

Levels of Helminth Infection of Small Rodents in Two Interspersed Habitats – the Forest and Beaver Sites

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Abstract

Helminth infection in small rodents (*Apodemus flavicollis* and *Myodes glareolus*) was studied in forest and in beaver sites that were tightly spatially interspersed. Beaver sites were regarded as disturbed habitats inhabited by these two typical forests dwelling small rodent species. We tested the hypothesis that beaver sites, as disturbed habitats, can influence the abundance of infections with helminths, typical of forest rodent species. The composition of helminth species overlapped fully in the two tested habitats for *M. glareolus* (8 species of parasites) and nearly completely also for *A. flavicollis* (7 species at beaver sites and 6 species in the forest). *M. glareolus* were more heavily infected with *Syphacia petrusewiczii* at beaver sites than in the forest in autumn and winter. The mean abundance of all helminths in *M. glareolus* was significantly higher at beaver sites only in the winter, with no significant differences in any other season. For *A. flavicollis*, we did not find statistically significant differences in mean abundance of helminths between the habitats neither for any particular helminth species nor for all helminth species taken together. We found some changes in the helminth community structure (as reflected in dominance hierarchies of helminths species) between the two studied habitats. *S. petrusewiczii* dominating markedly over the other helminth species in *M. glareolus* at the beaver sites, whereas dominance with this helminth was much weaker in the forest. For *A. flavicollis*, we found different dominant helminth species in each of the two habitats: *Syphacia montana* was strong dominant at beaver sites, but it was not found in mice dwelling in the forest, where *Syphacia stroma* dominated at the beaver sites. The patterns of taxonomic diversity of helminths and the distribution of parasites among host individuals did not reveal any significant differences between the tested habitats which were variable among host species and seasons. Our study has revealed habitat differences only in some aspects of the helminth infections of the two typical forest dwelling rodent species, whereas other tested parameters were highly variable and did not show significant inequalities between habitats. These findings suggest a rather weak impact of the beaver sites on the epidemiology of helminth infections in typical forest dwelling small rodent species.

Key words: *Apodemus flavicollis*, *Myodes glareolus*, helminths, helminth infection levels, forest, beaver sites, habitat disturbance

Introduction

Small rodents are good models for investigating the host-parasite relationship in different habitats due to abilities of these mammals to inhabit a wide range of different environments (Lafferty et al. 2008). Infections of small rodents caused by helminths are usually studied in separate habitats, such as forests, grasslands, mountains, rural and urbanized environments (Behnke et al. 2000, Ferrari 2005, Mažeika et al. 2003, Montgomery and Montgomery 1989) but few data exist on comparative helminthological analysis of small rodents in different habitats or habitat complexes with respect of their cardinal changes.

Forest habitats, despite their very uneven species composition and succession stage, can be regarded as the last stages of succession within a landscape ecosystem in temperate zones, thus, suggesting a rel-

atively long development of ecological relationships among various trophic levels, including the host-parasite system in small rodents. Various ecological factors might be considered to have the potential to influence the host-parasite relations, especially when the mature forest ecosystems are disturbed in some way.

Beavers (*Castor fiber* and *C. canadensis*) are widely acknowledged as a source of natural disturbance in forest ecosystems (Johnston and Naiman 1987, King and Antrobus 2001, Wright et al. 2002). Generally, beaver sites have very specific habitat qualities, and their numbers have increased significantly in the landscape ecosystem during recent decades (Ulevičius 2008). Beaver sites are usually tightly interspersed with forest habitats, thus, providing an opportunity for colonization by species of forest small rodent. The bank vole (*M. glareolus*) and the yellow-necked mouse (*A. flavicollis*) are considered as typi-

cal forest rodents (Juškaitis 1999, Juškaitis et al. 2001, Mažeikytė et al. 1999, Prūsaitė 1988) also inhabiting the environments of beaver sites with high relative abundance (Ulevičius and Janulaitis 2007).

