SHORT COMMUNICATION

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Life history told by a whale-louse: a possible interaction of a southern right whale *Eubalaena australis* calf with humpback whales *Megaptera novaeangliae*

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Abstract

Southern right whales (*Eubalaena australis*) are known to host three species of whale-lice, *Cyamus gracilis*, *Cyamus ovalis* and *Cyamus erraticus*. Such cyamids usually are generalists in toothed whales (Cetacea: Odontoceti) and host-specific in baleen whales (Cetacea: Mysticeti), and because they have no free-swimming stage, transmission only occurs by contact between whales. One southern right whale stranded at the southeastern coast of Brazil was found parasitized by a different species of cyamid. Over 300 specimens were collected and the only species identified based on morphological and molecular data was *Cyamus boopis*, a typical ectoparasite of humpback whales (*Megaptera novaeangliae*). This finding is the first record of *C. boopis* on the southern right whale. Both *E. australis* and *M. novaeangliae* are found in Brazilian waters and the presence of humpback's whale-louse together with the lack of the three specific parasites of right whales suggest an interspecific interaction between these whales based on the parasite's biology.

Keywords: Whale-lice, Interspecific interaction, Cyamus, Mysticeti

Background

Southern right whales *Eubalaena australis* (Desmoulins, 1822), which are observed in Brazil typically from May to December with peaks during August to October, prefer to use shallow and protected areas of the state of Santa Catarina (southern Brazil) as their wintering ground [1, 2]. Sightings of *E. australis* in southeastern and northeastern Brazil are reported occasionally [3–6] from early July to late October [3]. Although sightings of *E. australis* are periodic, strandings in southeastern Brazil are rare: only eight strandings, most of which were calves, were recorded between 1936 and 2001 [3]. Figueiredo et al. [7] reports on the last records of right whales in southeastern Brazil.

An unusual stranding of a southern right whale calf is reported here. On January 17th, 2004, a whale calf became stranded at Praia de Itaúna, Saquarema, which is located on the east coast of the state of Rio de Janeiro, Brazil, but it died shortly afterwards. The whale's body was in a carcass state of Geraci 1 (fresh) [8] for about 2 h and 30 min prior to the necropsy. The subsequent necropsy of this specimen revealed over 200 specimens of epibionts, including pedunculate barnacles *Conchoderma auritum* (Linnaeus, 1767) and cyamids.

The cyamids or whale-lice are crustacean parasites, which feed on the skin of cetaceans, that are represented by seven genera and 29 species [9–11]. They are considered generalists in toothed whales and dolphins (Odontoceti) and host specific in baleen whales (Mysticeti) [12, 13], therefore, new records of cyamids from slow moving great whales are unlikely, since most studies on the commercially valuable whales cyamid fauna was driven during the whaling period [9, 10]. For this reason, current

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works on new species and distributional records of Cyamidae are restricted to odontocetes parasites [10, 14].

Right whales are known to host three species of cyamids: *Cyamus ovalis* Roussel de Vauzème, 1834, *Cyamus erraticus* Roussel de Vauzème, 1834, and *Cyamus gracilis* Roussel de Vauzème, 1834 [15–17]. Each species has a different spatial distribution on the right whale's body [12, 15, 18].

Young calves of right whales usually present *Cyamus erraticus* on the head and these lice patches tend to disappear within a few months and migrate to soft skin such as genital and mammary slits in right whale adults, whereas the head patches are substituted by *Cyamus ovalis* and *Cyamus gracilis* [15, 17]. Thus, the calf's first louse infestation occurs by vertical transmission from its mother and subsequent horizontal transfer is expected to occur infrequently, which might reflect patterns of behavioral interactions among whales [17]. According to Page and Charleston [19], horizontal transfer is one of the events that contributes to patterns of association between hosts and parasites.

In this study, we report an atypical whale louse parasitizing a southern right whale stranded on the Southeastern coast of Brazil and we discuss the possible implications of this cyamid host change.

