



Infection rates and prevalence of metazoan parasites of the non-native round goby (*Neogobius melanostomus*) in the Baltic Sea

Heidi Herlevi · Riikka Puntila  · Harri Kuosa · Hans-Peter Fagerholm

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Abstract Studies in the Baltic Sea have identified over 30 parasite taxa infecting the invasive round goby (*Neogobius melanostomus* (Pallas, 1814)). In this study, we aimed at comparing parasite assemblages and infection rates (prevalence and intensity) in different populations across the invasive range in the Baltic Sea (Denmark, Lithuania, Estonia and Finland). Infection rates were 56–60% across all locations except Lithuania (28%). However, the parasite assemblages in the sampled populations were dissimilar, each location having unique parasites. In addition,

many of the parasites were generalists commonly infecting native fish species. Based on the results of this study and those previously conducted in the Baltic Sea, the round goby has not retained parasites from its area of origin, but instead has been successively colonized by native generalist parasites. Although variable, overall parasite richness is still quite low around the Baltic compared to the native areas (34 vs 71 taxa, respectively). Also, prevalence and mean infection intensities in the Baltic Sea are significantly lower than in the native areas. Therefore, the invasion success of the round goby in the Baltic Sea can at least partly be attributed to enemy release, in this case shedding a significant proportion of their native parasite load.

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H. Herlevi
Environmental and Marine Biology, Faculty of Science and Engineering, Åbo Akademi University, Tykistökatu 6, 20520 Turku, Finland

R. Puntila (✉)
Department of Aquatic Sciences, University of Helsinki, P.O. Box 65, Biocenter 3, Viikinkaari 1, 00014 Helsinki, Finland
e-mail: riikka.puntila@helsinki.fi

R. Puntila · H. Kuosa
Marine Research Center, Finnish Environment Institute, P.O. Box 140, 00251 Helsinki, Finland

H.-P. Fagerholm
Åbo Akademi University, Faculty of Science and Engineering, Environmental and Marine Biology, Laboratory of Aquatic Pathobiology, Tykistökatu 6, 20520 Turku, Finland

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Introduction

The round goby (*Neogobius melanostomus* (Pallas, 1814)) is one of the most widespread non-native fish species in the Baltic Sea (Kotta et al., 2016), with reported detrimental biological impacts in many locations (Ojaveer & Kotta, 2015; Ojaveer et al., 2015). Following its initial detection in the Gulf of Gdansk in 1990 (Skora & Stolarski, 1993), it spread throughout the southern Baltic Sea, eastward to the

Vistula lagoon and river system (Mierzejewska et al., 2011), westward to the German and Danish coastal areas and further north along the southeastern coast of Sweden (Florin & Karlsson, 2011; Sapota, 2011). In the northern parts of the Baltic Sea, round goby became a permanent component of the Lithuanian, Latvian and Estonian coastal fish fauna in the 2000s (Ojaveer, 2006; Sapota, 2011; Rakauskas et al., 2013; Strake et al., 2013). By 2011, the species had been recorded along the Finnish coast of the Gulf of Finland, in the Archipelago Sea and the Åland Islands. The northernmost observations to date were made in 2013 outside the town of Raahe (about 64°4'N) in the Bothnian Bay. Currently, the species has been observed along the coasts of all Baltic sub-basins, except for the Swedish coast of the Bothnian Bay (Kotta et al., 2016). Although originating from the brackish waters of the Black Sea and Sea of Azov, the species has also established viable and flourishing populations in many freshwater areas throughout European river systems (e.g. van Beek, 2006; Ondračková et al., 2010; Borcharding et al., 2011; Emde et al., 2012; Huyse et al., 2015) and the North American Great Lakes (Kornis et al., 2013).

The great invasion success of the species has partly been explained by its pronounced phenotypic plasticity in terms of life-history traits (MacInnis & Corkum, 2000; Gutowsky & Fox, 2012; Brandner et al., 2013a), feeding habits (Brandner et al., 2013b) as well as tolerance of different salinity regimes (Karsiotis et al., 2012; see review by Kornis et al., 2012). Many researchers have also suggested the lack of natural enemies as a major contributing factor (Sapota & Skora, 2005; Kvach & Skóra, 2006; Kvach & Stepien, 2008; Gendron et al., 2012). Parasites are important biological regulators and occur as an inherent part of any biological community (e.g. Williams et al., 1992). They can also be seen as a good indicator of the ecosystem health (Hudson et al., 2006; Palm, 2011). The lack of such natural regulators has—in accordance with the enemy release hypothesis—been pointed out as one of the reasons to the successful colonization and population expansion of many introduced species (Torchin et al., 2003; Colautti et al., 2004). Introduced species are known to have an impact on the native fauna in not only many direct ways, e.g. through competition and predation, but also indirectly by acting as a vector for new parasites and pathogens (Ruiz et al., 1999; Emde et al., 2012). This spillover

effect of non-indigenous parasites being introduced to new ecosystems together with their non-indigenous hosts has been discussed in numerous studies (e.g. Johnsen & Jensen, 1991; Prenter et al., 2004; Peeler et al., 2010; Lymbery et al., 2014). Kelly et al. (2009) highlighted the significance of the spillback of native parasites, i.e. when non-native species act as a new competent intermediate and/or paratenic host for native parasites. This spillback may be of pronounced ecological significance, as it facilitates the transmission of parasites to native hosts and can increase the prevalence of diseases in a population (Kelly et al., 2009 and references therein). However, non-native species may also alleviate the parasitic load of native species by ‘diluting’ the intensity of infection, e.g. when replacing native host species and acting as a “resistant target” (Kopp & Jokela, 2007). Thus, non-native species can simultaneously play many different roles in the parasite–host cycle.

The round goby was most probably transported to the Gulf of Gdansk in ballast water as eggs or larvae (Sapota & Skora, 2005). It is not known whether the source population is from the Black, Azov or Caspian Sea and whether it arrived via the inland river systems of Don/Volga or Dnieper–Vistula, but the species was most likely transported to the Baltic with the ballast water of vessels (Sapota, 2004). This comparatively rapid translocation, together with a steep decline in salinity (16–18 PSU in the Black Sea vs. 7–8 PSU in the Gulf of Gdansk) may have caused the loss of most of the native parasite fauna (Kvach et al., 2014). In a recent study, Kvach et al. (2014) found that the round goby had relatively many (15 species) metazoan parasite species in its native area (the Danube estuary) compared to other gobiid species, but had retained very few when the round goby invaded Vistula delta (5 species). Conversely, the racer goby (*Babka gymnotrachelus*, Kessler, 1857) has lower parasite richness in the native than in the invasive range (Kvach et al., 2014). Many previous results, in addition to the above, show that the round goby may be benefitting from decreased parasite load, supporting the enemy release hypothesis (ERH) (Kvach, 2001; Kvach & Skóra, 2006; Francová et al., 2011; Emde et al., 2012).