Due to the wet and swampy conditions of beaver sites and abundant intermediate hosts of helminths (snails and other mollusks) the parasitological state of small rodents may be different than in the forest (Ferrari 2005, Pulido-Flores et al. 2005). Mobile rodents have the potential to transfer helminths to the forest and other habitats, in this way influencing the parasitological state of the whole landscape ecosystem. The relationships between the individual biota components of the biota of beaver modified habitats, including the host-parasite system, have not been investigated across the whole range of the Eurasian beaver.

The aim of this study was to compare the helminth fauna notably the abundance of helminths in small rodents in the forest and beaver sites. We tested the hypothesis that beaver sites, as disturbed habitats, influence infection levels of helminths, typical of the forest rodent species.

Study area

The study sites were located in Vilnius, Molėtai and Širvintos districts, Eastern Lithuania (Fig. 1). The geographic co-ordinates of the approximate centre of this territory: 55°00'N, 25°14'E.

Morainic hills with numerous lakes are characteristic of the study area. Forests are highly fragmented (Fig. 1) and successional mixed stands (usually *Picea abies* with some deciduous species, *Alnus incana*, *Populus tremula*, *Betula pendula*) prevail. Average forest cover of the study area is ca. 29%. A number of abandoned meadows and extensively used pastures intersperse with fragmented forests. Fens, usually overgrown or fringed by *Salix* spp., *Alnus* spp. and *Betula* spp. stands, are common in depressions between hills. The majority of beaver sites are located

in fens at the margins of small forests. At beaver sites, usually *Salix* spp. or *Frangula alnus* shrubs grow, and in the grass layer *Carex* spp. dominate. Mean density of beaver sites was 4.5 sites/1000 ha. The age of beaver sites varied between 10 – 25 years. Beavers disturb forests by raising the water level of the ground and leaving trees to die or removing the majority of large trees by cutting them.

Materials and methods

Small rodents were sampled in the forest and at beaver sites from 2007 to 2009 four times per year: in spring (April), in summer (July – August), in autumn (October) and in winter (February). Snap traps were set in standard lines (25 traps at intervals of 5 meters) or trap quadrates (five traps, one in the centre and four in the corners of an approximately 3x3 m square) for three days and nights. Trap lines and trap quadrates were used in the forest, whereas trap quadrates were used only at beaver sites. Trap quadrates were set on beaver lodges (one trap on the top and four around the base of a lodge) (Ulevičius and Janulaitis 2007). Small pieces of brown bread crust moistened with sunflower oil were used as bait. Sampling of small mammals by snap traps were permitted by the Ministry of Environment (license No. (11-)-D8-3650).

In total 390 individual small rodents were caught and examined helminthologically: *M. glareolus* – n = 128 at beaver sites and n = 159 in the forest; *A. flavicollis* – n = 21 and n = 82, respectively. Small rodents of other species were caught but these were not studied helminthologically due to their low relative abundance and frequency of occurrence.

We dissected the entire intestinal tracts of small rodents and examined these for helminths. The content of the intestines was studied by the method of consistent flushing. The helminths were fixed in 70% ethanol. Nematodes and trematodes were studied on temporary water – glycerin preparations (Ивашкин и др. 1971).

Two indices of infection level were used. Mean abundance is the total number of individuals of particular parasite species in a sample of a particular host species divided by the total number of hosts of that species examined (Bush et al. 1997). The other indicator used was prevalence of infection which was calculated as the percentage of the infected individuals from among all the dissected rodents (Bush et al. 1997). Significance of differences of mean abundance and prevalence of infection was tested using the Mann-Whitney U test and the Chi-square test, respectively ($p < 0.05$).

Age (juveniles, subadults, adults) and gender difference were tested using tests of independence on

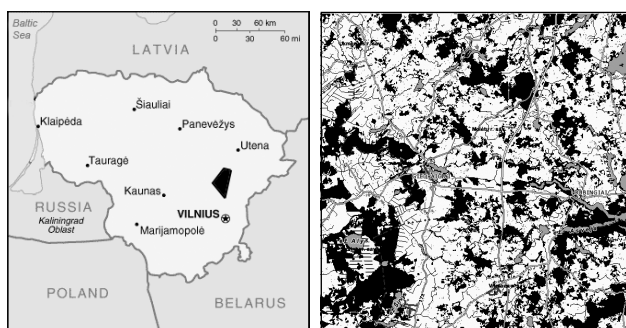


Figure 1. Study area. The black polygon on the left map covers the whole study area. On the right map is shown the characteristic pattern of the forest (black patches) fragmentation in the study area

