., 2015; Luz and Boehs, 2016).

Oysters with severe infection (level 4, Ray's scale) showed disorganization of gastric and intestinal epithelia and were assessed as impaired by the infection. In turn, both in the mantle and adductor muscle, as in oysters that had a low level of infection, there was no tissue change. Sanil et al. (2012) reported destruction of digestive tubules in some cases of infection by *P. beihaiensis*. Moss et al. (2008) reported epithelial lesions and necrosis in the stomach, intestine and digestive gland. According to Villalba et al. (2004), in more severe *Perkinsus* infections, such as those caused by *P. marinus*, maceration of the adductor muscle may occur, with consequent involuntary opening of valves, weight loss, shrinkage of the mantle, impairment of the immune system and death of the host.

Comparing the results of prevalence of *P. beihaiensis* of this study with others obtained for *Crassostrea* spp. in Brazil, it was found that the values obtained on the coast of Bahia are higher. Considering only the results obtained by histology, Sabry et al. (2013) found prevalences between 3 and 20% in the State of Ceará, in a region with a mean salinity of 33.1‰ and Queiroga et al. (2015) found mean prevalence of 48.9% in the State of Paraíba, but did not mention salinity. In the State of Bahia, Luz and Boehs (2016) previously found an overall mean of 93.3% in the Camamu Bay, in a region with a mean salinity of 28.4‰. In this study, the salinity values were generally above 20‰, and the overall mean prevalence of *P. beihaiensis* was 78.7%, considering the two species of oysters.

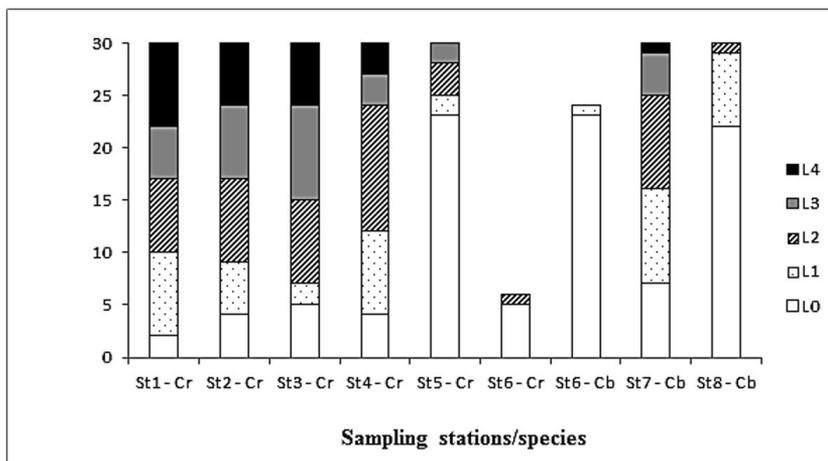


Figure 2. Absolute frequency of infection levels of *Perkinsus beihaiensis* of *Crassostrea rhizophorae* (Cr) and *C. brasiliana* (Cb) from coast of Bahia, observed under light microscopy after cultivation in Ray’s fluid thioglycollate medium - RFTM. n = 240. Infection levels according to the classification of Ray (1954), where: L0 = Nil infection; L1 = Very mild infection; L2 = Mild infection; L3 = Moderate infection; L4 = Severe infection.

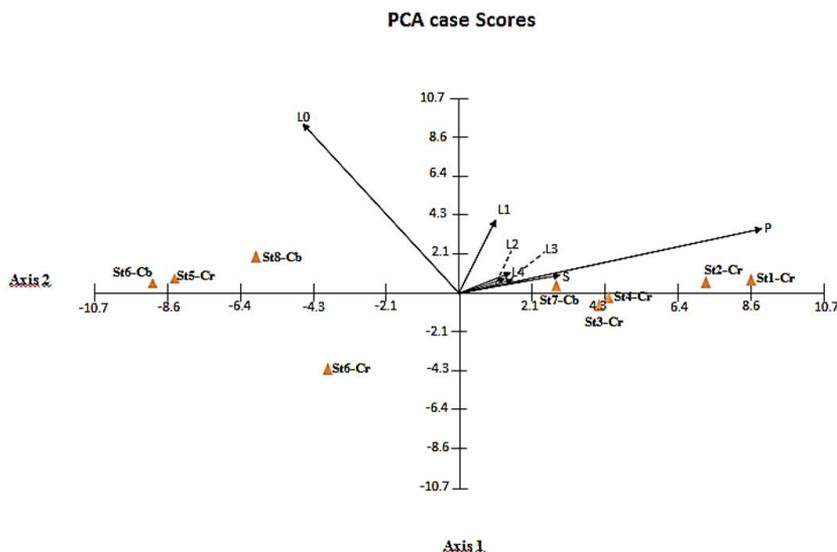


Figure 3. Graphic result of Principal Component Analysis (PCA), showing a positive correlation between the prevalence (P) and intensity of infection (levels L0-L4) of *Perkinsus beihaiensis* with salinity (S) in the oysters: *Crassostrea rhizophorae* (Cr) and *C. brasiliana* (Cb). St = sampling stations (St1-St8).

Our results showed positive correlations between salinity and both prevalence and infection intensity, which reinforces results from previous studies (Volety, 2008; La Peyre et al., 2010). According Villalba et al. (2004), in some places, the seasonal variations of salinity and of temperature modulate the occurrence of species of the genus *Perkinsus*. In Brazil, this pathogen has been detected so far only in the Northeast region of the country where, coincidentally, temperatures are high throughout the year.

With reference to the prevalence and levels of infection generally higher in *C. rhizophorae* in relation to *C. brasiliana*, we conclude that this result was in

principle affected by salinity levels, since the first species was collected in more saline places. However, a higher prevalence among *C. rhizophorae* was also seen in station St6 where there was the simultaneous occurrence of the two species, suggesting that perhaps *C. rhizophorae* is more severely affected by *P. beihaiensis* than *C. brasiliana*. Thus, ecological differences between the two oyster species (*C. rhizophorae* inhabits intertidal levels slightly higher than *C. brasiliana*) may have influence on susceptibility to *P. beihaiensis*.

As for the higher sensitivity of the detection of *P. beihaiensis* in histology relative to RFTM, we conclude

that this difference is due to the use of different organs in each of the techniques. The protocol currently used (OIE, 2012) indicates the use of the rectum and the gills for RFTM, where seems to be a lower concentration of the pathogen in relation to the stomach, organ usually included in the histological analysis. This difference was already observed in previous studies (Sabry et al., 2009, 2013; Ferreira et al., 2015; Luz and Boehs, 2016). We suggest the use of same organs in histology and RFTM (and also in PCR), to obtain greater uniformity in the diagnosis.

The presence of pustules on an infected oyster in station St7 indicates the possibility of using presumptive diagnosis as a diagnostic complement, however, as we observed the event in a single oyster, the reliability of occurring macroscopic evidence of the pathogen in question is still low.

It is concluded that *P. beihaiensis* does not affect massively the health of the oysters in the region at this moment. However, in order to the development of oyster farming, periodic monitoring of this pathogen should be emphasized.

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