Studies have shown that so far the round goby has relatively low infection rates in the Baltic (some 7–20 taxa; Rolbiecki, 2006; Rakauskas et al., 2008) compared to its native range (Kvach 2005) (up to 71 taxa and overall infection rate of up to 97%; Rolbiecki,

2006; Özer, 2007). All parasites found in the Baltic populations have also been previously recorded in the Baltic Sea (Rolbiecki, 2006; Kvach & Winkler, 2011). Therefore, there is no indication of spillover of non-native parasites into the Baltic Sea, which was an obvious concern knowing that some non-indigenous fish have introduced invasive and severely harmful parasites to native species (e.g. Johnsen & Jensen, 1991; Lymbery et al., 2014). However, in the Vistula Lagoon, the round goby has been reported to host larvae of the introduced swim bladder nematode *Anguillicoloides crassus* (Kuwahara, Niimi & Itagaki, 1974) (Kvach, 2004a; Rolbiecki, 2006), which may lead to further spread and transmission to its definitive host, the European eel (*Anguilla anguilla*, L., 1758), whose populations are already declining in the Baltic Sea (Kirk, 2003; Kvach, 2004a; Rolbiecki, 2006). While not currently found parasitizing the round goby in the Baltic Sea, the goby-specific Ponto-Caspian monogenean *Gyrodactylus proterorhini* (Ergens, 1967), has been transferred to the Vistula basin by other Ponto-Caspian gobies (Mierzejewska et al., 2011) and is more abundant in the non-native than in the native areas (Kvach et al., 2014). In other invaded fresh water habitats, like the Rhine River, round goby is also suspected to aid in the spreading of the non-native acanthocephalan parasite *Pomphorhynchus tereticollis* (Rudolphi, 1809) (Emde et al., 2012), and although this particular species is already a part of the parasite fauna in the Baltic Sea (Špakulová et al., 2011), it shows that the round goby has a potential of transmitting parasites to new areas as it continues to spread.

Populations of the round goby across the Baltic Sea are very different. They have a very different invasion history primarily in terms of time since introduction: the most recent introductions and oldest ones are more than 20 years apart (Sapota, 2011; Kotta et al., 2016). Also, the populations are exposed to very different abiotic conditions due to the pronounced gradients in both salinity and temperatures in the Baltic Sea and, therefore, different local parasite assemblages. The population age, i.e. time since introduction, is often reflected in the parasite loads as the infection rates increase the older the population. This has been true, for example, in the Gulf of Gdansk, where the parasite abundance of the round goby increased from six species in the first studies (Rokicki & Rolbiecki, 2002) to at least 12 metazoan parasite species by 2006

(Kvach & Skóra, 2006). In the most recently established populations, the infection rates are presumably still low due to fewer parasitic species adapted to this new host.

Studies focusing on round goby parasites in the Baltic Sea have been quite few, spatially limited to local studies on German, Polish and Lithuanian populations and completely lacking in the northern Baltic Sea (Kvach, 2001, 2004a; Kvach & Skóra, 2006; Rakauskas et al., 2008; Kvach & Winkler, 2011). The aim of the present study was to identify, quantify and compare the most common metazoan parasites (concentrating on metazoan endoparasites) infecting the round goby in four distinct locations in the Baltic Sea. In addition, we also review the existing literature on the round goby parasites in the region, and compare and contrast them to the results of this study.

Material and methods

Round gobies were collected from four locations around the Baltic Sea (Denmark, Lithuania, Estonia and Finland [the Åland islands]) (Fig. 1). The sampling was conducted in June 2015 using identical trap assays at each site (see below). The sites were chosen based on the previous observations of the round goby as these are the locations where the species was first discovered and populations are established and round gobies are abundant (Sapota, 2004; Ojaveer, 2006; Azour et al., 2015; Kotta et al., 2016). The habitat at all sampling locations consisted of vegetated sandy bottoms, except in Finland and Estonia where the bottom substrate consisted mainly of coarse gravel and rocks interspersed with sandy patches. This reflects a general change in dominant habitats from southern to northern Baltic Sea. The locations in Lithuania, Estonia and Finland were also close to artificial structures such as piers or built embankments, which has been noted as the round gobies preferred habitat as they provide suitable nesting sites and shelter (MacInnis & Corkum, 2000; Sapota & Skora, 2005; Ojaveer, 2006). The populations differ as to the time of invasion: the Lithuanian population around Klaipeda was discovered in 2002 (Sapota, 2011) and the population in Muuga, Estonia (Gulf of Finland) was discovered two years later in 2004 (Ojaveer, 2006). The first records of the round goby from Guldborgsund

Fig. 1 Sampling sites around the Baltic Sea (see Table 1 for more information) and the prevalence of parasite infection at each location

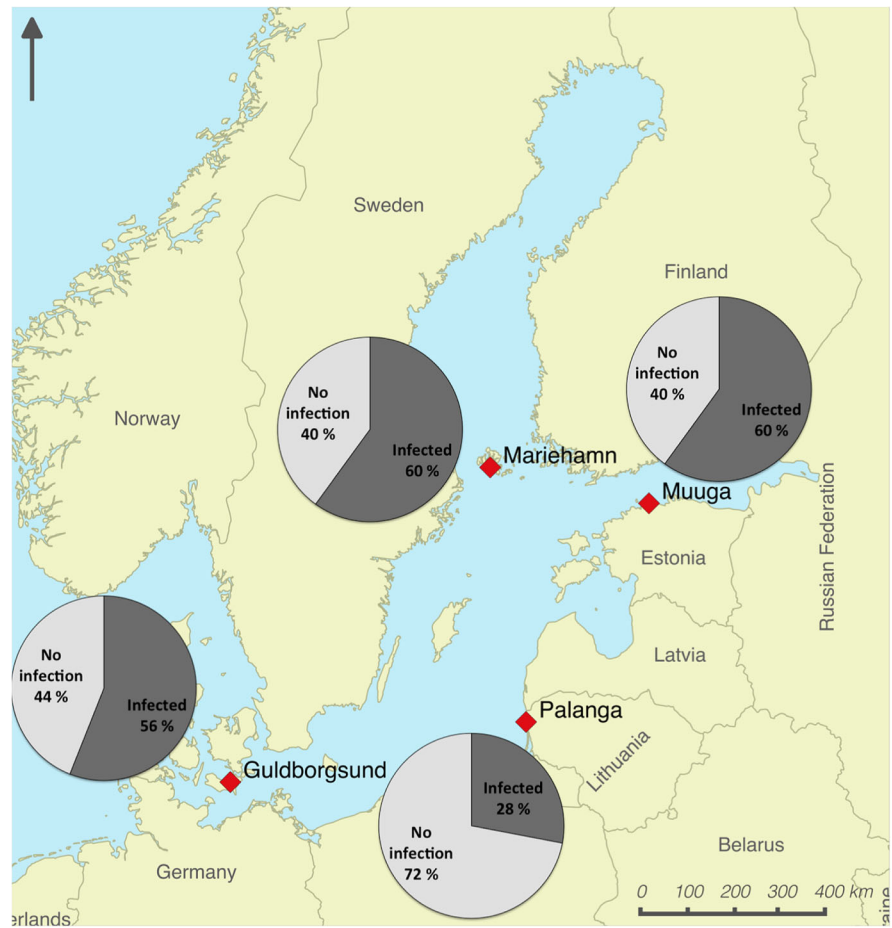


Table 1 Hydrographic data during sample collection at each site

Site	Sampling date	Temperature (°C)	Salinity (PSU)
1. Guldborgsund (DK)	25.05.2015	16.7	11.9
2. Palanga (LT)	08.06.2015	15.6	5.8
3. Muuga (EE)	16.06. and 09–17.07.2015	15.0–17.0	6.0
4. Mariehamn (FI)	26.06.2015	15.1	5.8

