

# Gastrointestinal parasite fauna of Emperor Penguins (*Aptenodytes forsteri*) at the Atka Bay, Antarctica

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**Abstract** In general, the knowledge on parasites infecting Antarctic birds is scarce. The present study intends to extend the knowledge on gastrointestinal parasites of Emperor Penguins (*Aptenodytes forsteri*) at the Atka Bay, Antarctica. Fecal samples of 50 individual Emperor Penguins were collected at the Atka Bay and analyzed using the sodium-acetate-formaldehyde (SAF) method for the identification of intestinal helminth eggs and/or protozoan parasite stages. In addition, coproantigen ELISAs were performed to detect *Cryptosporidium* and *Giardia* infections. Overall, 13 out of 50 penguins proved parasitized (26 %). The following stages of gastrointestinal parasites were identified: One *Capillaria* sp. egg, *Tetrabothrius* spp. eggs, *Diphyllobothrium* spp. eggs, and proglottids of the cestode *Parorchites zederi*. The recorded *Capillaria* infection represents a new host record for Emperor Penguins. All coproantigen ELISAs for the detection of *Cryptosporidium* spp. and *Giardia* spp. were negative. This paper provides current data on parasites of the Emperor Penguin, a protected endemic species of the Antarctica.

**Keywords** Emperor Penguin · *Aptenodytes forsteri* · Endoparasites · Antarctica

## Introduction

Only nine out of 156 currently recognized avian families are specialized as seabirds such as the penguins (Spheniscidae) (Lauckner 1985). The Emperor Penguin, *Aptenodytes forsteri*, is the most southerly situated penguin species on earth and a top endemic predator in the marine environment of the Antarctic region (Cherel and Kooyman 1998). The population size of the Atka Bay Emperor Penguin colony was composed of approximately 9,600 animals, estimated via satellite pictures by the British Antarctic Survey (BAS) (Fretwell et al. 2012). Moreover, the Atka Bay colony currently belongs to the ten largest colonies of Emperor Penguins of the Antarctic region. In general, birds of the genus *Aptenodytes* are the heaviest penguins (up to 40 kg) and the deepest divers (250 m, some deep dives up to 564 m) among seabirds (Kooyman et al. 1992; Kooyman and Kooyman 1995; Wienecke et al. 2007). Emperor Penguins (*A. forsteri*) feed on fish (mainly the Antarctic silverfish *Pleuragramma antarcticum*), squids (*Psychroteuthis glacialis* and *Alluroteuthis antarcticus*), and crustaceans (such as *Euphausiacea superba*), the proportions of which vary with the geographical location (Cherel and Kooyman 1998).

In general, knowledge on endoparasites of Antarctic birds is scarce (Vidal et al. 2012). So far, most reports focused on helminth infections of marine mammals whilst parasitoses of marine birds, particularly from polar regions, have been neglected in the last years (Lauckner 1985). Furthermore, ecological consequences of wide-spread infections in Antarctic bird populations have received few attentions, although a single outbreak may decimate animal populations (Barbosa and Palacios 2009). Since the Antarctic ecosystem is

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not beyond the risk of pathogens and/or parasites due to the geographical isolation and climatic conditions, Barbosa and Palacios (2009) voted for more research in this area, which is claimed necessary to establish general patterns of spatial and temporal variation in pathogens and parasites. Since infections with gastrointestinal parasites mainly depend on foraging habits (Hoberg 1996), modifications in host diet owing to climate change or anthropogenic impacts, such as fishing overexploitation, lead to changes in the occurrence of isolated parasites found in Antarctic penguins (Vidal et al. 2012). Especially, the issue of global warming (Steig et al. 2009), which may lead to increased survival of pathogens and potential outbreaks within such ecosystems (Barbosa and Palacios 2009), is believed to influence and possibly change the occurrence and diversity of prey organisms which in turn leads to changes in parasitoses prevalences. So far, studies on penguin parasites mainly focused on Chinstrap Penguins (*Pygoscelis antarcticus*, Palacios et al. 2012; Vidal et al. 2012), Gentoo Penguins (*Pygoscelis papua*, Fredes et al. 2007; Diaz et al. 2013), and on the closely related King Penguins (*A. patagonicus*, Mawson 1953; Prudhoe 1969; Jones 1988; Fonteneau et al. 2011), for reviews see Barbosa and Palacios (2009) and Clark and Kerry (2000). Some penguin species are known to harbor the anisakid ‘sealworm’ *Contracaecum* spp. (Mawson 1953; Fredes et al. 2007). However, information on gastrointestinal parasites of Emperor Penguins (*A. forsteri*) are scarce and are restricted to antarctic expeditions long time ago reporting on infections with two cestode species, *Parorchites zederi* and *Tetrabothrius wrighti* (Baird 1853; Leiper and Atkinson 1914, 1915; Fuhrmann 1921; Johnston 1937; Prudhoe 1969).