infection levels of *M. glareolus* (because of larger samples than *A. flavicollis*) within seasons. Statistically significant differences among age groups were found only in the autumn (Kruskal-Wallis test, $p=0.006$). Statistically significant differences between sexes of this host were found only in summer (Mann-Whitney test, $p=0.01$). In the remaining cases, we did not find statistically significant differences neither among age nor sex groups of *M. glareolus* probably due to remarkably smaller samples of individuals tested in other particular seasons. Thus, we have omitted the age and sex factors from further analysis by pooling together all groups of age and sex within seasons.

The Shannon – Wiener diversity index (H') (May 1975) and the Berger–Parker dominance index (May 1975) were used to express diversity of the helminth species and dominance of helminth species in small rodents from different habitats. The Jaccard's similarity coefficient (J) (Magurran 1988), which measures numerical similarity of helminth species, shared between component communities. The helminth aggregation was assessed using the variance-to-mean ratio (dispersion coefficient, CD) of helminths number per one host individual (Poulin 1998). Statistical significance of differences from random distribution of helminths ($CD=1$) (was tested using the Student's t -test (Kershaw 1978).

Results

In total, including both small rodent species (*M. glareolus* and *A. flavicollis*) and both treatments (beaver sites and forest), 66 % of the examined small rodent individuals were infected by helminths. Helminths of 9 species and 3 not identified to species level (5821 specimens) were found: 1 cestode (tapeworms), 6 nematodes (roundworm), 2 trematodes (fluke) species and 1 cestode and 2 nematodes not identified to species level (Table 1).

M. glareolus were infected by 8 species of helminths and there was no difference in the number of helminth species between *M. glareolus* caught at beaver sites and in the forest. All parasite species were the same in both habitats for *M. glareolus* (Table 1).

A. flavicollis were infected by 7 helminth species at beaver sites and 6 in the forest. The nematode *S. montana* was found only in mice dwelling at beaver sites (Table 1).

Comparing the two rodent species, they had three common taxa of helminths (*Cestoda g. sp.*, *Heligmosomum costellatum*, *Heligmosomum mixtum*), whereas other helminths were found to be rodent-specific. This pattern was identical between the two tested habitats (Table 1).

Table 1. Helminth species (or unidentified taxa) composition and their distribution among hosts from beaver sites and the forest

Class	Family	Helminth species or unidentified taxa	Life cycles	Beaver sites		Forest	
				<i>Myodes glareolus</i>	<i>Apodemus flavicollis</i>	<i>Myodes glareolus</i>	<i>Apodemus flavicollis</i>
Cestoda	Taeniidae	<i>Hydatigera taeniaeformis</i>	Indirect	+	-	+	-
		<i>Cestoda g. sp.</i>		+	+	+	+
	Capillariidae	<i>Capillaria sp.</i>		+	-	+	-
	Heligmosomatidae	<i>Heligmosomum costellatum</i>		+	+	+	+
<i>Heligmosomum mixtum</i>			+	+	+	+	
Nematoda	Syphaciidae	<i>Syphacia montana</i>	Direct	-	+	-	-
		<i>Syphacia petruszewiczi</i>		+	-	+	-
	<i>Syphacia stroma</i>	-		+	-	+	
	<i>Syphacia sp.</i>	-		+	-	+	
Trichocephalidae	<i>Trichocephalus muris</i>		+	-	+	-	
Trematoda	Plagiorchiidae	<i>Plagiorchis elegans</i>	Indirect	-	+	-	+
	Notocotylidae	<i>Notocotylus noyeri</i>		+	-	+	-
Total number of helminth species				8	7	8	6

Analyzing particular species of helminths in different seasons, the mean abundance index showed only slight and statistically insignificant differences between habitats (Table 2), except two cases, when *M. glareolus* were more heavily infected by *S. petruszewiczi* at beaver sites than in the forest (Mann-Whitney test: autumn, $p = 0.03$ and winter, $p = 0.02$). In other cases, the mean abundance of helminths in *M. glareolus* and *A. flavicollis* were not statistically significant between habitats (Table 2).