(GBS), Denmark, are from 2008 (Azour et al., 2015) and the population in Mariehamn, Åland islands, was first recorded in 2011 (Kotta et al., 2016).

Hydrographic information from each site at the time of sample collection is presented in Table 1. Fish were collected using three types of passive gear; 5 minnow traps (mesh size 6 mm), 10 collapsible crayfish traps (mesh size 12 mm) and 3 eel traps (mesh size in cod ends 10, 14 and 17 mm). This combination was used to provide a better size range of

catch than any trap type used alone. The traps were placed at approximately 2-m intervals parallel to the shoreline at 1–2 m depth and checked every 24 h during 3 days. A piece of frozen herring (or other locally available fish) was placed in sealed mesh bags inside the minnow and crayfish traps and baits were replaced as needed. The use of both baited and unbaited methods also gives a more representative sample of the population. The sampling procedure remained the same throughout the locations to enable

comparisons between locations. Fish from each sampling time ($n = 3$) and trap type ($n = 3$) were placed in separate bags and 3–4 fish were haphazardly taken from each bag for inspection so that a total number of 25 of round gobies per site, representing various sizes, were randomly selected for closer inspection. The collected fish were immediately terminated and frozen (-18 to -20°C) until examined. Before dissection, they were measured (total length, TL, in mm), weighed (W, in g) and their sex was determined. Their livers were extracted and weighed to obtain an estimate of health and energy reserves expressed as hepatosomatic index (HSI) ($\text{HSI} (\%) = 100 \times (\text{liver weight [g]} / \text{whole fish weight [g]})$). In addition, the condition of the inspected fish was expressed using the Fulton's K index (Fulton, 1904) as $K = 100 \times W [\text{g}] / \text{TL} [\text{cm}]^3$. Differences in fish condition between sites were tested using analysis of variances (ANOVA) and post hoc comparisons using the Tukey HSD procedure. Hepatosomatic index was log-transformed to fulfil the assumptions for parametric analyses.

The skins, fins and gills were carefully examined visually in case of signs of ectoparasites (or larval stages of endoparasites like metacercariae of digenean trematodes) and all anomalies in appearance were noted. All ectoparasite taxa are typed in italics (Tables 4 and 6) to allow for comparisons with the previously published studies. Opaque eyes can be a sign of *Diplostomum* spp. metacercariae larvae, which infect the lens of the eye and have been found in previous studies as one of the most abundant parasites of the round goby in the Gulf of Gdansk (Kvach & Skóra, 2006; Rolbiecki, 2006). The eyes of each fish were dissected and inspected carefully under a dissection microscope.

The entire intestinal tract was then removed for inspection of endoparasites on both inner and outer intestinal surfaces. The gut contents were removed and the digestive tract inspected carefully for parasites. Food items in the stomach and gut were also recorded by examining the contents under a microscope and identifying which taxa were present. The number of taxa in each stomach and the frequency of occurrence ($\text{FO} \% = \text{stomachs including prey item "i"} / \text{total number "n"} \text{ of stomachs with contents at location "x"}; \text{Hyslop, 1980}$) were calculated. This was done to get a picture of which taxa are being consumed at each location at the time of sampling. The body cavity and organs (gonads, kidney, liver, mesenteries and spleen)

were inspected under a dissection microscope. All parasites were counted and stored in 70% alcohol prior to a more detailed examination and identification. Digenean trematode larvae and Hirudinea were identified according to descriptions in Valtonen (2012). Often the Diplostomid larvae found in the lens of fish have been identified as *Diplostomum spathaceum* (Höglund & Thulin, 1982), but as studies have shown there are at least two species infecting the lens (Valtonen & Gibson, 1997), and *D. spathaceum* is most likely a conglomeration of species. Thus, all Diplostomid larvae in this study were assigned to *Diplostomum* spp. since no genetic verification was made. Nematodes and Acanthocephalans were cleared in lacto phenol before examination under a light microscope. Identification was done according to morphological descriptions of, e.g. foregut, head and tail structures of nematodes in Fagerholm (1982) and Moravec (1994). Acanthocephalans were identified according to a key (Arai, 1989) and descriptions by Valtonen (2012).

The parasitological indices were calculated according to Rózsa et al. (2000) as: prevalence (proportion of fish infected of all fish examined/site), mean intensity (MI) and median intensity (mean and median number of parasites in infected fish, MedI) and their ranges given as confidence limits obtained by bootstrapping (Rózsa et al., 2000) using the Quantitative Parasitology 3.0 software (Reiczigel & Rózsa, 2005). The differences in parasite prevalence between the sampling locations and sexes were analysed using the Fisher's exact test and the mean intensities between the locations and sexes analysed using the non-parametric Wilcoxon's test due to the skewed nature of the infection intensity data. Multiple comparisons (post hoc tests) were conducted using the non-parametric Steel–Dwass method (Critchlow & Fligner, 1991). In addition, the differences in parasite assemblages between the locations were analysed using analysis of similarities (ANOSIM) and taxa contributing the most to the observed differences were determined by similarity percentage analysis (SIMPER).

All statistical analyses were conducted using the JMP Pro 11 software (SAS Institute Inc., 2013), except the comparisons of the parasite assemblages in the infected fish between sampling locations, which were made using ANOSIM (analysis of similarities) and SIMPER (similarity percentages analysis) in Primer v6 software (Clarke & Gorley, 2006).

Results

Overall condition of the round gobies in different locations

Out of a total of 100 fish that were examined, 32 were females and 68 males (Table 2). Both males and females were the smallest in Guldborgsund (TL 64–133 mm), whereas males were the largest in Mariehamn (mean TL 152,5 ± 23,5 mm) and females in Palanga (TL 77–198 mm) (Table 2). No females

were caught in Mariehamn. The hepatosomatic Index (HSI) varied between the locations ($F(3,107) = 9.49$, $P = 0.0001$). Fish from both Muuga ($P = 0.0002$) and Mariehamn ($P = <0.0001$) had a highly significantly higher HSI than in Guldborgsund and significantly higher in Palanga ($P = 0.035$). The condition index (Fulton's K) was similar throughout the sampling locations ($F(3,96) = 0.587$, $P = 0.625$) and the infection intensity had no impact on either HSI or Fulton's K ($F(1,50) = 2.42$, $P = 0.126$ and $F(1,50) = 2.67$, $P = 0.109$, respectively).

