The occurrence of Echinorhynchus salmonis Müller, 1784 in benthic amphipods in the Baltic Sea

Benesh, Daniel; Aura, Raija-Liisa; Andersin, Ann-Britt; Valtonen, Tellervo


All material supplied via JYX is protected by copyright and other intellectual property rights, and duplication or sale of all or part of any of the repository collections is not permitted, except that material may be duplicated by you for your research use or educational purposes in electronic or print form. You must obtain permission for any other use. Electronic or print copies may not be offered, whether for sale or otherwise to anyone who is not an authorised user.
The occurrence of *Echinorhynchus salmonis* Müller, 1784 in benthic amphipods in the Baltic Sea

Daniel P. Benesh1,2, Raija-Liisa Aura1, Ann-Britt Andersin3 and E. Tellervo Valtonen1

1Department of Biological and Environmental Science, University of Jyväskylä, Finland; 2Marine Science Institute, University of California, Santa Barbara, CA, USA; 3Kappelisatamantie, Hanko Pohjoinen, Finland

**Abstract:** The acanthocephalan *Echinorhynchus salmonis* Müller, 1784 is a common parasite of salmonid fish, but it has rarely been reported from an intermediate host. Samples of benthic amphipods, *Monoporeia affinis* (Lindström), were taken from multiple, deep sites (usually below 70 m) in the Gulf of Bothnia over the course of more than a decade and examined for acanthocephalans. Overall, only 0.44% of 23,296 amphipods were infected, with just a single worm. This prevalence is consistent with several previous reports of acanthocephalans in deep-water, benthic amphipods, but it appears low compared to that often reported for acanthocephalan species infecting littoral amphipods. Parasite occurrence did not exhibit a clear regional pattern (i.e. northern vs southern sites) nor did it have any relationship with site depth. At sites sampled over multiple years, parasite abundance was consistently low (mostly <0.01), though two spikes in abundance (over 0.06) were also observed, indicating that infection can be substantially higher at particular times or in particular places. The median density of *E. salmonis* in samples containing the parasite was estimated as 8.4 cystacanths per m².

**Keywords:** Acanthocephala, aggregation, Gulf of Bothnia, cystacanth, density, Echinorhynchidae, *Echinorhynchus gadi*, intermediate host, repeatability

**MATERIALS AND METHODS**

The Baltic Sea is the world’s largest body of brackish water. Our sampling sites were mostly in the Gulf of Bothnia in the northern Baltic, which is characterised by lower salinity (<5 ppt) and no near-bottom oxygen deficiencies. Many taxa occurring in these areas are freshwater species (HELCOM 2009). Benthic amphipods were sampled from 12 sites in the Bothnian Bay (North Gulf of Bothnia), 4 sites in the Bothnian Sea (South Gulf of Bothnia) and one site in the Gulf of Finland (Fig. 1). A few sites were sampled multiple times, usually in different years. Sampling took place in early summer (32 of 45 samples were taken in June or

Address for correspondence: D.P. Benesh, Department of Biological and Environmental Science, POB 35, FI-40014 University of Jyväskylä, Finland. E-mail: daniel.benesh@lifesci.ucsb.edu
RESULTS

Monoporeia affinis was collected from all sites. Of 23,296 amphipods dissected, 103 were infected with the acanthocephalan Echinorhynchus salmonis (Table 1). All infections involved a single worm, so that the overall prevalence and abundance were equivalent (= 0.0044). Additionally, a second amphipod species was collected at the Gulf of Finland site, Pontoporeia femorata, and found to be infected with E. gadi (prevalence and abundance = 0.0088).

The distribution of parasite abundance across samples was clearly skewed (Fig. 2), such that no or few parasites were found in most samples, with a few samples having relatively high abundance (up to 0.128). Five sites were sampled multiple times (Table 1). The infection rate varied significantly between temporal samples in two of five sites (Fisher’s exact tests, site 1: χ² = 36.36, P = 0.008, E. gadi in site 17: χ² = 78.16, P < 0.001). However, when the sample with the highest infection rate in each of these two sites is excluded, then these tests were no longer significant (site 1: χ² = 12.85, P = 0.43, site 18: χ² = 0.59, P = 0.83), suggesting a pattern of temporal stability with an occasional spike in the infection rate (Fig. 3). In fact, these two ‘spikes’ were the only samples with prevalence above 3%.

Table 1. Samples of Monoporeia affinis (Lindström) from the Baltic Sea (for location of sampling site – see Fig. 1) examined for infection with Echinorhynchus salmonis Müller, 1784.

