ECOLOGY OF TRICHINELLOSIONS TRANSMISSION IN THE VORONEZH STATE NATURE RESERVE AND ADJACENT AREAS, RUSSIA

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Received: 12.05.2020. Revised: 22.01.2021. Accepted: 08.02.2021.

The article considers certain aspects of the morphology and biology of \textit{Trichinella}, and the ecology and epizootology of trichinellosis in the Voronezh State Nature Reserve and its adjacent areas (Black Soil Region of Russia). Original data were collected during 30 years (1990–2019) from potential hosts of \textit{Trichinella}, mainly carnivorous mammals. During this period, more than 200 specimens of wild and domestic carnivores of three families (Canidae, Mustelidae, Felidae) were studied. Six species of wild carnivores (\textit{Canis lupus}, \textit{Vulpes vulpes}, \textit{Nyctereutes procyonoides}, \textit{Meles meles}, \textit{Martes martes}, and \textit{M. foina}) were obligatory hosts. In addition, \textit{Trichinella} was found in \textit{Erinaceus concolor} and two species of domestic carnivores (\textit{Canis lupus familiaris} and \textit{Felis catus}). The highest prevalence of infection was observed in \textit{Vulpes vulpes}, \textit{Nyctereutes procyonoides}, \textit{Meles meles}, and \textit{Martes martes} (35.7–70.0%). These hosts play a leading role in the natural trichinellosis transmission. The carnivores can be divided into two groups according to the intensity of infection by \textit{Trichinella} larvae per gram of muscles (lpg). The first group includes native carnivore species (\textit{Vulpes vulpes}, \textit{Canis lupus}, \textit{Meles meles}, \textit{Martes martes}, and \textit{M. foina}), with an average of 10 lpg. The second group includes introduced species (\textit{Nyctereutes procyonoides}) and invasive species (immigrants) from anthropogenic areas (\textit{Felis catus}), with an average of 700 lpg. Due to its high pathogenicity, \textit{Trichinella} can be considered as a significant factor in regulating the number of alien carnivorous species. \textit{Trichinella} is also characterised by aggregated distribution in the muscles. In the Voronezh State Nature Reserve, the highest lpg values were observed in the muscles of the front and rear limbs of the carnivores. Over 50% of the local hemipopulation of \textit{Trichinella} is concentrated in these muscle types. Based on the assessment of the morphological features of the \textit{Trichinella} capsules and larvae from carnivores, two clusters of \textit{T. nativa} hosts were identified. These clusters comprise carnivorous species that have closer trophic relationships. The variability of quantitative and qualitative parameters in the hosts indirectly reflects the specifics of the relationships in the host-parasite system and shows nutritional preferences of the studied carnivores. In the Voronezh State Nature Reserve and its adjacent areas, \textit{Vulpes vulpes} is the main link in the stable transmission of trichinellosis. \textit{Vulpes vulpes} has high rates of infection intensity (35.7%) and large hemipopulations of \textit{Trichinella} larvae (3.3 million specimens). \textit{Vulpes vulpes} is the most abundant carnivorous species in the Voronezh State Nature Reserve and the most common hunting prey. So it can be used for monitoring the trichinellosis in European Russia. In the Voronezh State Nature Reserve, invertebrate animals can also contribute to the trichinellosis transmission. Carabid beetles (Carabidae) were registered as the disseminators of \textit{T. nativa}. At present, in the Voronezh State Nature Reserve and its adjacent areas, \textit{Trichinella} is transmitted by wild carnivores. The dominant position in the parasitic system of \textit{T. nativa} is occupied by \textit{Vulpes vulpes}. The other Carnivora species are subdominant. The specifics of the trichinellosis transmission and \textit{Trichinella} transmission factors depend on the trophic relationships between animal hosts. The main forms of trophic relationships between carnivores are predation, necrophagy and cannibalism. \textit{Erinaceus concolor} and insects (Carabidae) may also contribute to the preservation and dissemination of \textit{Trichinella}.