Total mean abundance of all parasites in *M. glareolus* showed a higher infection level of this rodent at beaver sites than in the forest, especially in winter (Mann-Whitney test: winter, $p = 0.049$). A similar tendency was observed in *A. flavicollis* showing higher abundance of all parasites in autumn; however these differences between habitats were not statistically significant (Table 2).

Similarly, analysis of the prevalence of infection did not reveal any significant differences between habitats in different seasons, except in two cases (Table 3). A contrasting pattern of prevalence of infection was found in *M. glareolus* infected by *S. petruszewiczi* and *H. mixtum*. This index of *S. petruszewiczi* in *M. glareolus* in autumn was significantly higher at beaver sites ($\chi^2 = 9,283$, $df = 1$, $p = 0.0023$), and prevalence of *H. mixtum* in the same host species and in season was higher in the forest ($\chi^2 = 6.493$, $df = 1$, $p = 0.0108$). For the rest of the parasite species differences in infection prevalence between two habitats in different seasons were not significant neither in *M. glareolus* nor *A. flavicollis* (Table 3).

Table 2. Mean abundance of helminths in small rodents at beaver sites and in the forest in different seasons (n – number of dissected individuals of small rodents, B – beaver sites, F – forest)

Helminth species or unidentified taxa	Habitat	<i>Myodes glareolus</i>				<i>Apodemus flavicollis</i>			
		Spring n=23(B) n=22(F)	Summer n=26(B) n=43(F)	Autumn n=66(B) n=80(F)	Winter n=13(B) n=14(F)	Spring n=3(B) n=12(F)	Summer n=8(B) n=18(F)	Autumn n=10(B) n=46(F)	Winter n=0(B) n=6(F)
<i>Capillaria</i> sp.	B	-	-	0.3	-	-	-	-	-
	F	-	-	0.3	-	-	-	-	-
<i>Cestoda</i> g. sp.	B	0.09	0.07	0.1	0.2	0.3	-	0.2	-
	F	0.1	0.2	0.09	0.07	0.3	0.06	0.1	-
<i>Heligmosomum mixtum</i>	B	6.3	0.5	1.2	4.2	1.3	0.3	1.0	-
	F	4.6	1.4	2.8	2.4	3.0	0.7	0.4	4.7
<i>Heligmosomum costellatum</i>	B	-	1.0	1.1	-	-	0.3	-	-
	F	-	2.6	0.8	-	-	0.2	0.4	-
<i>Hydatigera taeniaeformis</i>	B	0.09	-	-	-	-	-	-	-
	F	0.05	-	-	-	-	-	-	-
<i>Notocotylus noyeri</i>	B	13.4	0.7	0.08	-	-	-	-	-
	F	11.0	0.09	0.01	-	-	-	-	-
<i>Plagiorchis elegans</i>	B	-	-	-	-	-	0.4	-	-
	F	-	-	-	-	-	0.2	0.04	-
<i>Syphacia montana</i>	B	-	-	-	-	-	12.3	2.0	-
	F	-	-	-	-	-	-	-	-
<i>Syphacia petruszewiczi</i>	B	6.1	18.6	8.9*	9.8*	-	-	-	-
	F	-	14.2	2.4*	0.1*	-	-	-	-
<i>Syphacia stroma</i>	B	-	-	-	-	-	3.8	1.0	-
	F	-	-	-	-	2.8	11.8	3.6	7.7
<i>Syphacia</i> sp.	B	-	-	-	-	10.7	-	-	-
	F	-	-	-	-	0.3	0.7	4.0	0.5
<i>Trichocephalus muris</i>	B	0.04	-	0.02	0.08	-	-	-	-
	F	-	-	0.01	0.07	-	-	-	-
Totally for all helminth species	B	26.0	20.5	12.2	14.2	1.7	16.9	22.5	-
	F	15.8	18.5	6.6	2.7	6.4	13.6	8.5	12.8

* – statistically significant

Table 3. Infection prevalence (%) of helminths in small rodents at beaver sites (B) and in the forest (F) in different seasons (N - number of dissected individuals of small rodents)