Table 2 Number of fish, the total length (TL) and weight of round gobies inspected per site

Site	GBS		Palanga		Muuga		Mariehamn	
	Female	Male	Female	Male	Female	Male	Female	Male
N	12	13	9	16	11	14	0	25
TL (mm)	$x \pm SD$	Min–max	$x \pm SD$	Min–max	$x \pm SD$	Min–max	$x \pm SD$	Min–max
Female	96.9 ± 15.4	76–133	124.2 ± 26.9	98–180	147.2 ± 27.4	90–168	–	–
Male	95.6 ± 17.6	64–126	144.4 ± 41.6	77–198	128.1 ± 32.9	72–170	152.5 ± 27.5	93–205
Total	96.2 ± 16.3	64–133	137.2 ± 37.7	77–198	136.5 ± 31.5	72–170	152.5 ± 27.5	94–205
Weight (g)	$x \pm SD$	Min–max	$x \pm SD$	Min–max	$x \pm SD$	Min–max	$x \pm SD$	Min–max
Female	13.8 ± 7.8	5.5–34.0	30.1 ± 23.3	12–80.1	55.1 ± 23.7	8.3–76.4	–	–
Male	14.9 ± 9.8	3.1–37.9	55.2 ± 43.2	6.1–129.1	34.9 ± 24.4	4.8–70.5	55.2 ± 25.9	18.7–121.4
Total	14.4 ± 8.7	3.1–37.9	46.1 ± 38.7	6.1–129.1	43.8 ± 25.7	4.8–76.4	55.2 ± 25.1	18.7–121.5

Length and weight are presented as mean and standard deviation together with the range for each parameter. Each parameter is given separately for both sexes and as a total per location

Table 3 Number of infected and non-infected fish, prevalence (%) of infection, and mean and median intensities of infections (MI and MedI, respectively) at each location

Site	Sex	N	Infected	Prev.	Lower CL	Upper CL	MI	Lower CL	Upper CL	MedI	Lower CL	Upper CL
Guldborgsund		25	14	0.56	0.35	0.76	9.14	4.71	15.30	5	2	5
	M	13	6	0.46	0.19	0.75	7.50	2.17	21.80	3.5	1	31
	F	12	8	0.67	0.35	0.90	10.40	4.38	19.90	5	2	26
Palanga		25	7	0.28	0.15	0.54	2.12	1.00	4.75	1	1	3
	M	16	4	0.25	0.07	0.52	1.50	1.00	2.00	1	–	–
	F	9	3	0.33	0.14	0.79	2.75	1.00	4.50	1	–	–
Muuga		25	15	0.60	0.39	0.79	6.60	3.20	7.67	3	2	9
	M	14	7	0.50	0.23	0.77	5.29	1.86	10.10	2	1	16
	F	11	8	0.73	0.39	0.94	5.00	2.88	7.62	4	1	9
Mariehamn		25	15	0.60	0.39	0.79	7.93	4.00	16.50	3	1	7
	M	25	15	0.60	0.39	0.79	7.93	4.00	16.50	3	1	7
	F	0	–	–	–	–	–	–	–	–	–	–

The range is expressed as confidence limits (CL)

Parasite prevalence, infection intensity and assemblages in different locations

In total 51% of the fish examined were infected by at least one parasite taxa (Table 3). Overall, 383 specimens of metazoan parasites representing 10 taxa (3 species of Trematoda (Digenea), 3 Nematoda, 3 Acanthocephala and 1 Hirudinea) were identified (Table 4).

In Muuga, 73% of all females were infected, whereas the total prevalence was 60% in both Muuga and Mariehamn (Fig. 1). These two sites had the highest prevalence, compared to 56% in GBS and 28% in Palanga (Fig. 1, Table 3), although the differences were non-significant (Fisher’s exact test,

2-sided, $P = 0.164$). Similarly, there was no significant difference in prevalence between the sexes (Fisher’s exact test, 2-sided, $P = 0.199$). The observed numbers of parasite taxa were similar throughout the locations: three in GBS and Muuga, four in Palanga and five in Mariehamn (Table 4). The populations in Muuga and Mariehamn had the highest infection intensity (number of parasites in one fish) with a maximum of 99 and 121 parasite individuals observed. Overall infection intensity was very close to significantly higher in females than males ($Z = 1.93$, $P = 0.052$). The infection intensity (mean intensity) varied between the locations ($X^2(3, N = 52) = 8.29$, $P = 0.040$; Table 5). These differences are mainly driven by Palanga, which had significantly lower

Table 4 Parasites of *N. melanostomus* at four locations in the Baltic Sea

Parasite species, stage	Location	Guldborgsund			Palanga			Muuga			Mariehamn		
		P	MI	I	P	MI	I	P	MI	I	P	MI	I
Trematoda (Digenea)													
<i>Cryptocotyle</i> sp., met. ^{ba} (Creplin, 1825)	Skin, fins	40	9	5–30									
<i>Diplostomum</i> spp., met. (Rudolphi, 1819)	Eye, lens							52	6	1–16	40	10	1–36
<i>Tylodelphys clavata</i> , met. ^b (Nordmann, 1831)	Vitreous body							12	4	1–7			
Nematoda													
<i>Camallanus</i> sp., ad.	Intestine										8	2	1–2
<i>Contraecaecum</i> spp., L3	On intestine	12	2	1–4									
<i>Hysterothylacium aduncum</i> L3, L4, ad. (Rudolphi, 1802)	Intestine, mesentery				16	3	1–7						
Ascaridoidea indet., encysted larvae	Intestinal wall, mesenteries, body cavity	16	1	1	4	1	1	12	1	1	12	1	1
Acanthocephala													
<i>Corynosoma</i> sp., cysth.	Intestine										4	1	1
<i>Echinorhynchus gadi</i> , ad. (Müller, 1776)	Intestine				4	1	1						
<i>Pomphorhynchus laevis</i> , cysth. (Müller, 1776)	Intestinal mesentery				8	1	1						
Hirudinea													
<i>Piscicola geometra</i> ^c (L., 1761)	Skin										4	1	1–2
	Total	56	9	1–31	28	2	1–8	60	7	1–16	60	8	1–36

The primary site of infection, prevalence in percent (P), mean intensity (MI) and intensity (min–max) is given for each species/taxa, as well as, a total at each site

Met metacercariae larvae; *L3*, *L4* stage 3 and 4 larvae; *cysth* cystacanth larvae; *ad* adult