<table>
<thead>
<tr>
<th>Site</th>
<th>Latitude (N)</th>
<th>Longitude (E)</th>
<th>Depth (m)</th>
<th>Number of Amphipods dissected</th>
<th>Worms</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>65.14</td>
<td>25.34</td>
<td>15</td>
<td>8,084</td>
<td>56</td>
</tr>
<tr>
<td>2</td>
<td>65.00</td>
<td>25.15</td>
<td>4</td>
<td>924</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>64.42</td>
<td>25.06</td>
<td>1</td>
<td>33</td>
<td>0</td>
</tr>
<tr>
<td>4</td>
<td>64.34</td>
<td>25.20</td>
<td>1</td>
<td>120</td>
<td>0</td>
</tr>
<tr>
<td>5</td>
<td>64.25</td>
<td>25.55</td>
<td>1</td>
<td>85</td>
<td>0</td>
</tr>
<tr>
<td>6</td>
<td>64.04</td>
<td>25.55</td>
<td>b</td>
<td>402</td>
<td>2</td>
</tr>
<tr>
<td>7</td>
<td>64.56</td>
<td>22.21</td>
<td>2</td>
<td>1,684</td>
<td>12</td>
</tr>
<tr>
<td>8</td>
<td>64.48</td>
<td>23.29</td>
<td>1</td>
<td>112</td>
<td>0</td>
</tr>
<tr>
<td>9</td>
<td>64.17</td>
<td>22.21</td>
<td>10</td>
<td>3,304</td>
<td>7</td>
</tr>
<tr>
<td>10</td>
<td>64.14</td>
<td>22.36</td>
<td>9</td>
<td>432</td>
<td>2</td>
</tr>
<tr>
<td>11</td>
<td>64.04</td>
<td>21.27</td>
<td>9</td>
<td>211</td>
<td>1</td>
</tr>
<tr>
<td>12</td>
<td>63.59</td>
<td>21.46</td>
<td>1</td>
<td>854</td>
<td>0</td>
</tr>
<tr>
<td>13</td>
<td>61.07</td>
<td>18.00</td>
<td>b</td>
<td>233</td>
<td>3</td>
</tr>
<tr>
<td>14</td>
<td>61.11</td>
<td>18.14</td>
<td>74</td>
<td>634</td>
<td>9</td>
</tr>
<tr>
<td>15</td>
<td>61.10</td>
<td>18.50</td>
<td>65</td>
<td>326</td>
<td>0</td>
</tr>
<tr>
<td>16</td>
<td>61.05</td>
<td>19.35</td>
<td>126</td>
<td>1,006</td>
<td>7</td>
</tr>
<tr>
<td>17</td>
<td>59.51</td>
<td>23.16</td>
<td>3</td>
<td>4,847</td>
<td>0</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>-</strong></td>
<td><strong>-</strong></td>
<td><strong>45</strong></td>
<td><strong>23,296</strong></td>
<td><strong>103</strong></td>
</tr>
</tbody>
</table>

*a average across multiple samples; depth measurements were relatively consistent within sites; b coordinates were approximate for sites 6 and 13. Depth was not measured at these two sites, but it is presumably deeper than 70 m; c four samples of another amphipod species (Pontoporeia femorata Kroeyer) were taken at this site. Out of 1,695 dissected amphipods, 15 (0.88%) were infected with Echinorhynchus gadi Zoega in Müller, 1776.

spatial patterns. Generalised linear mixed effects models, with site as a random effect, binomial error distribution and a logit link (Wilson and Grenfell 1997, Bolker et al. 2009), were used to explore the effect of site depth and region (Bothnian Bay, Bothnian Sea, Gulf of Finland) on E. salmonis prevalence. All analyses were conducted in R 3.1.0 (R Development Core Team, Vienna, Austria).
The analysis of spatial patterns indicated that models with just ‘depth’ or just ‘region’ were slight improvements over models without these variables (likelihood ratio tests: $\chi^2_1 = 3.49, P = 0.06$ and $\chi^2_1 = 14.53, P < 0.001$, respectively). However, these effects were completely driven by the single Gulf of Finland site, which was the shallowest site and characterised by an apparent absence of *E. salmonis* (Table 1). When this site was removed, there was no relationship with depth ($\chi^2_1 = 0.004, P = 0.95$) and little difference between the northern and southern parts of the Gulf of Bothnia ($\chi^2_1 = 2.95, P = 0.09$; mean N vs S (95% CI): 0.38% (0.23–0.62%) vs 0.79% (0.36–1.75%)).

The estimated density of *M. affinis* (n = 38 samples) ranged from 237 to 4057 specimens/m² (median 1071) and there was no relationship between density and infection rate across samples (mixed model: $\chi^2_1 = 0.27, P = 0.60$). Amphipod density and parasite abundance were multiplied to estimate the density of larval acanthocephalans; when present in a sample, *E. salmonis* ranged from 2.1 to 44.8 specimens/m² (median 8.4).