The aim of the current study was to extend the knowledge of intestinal parasite infections of Emperor Penguins (*A. forsteri*). This study represents the first analysis of intestinal parasites of Antarctic Emperor Penguins for more than 40 years and the first molecular characterization of *P. zederi*.

## Material and methods

Fecal samples of 50 individual Emperor Penguins (*A. forsteri*) were collected (Fig. 1) at the Atka Bay, Weddell Sea, East Antarctica (coordinates 70° 36.664' S–70° 37.064' S; 008° 7.709' W–008° 8.769' W) (Fig. 2) in 2012 during the German Antarctica expedition of the Alfred Wegener Institute for Polar and Marine Research (Bremerhaven, Germany), under the compliance with statutory laws of the German Federal Environment Agency. Collected fecal samples were fixed in 70 % ethanol for conservation reasons. At the Institute for Parasitology, Justus Liebig University, Giessen, Germany, samples were analyzed at the coprological diagnostic laboratory unit using the sodium-acetate-formaldehyde (SAF) technique for identification of intestinal helminth eggs and/or

protozoan parasite stages. In addition, coproantigen ELISAs (*Giardia*/*Cryptosporidium*, Thermo Scientific, Oxoid ProSpecT™) were performed to detect *Cryptosporidium* spp. and *Giardia* spp. infections.

Shed cestode proglottids found in fecal material were stained in lactic acid-carmin (Blajin and Rukhadze 1929), differentiated in acid 70 % ethanol, dehydrated, mounted in Eukitt, and thereafter analyzed by light microscopy or alternatively cleared in lactophenol and examined without further treatment.

For the molecular characterization of the shed proglottids, DNA was extracted applying the DNeasy blood and tissue kit (Qiagen, Hilden, Germany) according to the tissue protocol. The near complete 18S ribosomal gene was amplified by PCR with universal forward and reverse primers: NC18SF-1 (5'-AAAGATTAAGCCATGCA-3'; Chilton et al. 2006) and WormB (5'-CTTGTTACGACTTTTACTTCC-3'; Littlewood and Olson 2001). The obtained amplicon was purified by agarose gel electrophoresis, isolated, and directly sequenced with the flanking primers and primer #652 (5'-GCAGCCGCGTAATTCCAGCTC-3'; Nadler et al. 2007) hybridizing to the internal region. The new sequence was submitted to GenBank (accession number KF705621). Sequence analysis for taxonomic classification was performed by BLAST search against the nucleotide collection database (nr/nt). Selected sequences of high score were aligned with MAFFT 7 (Katoh and Standley 2013) using the L-INS-i method and compared using MEGA6 (Tamura et al. 2013).

## Results

### Endoparasites in fecal samples

Parasitological calculations (prevalence in %) were made according to Bush et al. (1997). Thirteen out of 50 examined penguins proved parasitized (26 %). Overall, infections with four species of gastrointestinal parasites were diagnosed (illustrations of parasite stages are depicted in Figs. 3 and 4): *Capillaria* sp. egg (Nematoda) (2.0 % prevalence, 57 µm length/28 µm width), *Tetrabothrius* spp. eggs (Cestoda) (24 % prevalence, egg closed 73–74 µm length/48 µm width; egg open 41.0–41.5 µm length/48 µm width), *Diphyllobothrium* sp. eggs (2.0 % prevalence, 55–59 µm length/37–39 µm width), and cestode proglottids (2.0 % prevalence, eggs 51–52 µm length/37 µm width). It is worth noting that the diagnosis of *Capillaria* eggs in fecal samples represents a new parasite record for Emperor Penguins.

Neither protozoan parasites nor acantocephalan eggs was detected by microscopical examination of the SAF-concentrated fecal samples. Furthermore, all samples proved negative for *Cryptosporidium* spp. and *Giardia* spp. by coproantigen-ELISAs.