Helminth species or unidentified taxa	Habitat	<i>Myodes glareolus</i>				<i>Apodemus flavicollis</i>			
		Spring n=23(B) n=22(F)	Summer n=26(B) n=43(F)	Autumn n=66(B) n=80(F)	Winter n=13(B) n=14(F)	Spring n=3(B) n=12(F)	Summer n=8(B) n=18(F)	Autumn n=10(B) n=46(F)	Winter n=0(B) n=6(F)
<i>Capillaria</i> sp.	B	-	-	4.5	-	-	-	-	-
	F	-	-	5.0	-	-	-	-	-
<i>Cestoda</i> g. sp.	B	8.7	8.3	3.0	23.1	33.3	-	10.1	-
	F	9.1	20.9	8.8	7.1	33.3	5.6	6.5	-
<i>Heligmosomum mixtum</i>	B	43.5	19.2	21.1*	77.0	66.7	12.5	30.0	-
	F	68.2	32.6	42.5*	57.1	25.0	11.1	13.0	33.3
<i>Heligmosomum costellatum</i>	B	-	7.7	9.1	-	-	12.5	-	-
	F	-	20.9	10.0	-	-	5.6	6.5	-
<i>Hydatigera taeniaeformis</i>	B	4.5	-	-	-	-	-	-	-
	F	4.5	-	-	-	-	-	-	-
<i>Notocotylus noyeri</i>	B	17.4	7.7	1.5	-	-	-	-	-
	F	18.2	2.3	1.3	-	-	-	-	-
<i>Plagiorchis elegans</i>	B	-	-	-	-	-	25.0	-	-
	F	-	-	-	-	-	5.6	2.2	-
<i>Syphacia montana</i>	B	-	-	-	-	-	25.0	10.1	-
	F	-	-	-	-	-	-	-	-
<i>Syphacia petruszewiczi</i>	B	13.0	30.8	25.6*	30.8	-	-	-	-
	F	-	14.0	6.3*	14.3	-	-	-	-
<i>Syphacia stroma</i>	B	-	-	-	-	-	25.0	10.1	-
	F	-	-	-	-	33.3	5.6	13.0	16.7
<i>Syphacia</i> sp.	B	-	-	-	-	33.3	-	-	-
	F	-	-	-	-	16.7	5.6	15.2	16.7
<i>Trichocephalus muris</i>	B	4.3	-	1.5	7.7	-	-	-	-
	F	-	-	1.3	7.1	-	-	-	-
Totally for all helminth species	B	73.9	61.5	60.6	76.9	100.0	75.0	50.0	-
	F	86.4	74.4	70.0	64.3	75.0	33.3	34.8	50.0

* – statistically significant

Total prevalence of all helminths combined (all parasites) also did not differ significantly between the two tested habitats, although there were some differences between two rodent species with respect to this: the percent of infected *M. glareolus* was similar between habitats in all seasons, whereas the percent of infected *A. flavicollis* was somewhat higher (statistically not significant) at beaver sites in all seasons (Table 3).

There were 3 dominant species of helminths in *M. glareolus* and 4 species in *A. flavicollis*. The species abundance patterns of helminth species in *M. glareolus* varied more among seasons but less between the two tested habitats, except in autumn and winter (Table 4). *S. petruszewiczi* dominated in *M. glareolus* at beaver sites in summer, autumn and winter, whereas it was numerically dominant in the forest only in summer. These rodents contained the same dominant

Discussion and conclusions

The composition of helminth communities overlapped completely in the two tested habitats for *M. glareolus* (8 species of parasites) and nearly completely in *A. flavicollis* (7 species at beaver sites and 6 species in the forest). Among helminths infecting *M. glareolus*, only *S. petruszewiczi* was significantly more abundant at beaver sites than in the forest. No statistically significant differences were found for helminths of *A. flavicollis* with respect to habitat.

Taking into account total infection of rodents regardless of helminth species, the forest might be considered as a more optimal habitat than the disturbed environments of beaver sites at least for *M. glareolus*, as total mean abundance of parasites was significantly higher for this rodent species at beaver sites in winter. For *A. flavicollis*, we observed only tendencies towards higher abundance of parasites at beaver sites, these tendencies being more clearly pronounced in autumn. Infection prevalences also revealed a few significant differences between habitats but only in *M. glareolus* involving particular species of parasites, but no difference when all helminth species were pooled together.