**DISCUSSION**

Our data add to the handful of reports of *Echinorhynchus salmonis* (see Van Cleave 1920, DeGuisti and Budd 1959, Brownell 1970) and *E. gadi* (see Nybelin 1923, Valter et al. 1980, Valtonen et al. 1983) in various amphipod species. The overall prevalence of *E. salmonis* across all our Baltic Sea samples was 0.44%; the prevalence of *E. gadi* at site 17 was 0.88%. Several studies on acanthocephalans in amphipods from deep, benthic habitats have found comparable infection rates, e.g. 0.8% for *E. gadi* (see Marcogliese 1994), 0.1% for *E. salmonis* in lake Michigan (Amin 1978); 0.32–1.19% for several Antarctic species (Zdzitowiecki and Presler 2001, Laskowski et al. 2010). These values are low compared to those often reported from acanthocephalans in littoral lake or stream amphipods. For example, *Echinorhynchus truttae* Schrank, 1788, a parasite of brown trout, has been found at prevalences of 1 to 4% in amphipods (1–4% – Scheer 1935, 2.3% – Okaka 1984, 2.8% – Lassiere 1989, 4.1% – MacNeil et al. 2003, 1.3% – Dezfuli et al. 2008), whereas *Pomphorhynchus laevis* Zoega in Müller, 1776, a well-studied species usually infecting cypri
dinid fish, has often been reported to occur at prevalences above 10% in amphipods (Hine and Kennedy 1974, Brown and Pascoe 1989, Moravec and Scholz 1991, Dezfuli et al. 1999). In contrast, Okaka (1984) reported a low prevalence of *E. salmonis* (0.9%) in gammarids from a shallow river, while Nybelin (1924) found a high prevalence (8.8%) of *E. salmonis* in 160 *M. affinis* sampled off the east coast of Sweden (exact sampling location and depth were not given, but presumably it was sublittoral). Green (1965) also noted up to 12% prevalence of *E. salmonis* in the largest size class of *M. affinis* collected from the bottom of a glacial lake in New York. However, only 1–2% of individuals in Green’s (1965) samples were in the largest size class, so the overall prevalence of *E. salmonis* in his study was considerably lower (we roughly calculated it as about 1%). Thus, acanthocephalan infection rates are not always lower in benthic amphipods compared to littoral amphipods, and additional studies are needed to assess whether there is a general difference between habitats.

Although only 1 in 227 *M. affinis* was infected on average, prevalence was much higher in a few samples. The aggregated distribution of abundances across samples is a typical pattern in ecology (Taylor 1961, Taylor et al. 1978, Gaston et al. 2006), and it has also been noted for parasites (Morand and Guéant 2000, Krasnov et al. 2006, Pérez-del-Olmo et al. 2011). Both spatial and temporal variation in abundance of larvae *E. salmonis* contributes to this pattern. Sites that were sampled multiple times exhibited consistent, low parasite abundance, but occasional increases in
abundance were observed. Some sites may also be more or less conducive to parasite existence. For example, the parasite was not found at the shallow, near-coast Gulf of Finland site, despite intense sampling, while two sites in the Bothnian Sea seemed to have higher prevalence (Fig. 1, Table 1). However, we did not find infection rates to differ consistently across the Gulf of Bothnia (North vs South) or with sampling depth. These analyses had rather low power, though, and would benefit from further sampling, e.g. in the Bothnian Sea and at additional shallow sites. Valter et al. (1980) also found prevalence of *E. gadi* in amphipods to vary widely (from 0.5 to 6.3%), although the distribution of prevalence values was not explicitly investigated.

The precise spatiotemporal factors leading to higher larval acanthocephalan abundance remain to be determined. Environmental variables, like salinity and oxygen, vary across the study area, perhaps influencing parasite transmission. For example, *E. gadi* was only found at the southernmost site, presumably because the higher salinity there suits *P. femorata* (see Laine 2003) and cod, an important definitive host (Valtonen and Crompton 1990). Amphipod populations in the northern Baltic also exhibit both season- nal and long-term fluctuations in response to nutrient input (Lehtonen and Andersin 1998). However, parasite abundance appeared unrelated to the density of amphipods in the sample, so it is not obvious that there is a tight coupling between amphipod and acanthocephalan populations. Similarly, Valtonen et al. (2004) found that the abundance of three seal acanthocephalans in the Bothnian Bay varied little in response to sizeable fluctuations in the intermediate and definitive host populations.

Our study is one of the few to examine the distribution of an acanthocephalan in a benthic amphipod. We found that very low infection rates are the norm for this system (87% of our samples exhibited prevalence less than 1%), but we also found that at particular times or in particular places, infection can be substantially higher. This heterogeneous pattern of occurrence in amphipods is expected to translate into a more aggregated distribution in the fish definitive host (Janovy and Kutish 1988, Lotz et al. 1995). In the study area, *E. salmonis* is indeed overdetermined in its main definitive hosts (Valtonen and Crompton 1990). Infection rates in fish are also much higher than in amphipods, e.g. in *Coregonus lavaretus* widgreni Malmgren: prevalence 44% and abundance 9.9 (Valtonen and Crompton 1990). So even though the density of larval acanthocephalans seems quite low (at a maximum, just tens of individuals per m$^2$ vs thousands of amphipods per m$^2$), it is sufficient to maintain sizeable adult acanthocephalan populations in fish.

**Acknowledgements.** We thank the Tvärminne Zoological station, Helsinki University, for material and the Finnish Environment Institute (SYKE) for help with collecting and collating the data. Helpful reviewer feedback substantially improved the manuscript. DPB gratefully acknowledges support from the DFG (BE 5336/1–1).

**REFERENCES**