Key words: infection prevalence, morphometry, natural ecosystems, population structure, shape index, \textit{Trichinella nativa}, \textit{Vulpes vulpes}, wild carnivores

Introduction

Trichinellosis is a zoonotic disease, i.e. a disease affecting both humans and animals. It is very widespread and presents a serious social and general biology problem (Britov, 1982; Gajadhar & Gamble, 2000; Pozio, 2007). The causative agent of trichinellosis was discovered about 150 years ago, although its pathology in humans had been known long before. In any case, the co-evolution of people and \textit{Trichinella} began when humans started eating animals. In other words, the evolution of trichinellosis as a zoonosis proceeded simultaneously with human evolution (Boev, 1978; Britov, 1982; Zarlen-ga et al., 2006). Trichinellosis is mainly spread via two transmission cycles, natural and synanthropic (Britov, 1975; Boev, 1978; Pozio, 2001; Pozio &

Now, four *Trichinella* species are known in Europe, including *T. nativa* and *T. britovi* characterised by high incidence rates, and *T. spiralis* and *T. pseudospiralis*, which are sporadic (Pozio, 2000; Pozio & Murrell, 2006). *Trichinella spiralis* is spread via the synanthropic cycle largely involving *Sus scrofa domestica* Linnaeus, 1758 (Pozio & Zarlenka, 2005). *Trichinella nativa* is more widespread in the arctic and subarctic regions of Europe, Asia and North America. In Russia, *T. nativa* was registered in the regions with a cold climate, including the central part of European Russia, Siberia, and the Russian Far East (Britov, 1982; Pozio et al., 2001; Romashov et al., 2006; Gajadhar et al., 2009; Bukina & Odeovskaya, 2013; Odeovskaya et al., 2013; Tulovala, 2013; Solovieva et al., 2017; Andreyanov & Konyaev, 2018). It is also common in the North European countries (Murrell et al., 2000; Pozio, 2001; Oivanen et al., 2002; Airas et al., 2010; Asbak et al., 2010; Isomursu & Kunnasranta, 2011).

*Trichinella britovi* is found in regions with moderate climate, in Eastern and Western Europe, Asia, and North and West Africa (Pérez-Martín et al., 2000; Pozio, 2001; Schynts et al., 2006; Malakauskas et al., 2007; Blaga et al., 2008; Szell et al., 2008; Beck et al., 2009; Blaga et al., 2009; Frey et al., 2009; Gajadhar et al., 2009; Hurniková & Du-binský, 2009). *Trichinella nativa* and *T. britovi* occur mainly under natural conditions and occupy different areas depending on the climate. The January isotherm of -4°C is the southern boundary of *T. nativa* distribution, while the January isotherm of -6°C is the northern boundary of *T. britovi* distribution (Pozio & Murrell, 2006).

A particular role in the transmission of trichinellosis belongs to invertebrates involved in scavenging of carnivore carcasses. They contribute to the infective larvae transmission and act as disseminators of *Trichinella*. Certain species and groups of invertebrates can swallow *Trichinella* larvae and preserve them for a long time (Odeovskaya, 2011; Bukina, 2012c, 2014; Bukina et al., 2012; Bukina & Igitova, 2013; Zimmerman, 1970; Hulebak, 1980; Maroli & Pozio, 2000; Odeovskaya et al., 2013; Riva et al., 2015). Given the considerable proportion of insects in the diet of insectivores and carnivores, the possibility of the *Trichinella* transmission through insects is relatively high. The role played by insects involved in scavenging animal carcasses can be demonstrated by experimental studies. For example, larvae of *Lucilia sericata* Meigen, 1826 and species of *Sarcophaga* Meigen, 1826, and necrophagous larvae of *Dermestes maculatus* De Geer, 1774 are able to swallow *Trichinella* larvae and maintain their viability for several days. This increases the possibility of parasite transmission by arthropods in the wild (Maroli & Pozio, 2000; Riva et al., 2015).