**Fig. 1** (1–3) Emperor Penguin (*Aptenodytes forsteri*) colony at Atka Bay, Antarctica (approximately composed of 8,500–9,000 animals)



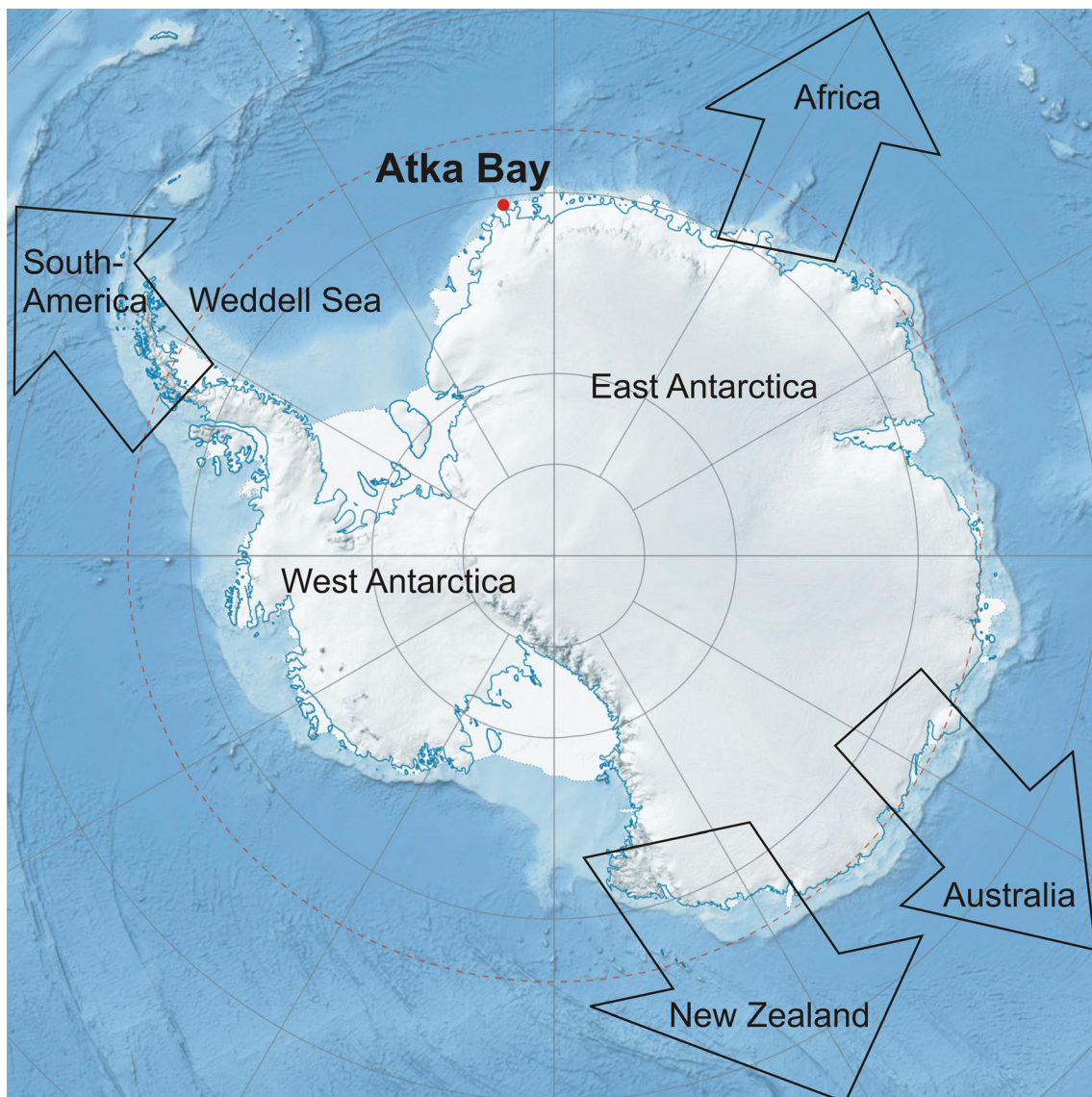
#### Morphological and molecular characterization of *Parorchites zederi*