^a Metacercariae capsules (black pigmentation) counted

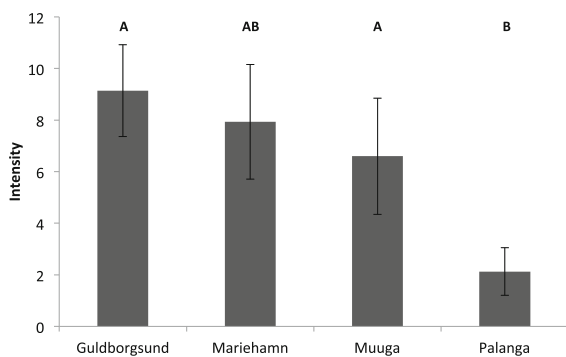
^b Identification based on ecological characteristics

^c Found unattached to fish, in the bag with sampled fish

Table 5 Results of non-parametric analyses of variances (Wilcoxon's test) and multiple comparisons (Steel–Dwass method) of the mean intensities of parasite infection between the different locations

Wilcoxon test	χ^2	DF	<i>P</i> value	<i>P</i> value
Overall	8.29	3	0.0404	
Multiple comparisons	Mean difference	SE dif	Z	
Muuga: Mariehamn	2.6	3.19	0.82	0.8474
Muuga: Guldborgsund	−1.38	3.13	−0.44	0.9713
Mariehamn: Guldborgsund	−2.21	3.12	−0.71	0.8938
Palanga: Mariehamn	−4.89	2.80	−1.75	0.2999
Palanga: Guldborgsund	−7.17	2.78	−2.58	0.0485
Palanga: Muuga	−7.95	2.92	−2.73	0.0324

Significant differences are indicated in bold

**Fig. 2** Differences in mean intensity (MI) of parasite infection between the sampling locations. Variability expressed as standard error. Same letters shared among the groups indicate no significant difference according to the Steel–Dwass method (post hoc test)

mean intensities than Guldborgsund and Muuga (Fig. 2; Table 5).

The non-parametric analysis of similarities shows that parasite assemblages were significantly dissimilar (Global $R = 0.442$, $P = 0.001$). Dissimilarity was largest between Muuga and Palanga ($R = 0.753$, $P = 0.001$; Table 6) and smallest (NS) between

Muuga and Mariehamn (Table 6). Analysis of Similarity Percentages (SIMPER) revealed that the differences were driven largely by the differences in abundance of *Diplostomum* spp. and their absence in Palanga, as well as, the presence of *H. aduncum* in Palanga (Table 4). Guldborgsund had the lowest similarity with all other sites, which was primarily explained by the presence of *Cryptocotyle* sp. in Guldborgsund and its absence elsewhere (Table 4). All sampled populations hosted at least one unique parasite taxa, which was not found in any of the other populations.

Parasite taxa found in the different locations

Guldborgsund, GBS (Denmark)

In Guldborgsund, 14 of the 25 fish inspected (56%) were infected by parasites (Table 3). Mean intensity was 9 and maximum intensity was 31 individuals (Table 4). The most common parasite in the round goby population, infecting 40% of the fish inspected, was the metacercariae larvae of *Cryptocotyle* sp.

Table 6 ANOSIM table of Global R values and their significance indicating differences in parasite assemblages between the sites

	Guldborgsund	Palanga	Muuga	Mariehamn
Guldborgsund		0.549***	0.747***	0.405***
Palanga	<i>0.549***</i>		0.753***	0.306***
Muuga	<i>0.747***</i>	<i>0.753***</i>		NS
Mariehamn	<i>0.405***</i>	<i>0.306***</i>	NS	

NS not significant

Italic values are indicated ($P > 0.05$)

*** $P < 0.0001$

infecting the skin and fins of the fish with an intensity of up to 30 capsules per fish (MI = 9) (Table 4). The infection causes a change in pigmentation of the host's skin around the cysts, which are seen as distinct black spots (black spot disease) on the skin. This is a typical infection of *Cryptocotyle concavum/lingua* (Creplin, 1825), which has been observed infecting the round goby, as well as, other fish in the southwestern Baltic Sea (Køie, 1999; Rokicki & Rolbiecki, 2002; Unger et al., 2014). All other parasites were larval stages of nematodes and occurred on the intestinal mesenteries or the body cavity at low intensities (1–4 individuals). Nematodes of the genus *Contracaecum* infected 12% of the round gobies in Guldborgsund. Based primarily on the size and site of infection, at least four of these were considered to be *Contracaecum rudolphii* A or *Contracaecum rudolphii* C ($P = 4\%$, $I = 4$). Four encapsulated larvae of an unidentified ascaridoidean nematode species were also found in the mesenteries and body cavity of four round gobies ($P = 16\%$) (Table 4). Most infected fish only hosted one species of parasite but three individuals were infected by both *Cryptocotyle* metacercariae and nematode larvae.

Palanga (Lithuania)

In Palanga, 7 of 25 fish inspected (28%) were infected by parasites. The species richness was four species with very low intensity, ranging between 1 and 8 individuals (MI = 2; Tables 3, 4). The most common parasite was the nematode *Hysterothylacium aduncum* ($P = 16\%$), which had a maximum intensity of 7 individuals in one host (MI = 3). Both larvae and adults were identified. One unidentified ascaridoidean larvae was found in the mesentery ($P = 4\%$). In addition to nematodes, two species of acanthocephalans were identified. Cystacanth stages of *Pomphorhynchus laevis* (Müller, 1776) were found in the intestinal mesenteries of two fish ($P = 8\%$). One adult *Echinorhynchus gadi* (Müller, 1776) was found in the intestine of a round goby ($P = 4\%$) (Table 4). One individual hosted both *H. aduncum* and *P. laevis*; all others were only infected by one species.

Muuga (Estonia)

In Muuga, 15 of 25 fish inspected (60%) were infected by parasites (MI = 7; Table 3). *Diplostomum* spp. was the most prevalent parasite species infecting 52%

of the round gobies (Table 4). Another digenean trematode species, *Tylodelphys clavata* (Nordmann, 1831), was found in 12% of the fish. The metacercariae larvae of *Diplostomum* spp. occurred in the eye lenses of the fish, whereas *T. clavata* infects the vitreous body. The mean and maximum intensities for these two parasites were 6 and 16, and 4 and 7 individuals, respectively (Table 4). In addition, a few encapsulated ascaridoidean larvae were found embedded in the intestinal wall and mesenteries of the fish ($P = 12\%$). In Muuga, 16% of the fish hosted either both species of digenean trematode larvae or digenean trematode larvae and nematode larvae simultaneously, at intensities varying between 2 and 15. The total mean intensity was 7 and the maximum observed was 16 individuals per one host (Tables 3, 4).