Methods

Cyamid specimens were collected randomly from the head of the stranded calf, fixed and preserved in ethanol 70%, and deposited at the Crustacean Collection of the Museu Nacional, Rio de Janeiro (MNRJ). Cyamids were absent in other body parts of the stranded calf. All specimens were observed with Zeiss Discovery V12 AxioCam MRc stereomicroscopes. Juveniles were not identified. One specimen was chosen randomly for scanning electron microscopy (SEM). Specimen preparation for SEM was made following the protocol adapted from Felgenhauer [20], dried at critical point, and sputtercoated with gold. Micrographs were taken with SEM JEOL JSM-6390LV at Laboratório Central de Microscopia Eletrônica (LCME/UFSC). Additionally, the cyamids were compared with type series of *Cyamus boopis* Lütken, 1870,

deposited at the Zoologisk Museum, Copenhagen (ZMUC), and comparative material of *Cyamus ovalis*, *Cyamus erraticus* and *Cyamus gracilis* from MNRJ and ZMUC (Table 1).

Five random specimens were subjected to molecular analyses. Total genomic DNA was obtained using a modified CTAB extraction [21]. A fragment of the cytochrome oxidase I gene (COI) was amplified using the Jercy and Patcy primers [17]. Amplification reactions included approximately 30 ng of genomic DNA, 1 U of GoTaq Flexi DNA polymerase (Promega), 3 µL of Green GoTaq Flexi Buffer (5×), 0.2 mM of dNTPs, 2.5 mM of MgCl2, 0.3 μ M of each primer, and 4 μg of BSA in a final volume of 15 μL. PCR conditions were: 1 cycle of 2 min at 95 °C followed by 35 cycles of 20 s at 95 °C, 30 s at 50 °C, and 50 s at 72 °C. All PCR products amplified were purified using the Agencourt AMPure PCR purification kit and sequenced in both directions using ABI 3500 automated DNA sequencers (Applied Biosystems). The sequences obtained were edited using Seqman 7.0 (DNAstar Inc.), aligned by the Clustal W algorithm implemented in Mega 6 [22], and deposited in GenBank (accession numbers KX100157 to KX100161). Previously published sequences for representatives of seven Cyamus species were included in the alignment. To identify each sample to species a Bayesian phylogenetic tree was inferred using MrBayes 3.2.6 (MCMC ngen = 10×10^6 generations, nchains = 4, nruns = 2, samplefreq = 1000, burninfrac = 0.25) [23]. In this analysis, an HKY + G + I model was chosen to be the best model as indicated by Bayesian information criterion (BIC) implemented in Mega with Caprela scaura Templeton, 1836 as the outgroup. Additionally, the intraspecific diversities and interspecific divergences were estimated in Mega, using the K2P distance model.

Results

Morphology

All 327 cyamid specimens were morphologically identified as *Cyamus boopis* Lütken, 1870, typical ectoparasite of humpback whales *Megaptera novaeangliae* (Borowski, 1781) [16, 24]. Based on the present data, this is a first record of *C. boopis* parasitizing *Eubalaena australis*.

Table 1 Comparative material of cyamids from Eubalaena sp

Ref. no.	Cyamid	Host	Stranding location	Date	
MNRJ 67664——67667	Cyamus erraticus	Eubalaena australis	Florianópolis, Brazil (fishing net)	Ago-14	
MNRJ 67668——67672	Cyamus ovalis	Eubalaena australis	Florianópolis, Brazil (fishing net)	Ago-14	
MNRJ 67673——67677	Cyamus gracilis	Eubalaena australis	Florianópolis, Brazil (fishing net)	Ago-14	
ZMUC CRU-7570	Cyamus erraticus	Eubalaena glacialis	British Columbia, Canada	1951	
ZMUC CRU-7679	Cyamus ovalis	Eubalaena glacialis	Iceland	1892	

Cyamus boopis was recently redescribed based on the type material, which was used for comparison on the present study, as well as compared with over 1000 *C.boopis* from sixteen humpback whales stranded off the Brazilian coast [25]. The identification of *C. boopis* was based on diagnostic characters as the presence of ventral acute processes on pereonites 5–7 and shape of accessory gills (Fig. 1).