*Trichinella*-infected larvae of insects can be found on the ground and become food for Insectivora species, which are registered as hosts of *Trichinella* too (Pozio & Murrell, 2006). As a result, the number of hosts for this nematode increases. Pozio (2001) described the infection process of herbivorous mammals feeding on pastures where there were carcasses of animals infected with *Trichinella* larvae. A high reproductive ability was also observed in *Trichinella* extracted from insect larvae, when used in the experiments on laboratory mice (Maroli & Pozio, 2000).

Therefore, the possibility of *Trichinella* transmission by insects is relatively high, given that they are an important element of the diet of insectivores and carnivores. Earlier, Zimmerman (1970) described mechanical transmission of *Trichinella* by necrophagous insects and demonstrated that this process could contribute greatly to the epidemiology of trichinellosis. Now, a growing number of studies focuses on the role of these insects as paratenic hosts. This is a major way for the *T. spiralis* transmission, since after rodent control activities, car-
casses of dead rodents remain close by pig farm, and paratenic hosts can easily disseminate *Trichinella* among domestic and wild animals (Pozio, 2007).

Currently, the epidemic and epizootic situation associated with trichinellosis is rather tense. The distribution areas of *Trichinella* tend to grow all over the world (Gajadhar & Gamble, 2000; Pozio, 2007), including Russia (Uspensky, 2004, 2007; Bukina & Odoevskaya, 2013). Trichinellosis is of current concern in European Russia including the Black Soil Region (Moskvitin et al., 2006; Romashov et al., 2006; Vagin et al., 2011). In the Black Soil Region, the first studies on trichinellosis were conducted by Merkushev (1955, 1965) and continued by Romashov (1959), Romashov et al. (1980). They gathered the information regarding the infection spread in wild animals in the region. More recent studies demonstrated that the first cases of endemic human trichinellosis in the Voronezh Region were assigned to ingestion of badger meat, as *Meles meles* Linnaeus, 1758 is a hunting species in this area (Romashov et al., 2003a,b). Special genetic studies conducted earlier determined that one species of *Trichinella, T. nativa*, is found in the Carnivora populations in the Voronezh State Nature Reserve and its adjacent areas (hereinafter – VSNR) (Romashov et al., 2006). Recent molecular genetic tests confirmed that «Voronezh»’s *Trichinella* belongs to *T. nativa* (Odoevskaya & Spiridonov, 2016). This species is characterised by small reproduction potential in females (due to a short uterus), low adaptation to *Rattus, Sus scrofa* Linnaeus, 1758 and high resistance to freezing (Britov, 1982; Kapel, 2000; Odoevskaya et al., 2009).

*Trichinella* has an «economical» (direct) life cycle. A single host organism combines the functions of the definitive and intermediate hosts, i.e. the phenomenon of amphixenia is observed (Sudarikov, 1971; Pozio, 2007). As a result, two *Trichinella* groups are formed in the host at the same time, i.e. imaginal and larval hemipopulations. Within a single host, they are defined as local hemipopulations, which successively replace one another. The first life cycle stage is the formation of an imaginal hemipopulation after the invasive larvae enter the host organism. The second stage is the formation of the larval *Trichinella* hemipopulation by migration of the larvae and their encapsulation into muscles. Fig. 1 shows a scheme of the population structure and the life cycle of *Trichinella*, where all stages of the development of the nematode proceed in the endogenous environment, i.e. in the definitive (= intermediate) host (Fig. 1).

Today, it is important to study the trichinellosis transmission into the natural environment in the forest-steppe conditions of the centre of European Russia. We collected new material on trichinellosis under similar conditions, focusing primarily on the data from the Voronezh State Nature Reserve. In this regard, the obtained results will be of great practical interest. This material will be able to characterise the distribution, transmission process and epizootology of trichinosis in the study area. This will enable us to assess the epidemic risks associated with this infection.