Mariehamn (Finland)

In Mariehamn, 15 of 25 fish inspected (60%) were infected by parasites. The most common parasite, *Diplostomum* spp. ($P = 40\%$), was the same as in Muuga (Table 4). In contrast, the parasite richness (5 species), together with the maximum intensity (36 individuals) were the highest among all sites (Table 4). Three of the parasite taxa were unique and one of these (*Corynosoma* sp.; Acanthocephala) has not been found in the round goby before. Only one immature individual of *Corynosoma* sp. ($P = 4\%$) was found in the intestine of a round goby. Another of the species not found elsewhere was the nematode *Camallanus* sp., which was found in 8% (MI = 2) of the round gobies and the third species was the hirudinean *Piscicola geometra* L., 1761. *Piscicola geometra* was also the only ectoparasite encountered. Although the two individuals found were no longer attached to the fish hosts when thawed, it is clear they had been attached at the time of capture. Three round gobies ($P = 12\%$, MI = 1; Table 4) also hosted encapsulated ascaridoidean larvae in their intestinal wall. The majority of round gobies hosted only one parasite species, but 8% were parasitized by both *D. spathaceum* and *Camallanus* sp.

Discussion

Currently, 34 metazoan parasite species have been found infecting the round goby within the invaded

Table 7 Infection rate by parasites of *N. melanostomus* around the Baltic Sea basin based on the existing literature and the present study

	Prior literature				Present study			
	Gulf of Gdansk ^{a,b,c}	Vistula lagoon/delta ^{d,g,h}	SW Baltic ^k (German coast) ^f	Curonian lagoon/Klaipeda strait ^e	Guldborgsund	Palanga	Muuga	Mariehamn
Monogenea								
<i>Gyrodactylus rugiensis</i> , ad.			X					
Cestoda								
<i>Bothriocephalus</i> sp.(<i>scorpii</i>), pl.	X	X	X					
<i>Eubothrium crassum</i> , pl.		X						
<i>Paradilepis scolecina</i> , pl.		X						
<i>Proteocephalus filicollis</i>		X						
<i>Proteocephalus gobiorum</i>		X						
<i>Proteocephalus</i> sp.		X		X				
Trematoda (Digenea)								
<i>Bucephalus polymorphus</i> , met.			X					
<i>Bunodera luciopercae</i>		X						
<i>Cryptocotyle</i> spp, met.	X	X			X ⁱ			
<i>Diplostomum</i> spp., met.	X	X	X	X			X	X
<i>Tylodelphys clavata</i> met.	X	X					X	
<i>Tylodelphys</i> sp., met.			X					
Nematoda								
<i>Anguillicoloides crassus</i> , L3	X	X						
<i>Camallanus truncatus</i>		X						
<i>Camallanus lacustris</i>				X				X
<i>Contraecum</i> spp., L3		X			X			
<i>Contraecum rudolphii</i> L3		X						
<i>Cosmocephalus obvelatus</i> L3		X	X					
<i>Cystidicoloides ephemeridarum</i>		X						
<i>Dichelyne minutus</i>	X	X						

Table 7 continued

	Prior literature				Present study			
	Gulf of Gdansk ^{a,b,c}	Vistula lagoon/delta ^{d,g,h}	SW Baltic ^k (German coast) ^f	Curonian lagoon/Klaipeda strait ^e	Guldborgsund	Palanga	Muuga	Mariehamn
<i>Eustrongylides excisus</i> L3			X					
<i>Hysterothylacium aduncum</i>, L3, L4, ad.	X	X		X		X		
<i>Paracuaria adunca</i> , L3			X					
<i>Agamonema</i> sp. L3		X						
Ascaridoidea indet.					X	X	X	X
Acanthocephala								
<i>Acanthocephalus anguillae</i>				X				
<i>Acanthocephalus lucii</i>	X							
<i>Echinorhynchus gadi</i>	X	X	X	X		X		
<i>Corynosoma</i> sp.								X
<i>Pomphorhynchus laevis</i>	X	X	X	X		X		
Hirudinea								
<i>Piscicola geometra</i>	X							X
Crustacea								
<i>Ergasilus sieboldi</i>	X							
Bivalvia								
<i>Unio</i> sp., glochidia		X	X					
Total	13	22	11	7	3	4	3	5
Infection rate	98.7% ³	18.3 ^d	58.10% ^{if}	–	56%	28%	60%	60%

The most common parasites are in bold and ectoparasites in grey

Met metacercariae larvae; L3, L4 stage 3 and 4 larvae; *cysth* cystacanth larvae; *pl* plerocercoid larvae; *ad* adult

^a Rokicki & Rolbiecki, 2002, ^bKvach, 2001, ^cKvach & Skóra, 2006, ^dRolbiecki, 2006 (and references therein), ^eRakauskas et al., 2008, ^fKvach & Winkler 2011, ^gKvach et al., 2014, ^hSzostakowska & Fagerholm, 2007

ⁱ Identification based on ecological characteristics

^j Value given in Kvach & Winkler, (2011) including microsporidians

^k Area of study includes: Szczecin Lagoon, Peenemünde (Peene river), Strelasund strait, Unterwarnow (Mecklenburg bight) and Kiel Canal

Baltic Sea basins and lagoons (Kvach, 2001; Rokicki & Rolbiecki, 2002; Kvach & Skóra, 2006; Rolbiecki, 2006; Rakauskas et al., 2008; Kvach & Winkler, 2011; Table 7). Most of the taxa have been detected in the Gulf of Gdansk, and Vistula lagoon and river delta (13

and 22 taxa respectively; Table 7). This is expected, as these areas have been inhabited by round gobies the longest, since the early 1990s. Previously, the lowest species richness was reported for the Curonian lagoon, where only 7 parasitic species were found (Rakauskas

et al., 2008). In the Gulf of Gdansk, round goby populations have shown a clear increase with time in the number of parasites and intensity of infection. The first study in the region only showed five taxa of metazoan endoparasites infecting the round goby (Rokicki & Rolbiecki, 2002), whereas, a few years later, 12 taxa (8 endoparasite species and 4 ectoparasites) were recorded (Kvach & Skóra, 2006). The later study also had an infection rate of 98.7% (Kvach & Skóra, 2006), which closely resembles the infection rate of 97.5% reported by Özer (2007) in the goby's native range in the southern Black Sea. However, the mean intensity of parasite infections was much higher (78.2 ± 23.1) in the Black Sea than in the Gulf of Gdansk. In the invasive range, the round goby also tends to have a lower prevalence of parasites compared to other gobiid species and other fish species (Kvach, 2001, 2004b; Ondračková et al., 2010; Gendron et al., 2012). In the southwestern Baltic region, both species richness (11 species; Table 7; Kvach & Winkler, 2011) and infection rates (57%; Table 7; Kvach & Winkler, 2011) are lower than in other gobiid species, which all host a range of 14–24 parasite species and have infection rates regularly attaining 100% (Zander, 2003).