The fecal sample from one penguin contained a strobila (Fig. 4 (1)) of postmature and gravid proglottids. The proglottids exhibited the following characteristics: much wider than long ( $4,700 \times 500 \mu\text{m}$ ), craspedote, each with single genitalia set. The genital pores lay lateral, alternating irregularly in consecutive segments. Proglottids of the present specimen had large genital atriums which are everted and appeared as prominent, funnel-shaped, muscular bulbs, some of which showing a protruding cirrus (close-up view Fig. 4 (1)). The ovary was situated anterior and next to the genital pore (anteroporal), irregularly lobed, and loose in tissue structure; vitellarium, small and compact, kidney-shaped, posterior to ovary. The uterus in the postmature proglottids (Fig. 4 (1–4)) was reticulate and with parenchymatous egg capsules in mature proglottids. These capsules measured  $150 \times 100 \mu\text{m}$  containing one

egg (monovular) of  $40 \times 50 \mu\text{m}$  with the oncophere surrounded by a thick egg membrane (Fig. 4 (4)). Testes were numerous ( $\sim 50$ – $100$ ) constituting a transverse band, as wide as the proglottide (Fig. 4 (3)). The cirrus sac was very small.

Given these features and the microscopical image, the strobila and proglottids were consistent with *P. zederi* Baird 1853 (Dilepididae) following the key of cestode guide books (Schmidt 1986; Kahlil et al. 1994). This identification was verified by comparison with the more detailed descriptions and drawings of *P. zederi* by Johnston (1937), Cielecka et al. (1992), and Georgiev et al. (1996), of those the last two authors isolated this cestode from *Pygoscelis* spp. penguins in the South Shetlands.

In addition, we performed a first molecular characterization of *P. zederi*. We amplified and sequenced a near complete 1,966 bp fragment of the small subunit ribosomal RNA (rRNA) gene by PCR using universal primers. BLAST search and phylogenetic analyses with this new sequence were in



**Fig. 2** Modified Antarctic map, original work under Creative Commons license: Alexrk2/CC-BY-SA-3.0

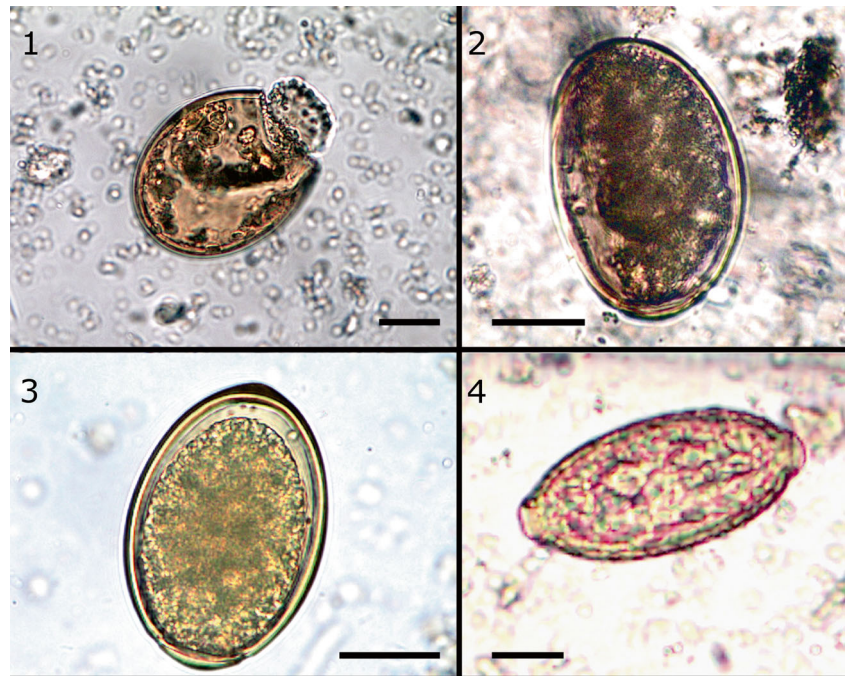
accordance with the morphological classification to the family Dilepididae. So far, *P. zederi* is known as the only member of this order within the pelagic system (Hoberg 2005).

## Discussion

Overall, a low number of parasite species were found in Emperor Penguins (*A. forsteri*), which is in accordance to the previous studies on penguin parasite fauna. In general, the here found prevalences were rather low. Highest prevalences were recorded for the cestode *Tetrabotheirus* spp. of which further species identification based on the morphology of early egg stages was not possible. Representatives of the genus *Tetrabotheirus* had already been recorded in Emperor