During our research some helminth species were not identified to species level but species richness was similar in both small rodent species (8 species in *M. glareolus* and 7 in *A. flavicollis*). At a larger regional scale, many more helminth species are recovered e.g. on the territory of Lithuania, more than 20 helminth species were described parasitizing *M. glareolus* and 10 helminth species found in *A. flavicollis* by Prūsaitė (1988), and 42 in *M. glareolus* and 17 in *A. flavicollis*, respectively, by Mažeika (1992). However, at a local scale helminth species richness can be somewhat lower. In a mountain locality of Serbia, *M. glareolus* were infected by seven species of nematodes (Bjelić-Čabrilović et al. 2009). Eleven species of helminths were found infecting *M. glareolus* in three localities in Northern Poland (Behnke et al. 2001). Similar helminth species numbers were reported for *A. flavicollis* on a local territory (Klimpel et al. 2006). In our research, helminth species fully overlapped in two tested habitats for *M. glareolus* and almost fully for *A. flavicollis*. This finding suggests that the species composition of a helminth community in small rodents at a local scale might not be affected by habitat.

Different species of parasites were sometimes characterized by different patterns of infection in rodent in both habitats tested e.g., the most pronounced differences of infection level (prevalence of infection) of *M. glareolus* by two different species of helminths (*S. petruszewiczi* and *Heligmosomum mixtum*) were

different in the two habitats. It might be expected these parasite species having different habitat requirements when infecting the same host species e. g., the crucial factor of reproduction success is humidity for some species of *Syphacia*, and temperature is critical for *Heligmosomum mixtum* (Haukisalmi and Henttonen 1999, Определитель ... 1979). In other studies, the *M. glareolus* infection levels by these two parasites were found to be different in similar habitats from different neighbouring localities (Behnke et al. 2001, Kuliś-Małkowska 2007), indicating probable intrinsic (age, sex) and extrinsic (time, season) factors in both the parasite and the host populations, as well as synergistic and antagonistic interactions between parasite species (Behnke et al. 2005, Ferrari et al. 2003, Ferrari et al. 2004, Kuliś-Małkowska 2007).

We found changes in the numerical composition of helminth communities between habitats. Generally, *S. petruszewiczi* strongly dominated other helminth species in *M. glareolus* at beaver sites, whereas domination of this helminth was much weaker in the forest. For *A. flavicollis*, we found completely different dominant species among the two habitats. *S. montana* was strongly dominant at beaver sites but it was not found in mice dwelling in the forest where *S. stroma* dominated.

The domination of one species in a helminth community is rather a common phenomenon and usually one or few species dominate over many other species which are present at low abundance levels (Poulin et al. 2008). Presence or absence of a helminth species may be influenced by abiotic factors to the intermediate hosts (Krasnov et al. 2008), and difference between the species in patterns of helminth domination can be explained by the peculiarities of life cycles in different habitats.

The diversity indices of the helminth community varied considerably among seasons. Nevertheless, we cannot find any regularity in these seasonal patterns that would logically explain differences between habitats. This suggests that parasite diversity varied randomly in both tested habitats.

Our findings did not reveal any obvious differences in parasite distribution among host individuals between habitats and seasons. Parasite distribution among host individuals is determined by many factors (Anderson and Gordon 1982). In our study, helminths of *M. glareolus* and *A. flavicollis*, were mainly aggregated distribution in their host populations (Haukisalmi and Henttonen 1999). The reasons for aggregated distribution in parasites are not well understood and can include: 1) host age at first contact with helminths; 2) social status of host individuals; 3) infection with other helminth species; 4) genetics of host and para-

site; 5) diet of host; and, 6) host behavior, as well as many other factors (Anderson 1991). Parasites of small rodents are characterized by an aggregated distribution in host populations, which is considered one of the stabilizing factors of the host-parasite interactions (May and Anderson 1978, Wakelin 1986). In addition, the aggregate distribution of helminths can reduce interactions among helminths (Wakelin 1986).