The parasite taxa richness in the present study is quite low (3–5 species; Tables 4, 7). Nevertheless, many of the most common species found in the present study are the same as in the previous investigations. Digenean trematode larvae of the species *Diplostomum* spp. have been observed as the most prevalent parasites in many previous studies in the Baltic Sea (Kvach & Skóra, 2006; Kvach & Winkler, 2011; Kvach et al., 2014; Table 7), as well as, other native and non-native areas (Kvach & Skóra, 2006; Kvach & Stepień, 2008; Francová et al., 2011; Gendron et al., 2012). In this study, it was found to be the most common parasite with the highest prevalence and intensity in Muuga and Mariehamn (Table 3), but absent from the two other sites. *Diplostomum* spp. together with another digenean trematode species, *Cryptocotyle* sp. observed in Guldborgsund, were the only species reaching high prevalence and intensities (Table 4).

Of the nematode species observed in this study, *Hysterothylacium aduncum* has frequently been reported infecting the round goby in all areas except the southwestern Baltic Sea (Kvach & Winkler, 2011; Table 7). This is despite the fact that *H. aduncum* does

occur quite commonly in the area and has been recorded in native goby species (Zander, 2003). The acanthocephalans *Echinorhynchus gadi* and *Pomphorhynchus laevis*, which were found in Palanga (Table 4), are also common species reported for the round goby in the Baltic Sea, as well as, other brackish and freshwater native and non-native areas (Kvach & Skóra, 2006; Francová et al., 2011; Table 7). *Hysterothylacium aduncum* and *E. gadi* are both fish parasites of marine origin which are commonly found in eelpout (*Zoarces viviparus* L., 1758) and cod (*Gadus morhua* L., 1758), but can also be found in other fish species (Fagerholm, 1982; Valtonen, 2012).

Larvae of *Contracaecum osculatum*, a mammalian parasite infecting mostly seals, have been observed in the round goby in the Vistula lagoon (Rolbiecki, 2006), but the distinction between *C. osculatum* and *C. rudolphii* larvae is often difficult and thus some specimens may have been misidentified. However, *C. rudolphii* found in this study in Danish and Lithuanian round gobies is an avian parasite, maturing primarily in cormorants (Szostakowska & Fagerholm, 2007, 2012). Cormorants have been known to prey actively on round gobies where they co-occur, e.g. in the Gulf of Gdansk (Bzoma, 1998) and the Curonian Lagoon (Rakauskas et al., 2013). Thus, round goby is a potential paratenic host in completing the life cycle of this *C. rudolphii* in the Danish and Lithuanian populations (Table 4).

This study gives a new host record for the acanthocephalan *Corynosoma* sp. parasitizing round gobies in the Baltic Sea. The taxon was observed in Mariehamn, Finland (Table 4). The three species in the genus *Corynosoma* occurring in the Baltic Sea are all primarily seal parasites and mainly use the amphipod *Monoporeia affinis* (Lindström, 1855) and sculpins or other fish as the intermediate and paratenic hosts (Valtonen, 2012). So far, there has been no evidence of seals using round gobies as prey. However, the migration patterns of the round goby in the Baltic Sea are still poorly understood and it is possible that during autumn and early winter as they migrate to deeper water and possibly further offshore, seals could prey on round gobies. Thus, the round goby may act as an alternative host in the life cycle of *Corynosoma* sp. It is also possible that the round goby is merely a “dead-end” paratenic host for these species and could thus contribute to a diluting effect for these parasites if they do not reach their definitive hosts. However, with

such low infection intensities in the round goby, the aforementioned ecological effects are likely to be restricted.

The observed parasite species are all generalists and occur commonly in the Baltic Sea. The fact that the round goby shows lower infection rates than other benthic fish species in the invaded area (Kvach & Winkler, 2011), together with the lack of any specialized and/or non-native parasites (Kvach, 2001; Kvach & Skóra, 2006), complies with the theory of enemy release in the Baltic populations (Torchin et al., 2003; Colautti et al., 2004). We suggest that a similar increase in parasite infection prevalence and intensity, as was seen in the Gulf of Gdansk, may be expected in the northern populations, sampled in this study, as the native parasites gradually adapt to and colonize this new host species.

The Baltic Sea parasite fauna, similarly to the Baltic Sea fauna and flora in general, is a heterogeneous assemblage of species of freshwater, marine and brackish water origin (Fagerholm, 1982; Kjøie, 1999; Zander & Reimer, 2002). Some changes in the parasitic fauna of the round goby can thus be observed due to differing salinity in the southern/southwestern population (Denmark) compared to the northern populations (Estonia and Finland). The increase of *Diplostomum* spp. and decrease of *Cryptocotyle* sp. has often been observed in studies following a southwest to northeast salinity gradient in the Baltic Sea (Kjøie, 1999; Unger et al., 2014). The low similarity between the locations (Tables 3, 4 and 7) might thus be explained by the difference in salinity, which causes significant differences in benthic fauna composition and, consequently, in parasite fauna that rely on these species as intermediate and paratenic hosts (Williams et al., 1992; Zander & Reimer, 2002). The parasite composition is probably a reflection of the salinity tolerance of both the parasite species and its intermediate host species. It would seem that in general the infection rate is higher in low and intermediate salinities, like in the Vistula and Szczecin lagoons (Rolbiecki, 2006; Kvach & Winkler, 2011; Table 7), as well as, in Mariehamn and Muuga (Table 3). The round goby seems to host a mix of parasites of both limnic and marine origin throughout the Baltic Sea, albeit clearly dominated by a few limnic species.

Due to complex life histories of many parasites, completing their life cycle depends on access to

suitable hosts in the system. In this study, the lack of gastropods in the diet of the round goby in Palanga (FO = 5.6%; Table 8), indicates an absence of gastropod species in the habitat and thus appeared to be reflected in an absence of some parasite species, especially digenean trematodes of the genera *Diplostomum*, *Tylodelphys* and *Cryptocotyle*, which use gastropods as intermediate hosts for the metacercariae larvae (Zander, 2003; Valtonen & Gibson, 1997). The lower infection rate in Lithuania (28%) is probably a result of this absence of digenean trematode larvae, which account for the highest prevalence at all other sites (Table 4). The absence of *Diplostomum* spp. metacercariae in Palanga is noteworthy, since it was the most prevalent parasite (13.5%) in 2007 (Rakauskas et al., 2008) and has also been reported in other fish species along the Lithuanian and Latvian coasts (Tabolina, 1994; Kjøie, 1999).