Penguins (Prudhoe 1969). The complete life cycle of those cestodes is to date still unclear and further investigations are needed. It is known that plerocercoid larval stages of penguin cestodes use a variety of prey fishes/crustaceans (Williams 1995; Hoberg 2005; Vidal et al. 2012; Diaz et al. 2013) as intermediate hosts and birds as final hosts (Fredes et al. 2007); therefore, it can be expected that the cestode *Tetrabotheirus* also have either prey fishes or crustaceans as intermediate hosts in the Antarctic region. On the other hand, the second cestode species found in this study, *P. zederi*, is a well known parasite of *A. forsteri* collected by several Antarctic expeditions. This cestode is widely distributed among other Antarctic penguins (Hoberg 2005; Barbosa and Palacios 2009; Vidal et al. 2012) due to a broad oceanic distribution of euphausiids, such as the pelagic crustacean *E. superba*, which serves as main intermediate hosts for this cestodes

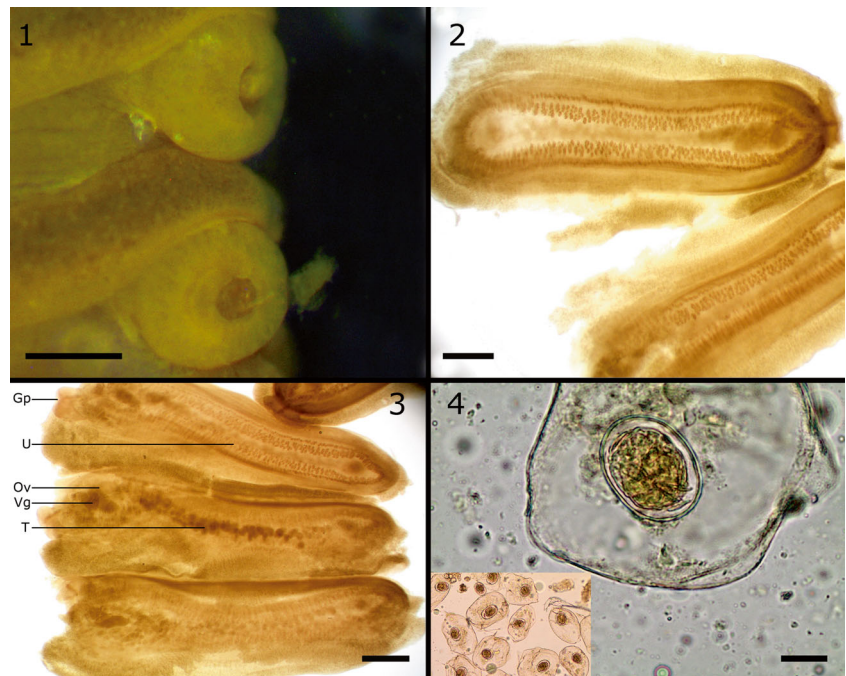
**Fig. 3** Endoparasites from fecal samples of *Aptenodytes forsteri*. 1–2 *Tetrabothrius* spp. eggs. 3 *Diphyllobothrium* sp. egg (Cestoda) (Scale bars 1 and 3; 20  $\mu$ m, 2; 25  $\mu$ m). 4 *Capillaria* spp. egg (Nematoda) (Scale bar 15  $\mu$ m)



(Williams 1995; Hoberg 2005; Vidal et al. 2012; Diaz et al. 2013) and is known as prey organism for Emperor Penguins (Cherel and Kooyman 1998). Despite of an analysis of two large samples of Antarctic krill, Cielecka et al. (1992) did not find any *Parorchites* stages in krill. However, it was isolated from the intestine of crab and leopard seals. Georgiev et al. (1996) supposed that *P. zederi* is a complex of species because Cielecka et al. (1992) reported considerable differences in the size of the rostellar hooks from *Parorchites* isolated from different host species. The SSU rRNA gene sequence of the

specimen from *A. forsteri* might contribute to solve this issue in future experiments. Here, detected *P. zederi* and *Tetrabothrius* eggs in Emperor Penguins (*A. forsteri*) can be distinguished from other parasitic eggs by their characteristic morphology features such as size, egg shell, form, content, and form of operculum, but the contrast is true for other helminth eggs occurring in seabirds such as digenean eggs. To the best of our current knowledge, we here describe the first record of *Capillaria* spp. eggs in Emperor Penguins (*A. forsteri*). The genus *Capillaria* includes a diversity of