H. taeniaeformis was found at the larval stage (strobilocercus fasciolaris) in *M. glareolus*. The definitive host of this parasite is a carnivorous mammal but it can also be found in humans. The intermediate host of *H. taeniaeformis* is usually a rodent (Козлов 1977). In Lithuania, this helminth is also found in *Mus musculus*, *Apodemus agrarius*, *A. flavicollis*, *Microtus arvalis*, and *Rattus norvegicus* in Kėdainiai district and Vilnius suburbs (Mažeika 1992), and *Ondatra zibethicus* in Rusnė Island (Mažeika 2009). We did not find any differences at infection levels of this parasite in the bank vole between habitats, and thus beaver sites cannot be regarded as more risky habitats than forests for human or predator health with respect of the *H. taeniaeformis* infection.

Our study revealed differences in some aspects of helminth infection in two forest rodent species (*M. glareolus* and *A. flavicollis*) inhabiting relatively undisturbed forest and beaver sites that intersperse with the forest fragments. This holds true for differences of helminth abundance in *M. glareolus* in winter, as well as helminth community structure (expressed by domination of helminth species) between the two studied habitats for both rodent species. Many other parameters examined were highly variable but did not show significant inequalities between habitats. These findings suggest a rather weak impact of beaver sites on a parasitological state of forest rodent species.

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УРОВЕНЬ ЗАРАЖЕНИЯ ГЕЛЬМИНТАМИ МЕЛКИХ ГРЫЗУНОВ В ДВУХ ПЕРЕМЕЖНЫХ СРЕДАХ ОБИТАНИЯ – В ЛЕСУ И БОБРОВЫХ ПОСЕЛЕНИЯХ

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Резюме

Уровень заражения гельминтами мелких грызунов (*Apodemus flavicollis* и *Myodes glareolus*) изучали в лесу и бобровых поселениях, в средах, которые тесно перемежаются между собой, а бобровые поселения могут рассматриваться как своеобразные нарушения лесной среды обитания двух типичных мелких грызунов леса. Мы тестировали предположение о том, что бобровые поселения, являясь нарушенными биотопами, могут влиять показатели зараженности гельминтами двух типичных мелких грызунов леса. Видовой состав гельминтов полностью перекрывался между бобровыми поселениями и лесом у *M. glareolus* (8 видов паразитов) и почти полностью у *A. flavicollis* (7 видов в бобровых поселениях и 6 видов в лесу). Анализируя отдельных видов гельминтов, *M. glareolus* были достоверно сильнее заражены только *Syphacia petrusewiczii* в бобровых поселениях нежели в лесу осенью и зимой. Средняя численность всех гельминтов вместе взятых у *M. glareolus* было достоверно выше в бобровых поселениях только зимой, без достоверных различий между биотопами в другие сезоны. У *A. flavicollis* мы не нашли достоверных различий ни по средней численности гельминтов, ни по отдельным видам ни по их совокупности между биотопами. Были отмечены различия по структуре сообщества гельминтов между тестируемыми биотопами. *S. petrusewiczii* сильно доминировало над другими гельминтами у *M. glareolus* в бобровых поселениях летом, осенью и зимой, в то время как в лесу – только летом. У *A. flavicollis* были обнаружены совсем другие доминанты в сравниваемых биотопах: *Syphacia montana* доминировало в бобровых поселениях, но вообще не обнаружено в лесу, где доминантом было *Syphacia stroma*. Небыло выявлено существенных различий между биотопами ни по разнообразию гельминтов, ни по их распределению между индивидами хозяев. Эти показатели сильно варьировали между видами хозяев и между сезонами. Наши исследования показали средовые различия только по нескольким аспектам заражения гельминтами двух типичных видов лесных грызунов. Многие другие параметры их заражения были крайне изменчивыми и не показали значимых различий между биотопами. Полученные результаты предполагают скорее слабое влияние бобровых поселений на паразитологический статус здесь обитающих типичных грызунов леса.

Ключевые слова: *Apodemus flavicollis*, *Myodes glareolus*, гельминты, экстенсивность заражения, интенсивность заражения, сезонная динамика, лес, бобровые поселения, нарушение среды обитания.