Diagnostic features were compared with *Eubalaena*'s typical whale-lice, and it differs from *C. ovalis* by the shape of accessory gills, which are larger and asymmetrical in the later species. *Cyamus boopis* also presents a slender body compared to *C. ovalis*, however broader than *C. gracilis*. *Cyamus gracilis* lacks acute ventral processes on the pereonites 5–7, while *C. boopis* shows a 1/1/1 pairs of acute ventral processes arrangement on pereonites 5–7, respectively, and *C. erraticus* presents a 2/2/1 arrangement. Due to the cyamid succession in right whales, it was expected that the calf would host blooms of *C. erraticus* on its head, instead, the only species found was *C. boopis*.

Barcoding

Our molecular analyses suggest that the five sequenced specimens of the right whale calf belong to the species *Cyamus boopis*, thereby corroborating the morphological results. In the Bayesian phylogenetic tree, the analysed specimens are not closely related to none of the reported

Eubalaena's specific cyamid species, contrariwise, they formed a monophyletic group well supported with the sequences of *Cyamus boopis* (Fig. 2). Furthermore, the value of divergence between the analysed specimens and *C. boopis* was 1.4%, value that is within the rank of intraspecific distances (0.9–9.0%) and far below of the rank of interspecific distance between species of *Cyamus* (14.2–21.0%, Table 2).

Discussion

Based on morphological and molecular data, the only species that was found in the right whale calf was *Cyamus boopis*, a typical ectoparasite of humpback whales (*Megaptera novaeangliae*). The fact that we only found *Cyamus boopis* suggests that the calf might have interacted likely by chance with a humpback whale. Thus, the high concentration of gravid females of *C. boopis* (n = 73) found on the right whale calf agrees with Rowntree [15] assumption that 'gravid females may move to areas where there is less competition for their younger to "launching" sites where there is more contact between whales and the cyamids have a greater chance of moving to a new host.'

Cyamus boopis is the only cyamid that parasitizes humpback whales, and it colonizes areas with reduced water flow, such as the blowhole, wounds, around barnacles and in the ventral grooves [15]. Also, *C.boopis* is only found in humpback whales. Unlike the right whales lice, there is no succession of cyamid species in humpbacks,

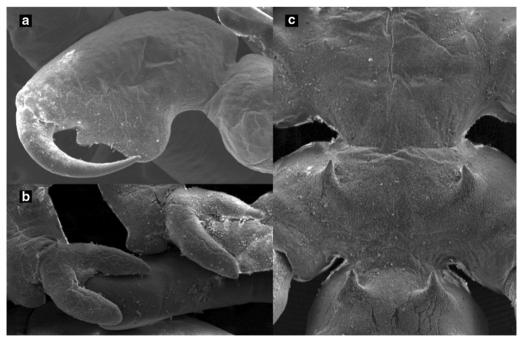


Fig. 1 Morphological diagnostic features of *Cyamus boopis* found in the southern right whale GEMM-Lagos —0051. **a** Shape of gnatopod 2; **b** shape of accessory gills; **c** number of ventral acute processes on pereonites 5, 6 and 7

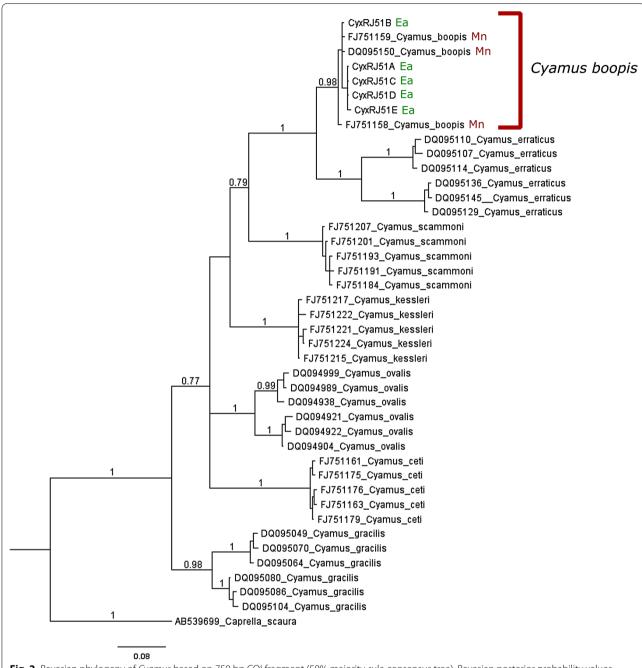


Fig. 2 Bayesian phylogeny of *Cyamus* based on 750 bp COI fragment (50% majority-rule consensus tree). Bayesian posterior probability values above 70% are shown. *Cyamus boopis* is shown in the *red bracket* and the CyxRJ51 samples correspond to *Cyamus boopis* collected from *Eubalaena australis* calf GEMM-Lagos 0051. Ea = *Eubalaena australis*; Mn = *Megaptera novaeangliae*

and the only movement of *C. boopis* observed by Rowntree [15] is between small patches of white and black skin within a humpback, which might be driven by the male competition for mates.