The lack of copepods in the diet of *N. melanostomus* at the study locations (Table 8) is a probable cause of the lack of cestode parasites observed in this study, since planktonic crustaceans are their primary intermediate hosts (Scholz, 1999). Together with the absence of cestode species, the lack of metazoan ectoparasites in this study is also contributing to the low species richness. The most likely reason is salinity as well as time since introduction. Most metazoan ectoparasites that have previously been found infecting the round goby in the Baltic Sea (crustacean *Ergasilus sieboldi*, Nordmann 1832 and the glochid stages of unionid bivalves) are freshwater species. The salinity in the sampling areas (5 to 7 PSU; Table 1), is thus probably too high for these species to occur. The monogenean *Gyrodactylus rugiensis*, on the other hand, is a marine parasite normally infecting *Pomatoschistus microps*, and it has previously been found infecting the round goby in salinities above 9 PSU in the Baltic (Kvach & Winkler, 2011). Monogeneans are in general quite host specific (Poulin, 2002; Huyse et al., 2003), and the introduction of the round goby in these locations happened 10–15 years ago, which is likely not enough time for most native monogeneans to adapt or switch host. The goby-specific monogenean, *Gyrodactylus proterorhini*, infects the round goby in its native areas in Black sea and Sea of Azov, but has to date not been observed in the Baltic round gobies (Mierzejewska et al., 2011). Other metazoan ectoparasites, mainly copepods, e.g. *Ergasilus sieboldi*, are found only occasionally even in

Table 8 The frequency of occurrence (%) of prey items (taxa) in the stomach contents of round gobies examined for parasites at four different locations in the Baltic Sea

Taxa	Guldborgsund			Palanga			Muuga			Mariehamn	
	Total 22	F 10	M 12	Total 18	F 7	M 11	Total 22	F 11	M 11	Total 21	M 21
<i>Mytilus</i>	4.5		8.3	22.2	14.3	27.3	13.6		27.3	9.5	9.5
<i>Macoma</i>							45.5	72.7	18.2	19.0	19.0
Bivalvia indet.	4.5		8.3								
Amphibalanus	9.1	10.0	8.3	50.0	14.3	72.7	54.5	27.3	81.8	42.9	42.9
Gastropoda	50.0	50.0	50.0	5.6		9.1	13.6	9.1	18.2	66.7	66.7
Amphipoda				61.1	28.6	81.8	18.2		36.4	9.5	9.5
Isopoda	9.1	10.0	8.3	22.2	28.6	18.2					
Decapoda	9.1	10.0	8.3	16.7		27.3				4.8	4.8
Polychaeta	22.7	30.0	16.7	11.1	28.6		9.1	9.1	9.1		
Oligochaeta	4.5	10.0		5.6	14.3		9.1	9.1	9.1	4.8	4.8
Insecta	18.2	10.0	25.0	5.6		9.1	18.2	9.1	27.3	9.5	9.5
Other	9.1	10.0	8.3								
<i>N. melanostomus</i> ^a	72.7	60.0	83.3	33.3	14.3	45.5	22.7	9.1	36.4	23.8	23.8
Pisces undetermined ^a	9.1		16.7				9.1	18.2		14.3	14.3
Pisces eggs	4.5	10.0									

At each location, the number of fish with stomach contents is given (*n*). All values are given separately for males and females as well as in total for each location

^a Scales and bones

its native area (Kvach, 2004b; Rolbiecki, 2006 and references therein; Özer, 2007). The only ectoparasite in this study, the hirudinea *P. geometra*, is very common in the Baltic Sea and has low host specificity as it infects several species of fish, e.g. eel (Rolbiecki, 2006), eelpout *Zoarces viviparus* (L., 1758), and Baltic flounder *Platichthys flesus* (L., 1758) (Køie, 1999). However, it has to be noted that the relatively small sample size (25 fish/site) can have caused some rare species not to be detected.

The most prevalent parasites in Guldborgsund, Muuga and Mariehamn, digenean trematodes, also correspond to the predominant prey items, i.e. gastropods, at these sites (FO = 50, 13.6, and 66.7%, respectively; Table 8). Likewise, the occurrence of amphipods in the stomach contents in Palanga (FO = 61.1%; Table 8) is seen in the presence of acanthocephalan parasites (Table 4). Although characterized as an opportunistic feeder (Rakauskas et al., 2008; Järv et al., 2011), the main components of the round goby diet are usually molluscs and adult barnacle *Amphibalanus improvisus* Darwin, 1854,

which can locally be a dominating prey species (Herlevi, unpubl. data; Table 8). *Contracaecum osculatatum* may infect larval stages of *A. improvisus*, but parasites which would use adult *A. improvisus* as an intermediate host are not known. There are also no other known predators for adult *A. improvisus*, although it is occasionally consumed by benthic predators as a fouling species on, e.g. *Mytilus trossulus* L., 1758 (Laudien & Wahl, 1999). A few freshwater digenean trematodes parasitizing fish and using bivalves as intermediate hosts have been found in the round goby (*Bucephalus polymorphus*, *Bunodera luciopercae*; Table 7), but these species require a freshwater bivalve species as vectors (*Dreissena polymorpha* Pallas, 1771 or *Pisidium* spp.). None of the common parasites infecting bivalves such as *Mytilus trossulus*, *Macoma balthica* L., 1758 or *Cerastoderma* spp. in the Baltic have been observed to infect round gobies (Zander & Reimer, 2002). As these species most commonly share habitat and are important prey of the round goby, it is possible that the round gobies are not exposed to many parasites

capable of infecting them. This can explain the low intensity of infection in the Baltic Sea. The diet composition of the round goby also explains some of the differences compared to native gobies, since other native goby species are parasitized mainly through ingesting intermediate hosts such as planktonic crustaceans, amphipods and oligochaetes. Thus, the reason for lower parasitization may be the round goby's main prey items and habitat (Zander, 2003; Emde et al., 2014).

The diet of the round goby and the regional differences therein thus seem to explain the intensity of parasitization and parasite assemblages observed in this study. The pattern of parasitization would indicate that the round goby has found a niche, which enables it to escape parasitization by many common parasites in the Baltic Sea. However, since the diet of the round goby has been shown to vary both seasonally and between size groups (e.g. Skora & Rzeznik, 2001; Rakauskas et al., 2008, 2013; Brandner et al., 2013b; Ustupis et al., 2015), future parasitological studies should investigate both seasonal and size-related differences in the infection rates of the round goby in the Baltic Sea.

Conclusions

The round goby acts as an intermediate and paratenic host for most of the parasite species found in this study. However, the presence of adult individuals of *Hysterothylacium aduncum*, *Camallanus* sp., and *Echinorhynchus gadi* shows that it can also function as a definitive host for these parasite species. Based on this study and others previously conducted in the Baltic Sea, it seems that overall the round goby has not retained parasites from its area of origin, but instead has been successively colonized by generalist parasites in the introduced areas. Although variable, overall parasite richness is still quite low in the Baltic compared to the native areas and a similar pattern can be seen in the prevalence and mean intensities of infection (Kvach, 2001, 2004b; Özer, 2007). The present study thus adds to the evidence supporting the enemy release hypothesis (e.g. Kvach & Stepien, 2008; Emde et al., 2014; Kvach et al., 2014), as no non-native parasites were detected, and the infection rates remain quite low throughout the invaded areas. The diet and opportunistic feeding behaviour, which

enable the round goby to exploit prey items largely unutilized by other predators, are probably a key factor for the observed low parasite intensities, and continued success as an invasive species in the Baltic Sea.

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