**Fig. 4** (1–4) *Parorchites zederi*. 1 Proglottids wider than long. Genital pores irregularly alternating with prominent everted genital atrium, inset close-up view showing protruding cirrus (Scale bar 250  $\mu$ m). 2–3 Detail view of postmature proglottids from *P. zederi* (Scale bars 500  $\mu$ m). Gp genital pore, Ov ovary, T testes, U uterus, and Vg vitelline gland. 4 Isolated egg capsules from gravid proglottids containing one single egg with an oncosphere (Scale bar 20  $\mu$ m)



widespread nematode parasite species, which can have different vertebrates as final hosts, such as fishes, birds, and mammals (Anderson 2002). More than 300 different species of *Capillaria* have been described so far worldwide, many of them occurring in wildlife (Anderson 2002). For some marine *Capillaria* species, the life cycle is fortunately known. The final host shed non-embryonated eggs into the marine environment. These *Capillaria* eggs embryonate thereafter and become infectious. Some species will include fishes as intermediate hosts and these infected fishes can become prey organisms for the penguins, which become infected after consumption and within the final host ingested larva will develop after two molts into adult nematodes and the life cycle is completed. The species *C. convoluta* has previously been recorded from marine seabirds, such as the Southern Giant Petrel (*Macronectes giganteus*) (Mawson 1953) but until now never been described in penguins (for review see Barbosa and Palacios 2009). The low occurrence of *Capillaria* in Emperor Penguins (*A. forsteri*) could be explained by the strict dependency of the life cycle of this nematode species on the aquatic environment and the stenophagic and pelagic diet of Emperor Penguins (see Vidal et al. 2012). However, we can not entirely rule out the possibility of a false-positive finding through intestinal passage or accidental contamination by feces from the Southern Giant Petrel or South Polar Skua.

Furthermore, the low parasite richness observed in our current study, especially for trematodes, could be related to the very low temperatures observed in the Antarctic environment, which might inhibit survival of larval parasitic stages (Lauckner 1985). Despite hostile environmental and climate conditions, it was possible to isolate four different parasite species. Some other related studies on penguin parasites have also shown similar diversities. Vidal et al. (2012) likewise observed four helminth parasite species: two cestodes (*Tetrabothrius pauliani* and *P. zederi*), one nematode (*Stegophorus macronetes*), and one (*Corynosoma* sp.) in Chinstrap Penguins.

Diaz et al. (2013) also revealed four parasite species in Gentoo Penguins: one cestode (*P. zederi*), two nematodes (*Stegophorus macronectes* and *Tetrameres wetzeli*), and one acanthocephalan species (*Corynosoma shackeltoni*). Fredes et al. (2007) detected in total two helminth parasite species, the cestode *Tetrabothrius* sp. and the ascarid nematode *Contraecaecum* spp. Hoberg (1996) stated that the core parasite composition in pelagic birds consists of cestodes, mainly Tetrabothriidae, but with the present study we could clearly expand the spectrum of endoparasites occurring in Emperor Penguins (*A. forsteri*). Within the present study, the nematode *Capillaria* sp. could be identified as one new host record for *A. forsteri*. The isolated cestoda eggs, *Tetrabothrius* spp., were already recorded for Emperor Penguins (Prudhoe 1969) as well as *P. zederi* (Baird 1853). Minor changes/differences

within the parasite composition of Emperor Penguins could be detected within this study. Unfortunately, it was not possible to identify all isolated parasite eggs, at least, on genus level and therefore it is difficult to draw conclusions on the life cycles of all isolated parasites. Changes within the gastrointestinal parasite composition in marine environments are directly linked to the surrounding invertebrate and vertebrate communities. These communities are on their part depended from climate and/or anthropogenic impacts (MacKenzie et al. 1995; Dzikowski et al. 2003; Marcogliese 2003, 2005; Hudson et al. 2006; Hechinger et al. 2007). Therefore, marine parasites with complex multiple-host life cycles (Hechinger et al. 2007) are known to be sensitive bioindicators of aquatic ecosystem health (Overstreet 1997; Dzikowski et al. 2003). They require unaffected environmental conditions to get access to the full range of potential parasite intermediate hosts, whereas monoxenous parasite species (single-host) may persist in highly perturbed, extreme environments (Dzikowski et al. 2003; Hechinger et al. 2007). Under natural conditions, marine hosts can accumulate the highest possible parasite load (endoparasites). Within impacted habitats, this endoparasite load will decrease, whereas the ectoparasite load will increase (MacKenzie et al. 1995; Diamant et al. 1999; Dzikowski et al. 2003; Marcogliese 2003, 2005; Hudson et al. 2006; Hechinger et al. 2007).

The parasitological information presented in this study can be used for conservation purposes of pelagic bird species in the Antarctica. Over time, possible ecological long-term changes could be expected and we therefore call for further investigations on parasitosis occurring in the fragile ecological environment of the Antarctica.

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