Humpbacks are commonly sighted off the east coast of Rio de Janeiro during winter and spring (July–October), though mother and calves of humpback whales as well as yearlings have also been observed along the same coast during austral summer months (December–March) (Salvatore Siciliano, pers. observ.). Reports about interactions between two mysticetes species are less common than interactions between mysticetes and odontocetes [26, 27].

Since cyamids do not have a swimming larval stage and the transmission only occurs by contact between whales

Table 2 Summary of average Kimura two-parameter distances for Cyamus species

	CyxRJ51	C. boopis	C. ceti	C. scammoni	C. kessleri	C. ovalis	C. gracilis	C. erraticus
C. boopis (CyxRJ51)	0.006							
C. boopis	0.014	0.009						
C. ceti	0.197	0.203	0.009					
C. scammoni	0.169	0.165	0.171	0.010				
C. kessleri	0.171	0.168	0.188	0.152	0.013			
C. ovalis	0.173	0.172	0.175	0.161	0.158	0.057		
C. gracilis	0.186	0.181	0.182	0.162	0.175	0.162	0.055	
C. erraticus	0.142	0.142	0.210	0.200	0.197	0.195	0.207	0.090

Intraspecific diversities (on the diagonal) and interspecific divergences (below the diagonal)

[15, 28], the presence of over 300 Cyamus boopis suggest that the right whale calf unexpected contacted with a humpback whale for a considerable period of time, and the absence of three typical right whale cyamids species raises an inquiry about the nature of the interaction between this right whale calf and a humpback whale and whether it is possible for different baleen whale species to nurse each other. Thus, the transmission of cyamids from Eubalaena to Megaptera whales just recently could be related to the population expansion of humpback whales in the Southwestern Atlantic Ocean [29] and of southern right whales recovering from exploitation [30], which increases the chance of a causal encounter between these two baleen species. While the origin of the cyamid movement onto the right whale calf cannot be known, it is indicative of the first movement of C. boopis to a new baleen whale host species.

Our findings show that even though the mysticetes' fauna of cyamids might be well described [31], new occurrences are possible. More information about the cyamid hosts' ecology may yet be revealed by documentation of these ectoparasites. Nonetheless, more effort should be given to collection and sampling of ectoparasites in any dead whale carcass. Collections should be made from fresh carcasses, sampling all the specimens as possible in different areas of the host, and placed into preserved ethanol P.A. for further morphological and molecular studies. This may reveal other new cyamid occurrence and transfers.

Abbreviations

GEMM: Grupo de Estudos de Mamíferos Marinhos; MNRJ: Museu Nacional, Rio de Janeiro; SEM: scanning electron microscopy; LCME/UFSC: Laboratório Central de Microscopia Eletrônica/Universidade Federal de Santa Catarina; ZMUC: Zoological Museum of the University of Copenhagen; COI: cytochrome oxidase I; PCR: polymerase chain reaction.

Authors' contributions

TIA conceived the project and performed the taxonomic identifications and molecular experiments. GTRR designed and performed the molecular experiments, and interpreted the data. SS collected the material and field data and was a major contributor in writing the manuscript. CSS performed the taxonomic identifications. All authors read and approved the final manuscript.

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Competing interests

The authors declare that they have no competing interests.

Availability of data and materials

All data generated or analysed during this study are included in this published article. DNA and RNA sequences are deposited at GenBank mentioned by the accession numbers in the main text.

Consent for publication

Dr. Salvatore Siciliano, co-author of this manuscript, consent the publication of personal observation and Fig. 1 source. Contact of the author for consent form: gemmlagos@gmail.com

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