We aimed to study certain aspects of the morphology and biology of *Trichinella*, and the ecology and epizootology of trichinellosis under natural conditions in the Black Soil Region, represented by the VSNR. For this purpose, we performed the following tasks: a) to study the morphological and taxonomic features of the local *Trichinella* species; b) to estimate the *Trichinella* larvae distribution in the host’s muscles and assess the local hemipopulation of *Trichinella*; c) to study the ecological and biological patterns of the trichinellosis transmission and the formation of an ecological model of trichinellosis in the study area.

**Material and Methods**

The material was collected in the VSNR (between 51.083° N and 52.0° N and between 39.033° E and 39.083° E). The Voronezh State Nature Reserve is located in the central part of the Black Soil Region. This is a predominantly forest-steppe area, with vast steppe areas combined with forest outliers. The area of the Voronezh State Nature Reserve is 320 km². The Protected Area is almost completely
covered by forest. It occupies the northern half of the Usman forest, one of the largest forest outliers in the Black Soil Region. The adjacent areas occupy the southern half of the Usman forest (280 km²). The Voronezh State Nature Reserve borders with natural and agricultural areas, as well as two administrative regions (Voronezh Region and Lipetsk Region) and five districts (Verkhnekhavsky district, Ramonsky district, Novousmansky district, Usmansky district, and Voronezh urban area).

During the study, we processed and analysed original material collected over 30 years (1990–2019) in the VSNR from potential *Trichinella* hosts, mainly carnivores, and some Insectivora species (e.g. *Erinaceus concolor* Martin, 1838). More than 200 specimens of wild and domestic carnivores (Canivora) of three families (Canidae, Mustelidae and Felidae), belonging to 12 species, were studied, including *Canis lupus* Linnaeus, 1758 (34 specimens), *Vulpes vulpes* Linnaeus, 1758 (157 specimens), *Nyctereutes procyonoides* Gray, 1834 (ten specimens), *Meles meles* (eight specimens), *Martes martes* Linnaeus, 1758 (14 specimens), *M. foina* Erxleben, 1777 (12 specimens), *Neovison vison* Schreber, 1777 (19 specimens), *Lutra lutra* Linnaeus, 1758 (two specimens), *Mustella eversmanni* Lesson, 1827 (one specimen), *Mustela nivalis* Linnaeus, 1766 (six specimens), *Canis lupus familiaris*, Linnaeus, 1758 (30 specimens), *Felis catus* Linnaeus, 1758 (18 specimens). In the Voronezh State Nature Reserve, the material was obtained from living host animals during control activities, as well as from dead animals. In the adjacent areas, material was obtained from carnivores brought by hunters (in areas where hunting is allowed) and from dead animals (mostly the ones killed by cars in traffic). More than 2000 specimens of *Sus scrofa* were studied on *Trichinella* larvae in the Voronezh State Nature Reserve in the 1980–1990s.

We diagnosed and detected *Trichinella* sp. larvae in the muscle tissue using trichinelloscopy and artificial digestion (Vladimirova, 1965; Gamble et al., 2000). The prepared material (compressorium and watch glasses) were examined using MBS-10 and Motic SMZ161-TLED microscopes at a magnification of 10–45×. The relative size of the local hemipopulation of *Trichinella* (in one host) was calculated by the number of larvae per 1 gram (lpg) of the muscle tissue (relative intensity of infection).

We investigated the distribution of *T. nativa* larvae in various muscle groups of carnivores. These data were obtained in the VSNR from four carnivore species (*Vulpes vulpes, Nyctereutes procyonoides, Martes martes*, and *Felis catus*). We took a 1-g sample from each muscle and counted the number of larvae in them. Forty skeletal muscles combined into six groups were examined using trichinelloscopy.

Based on the relative *Trichinella* number in the muscles of an individual host, we can determine the size of local hemipopulations (Fig. 1). This indicator allows us to calculate the approximate size of the *Trichinella* hemipopulation in the population of a certain host species per limited area. The size of *Trichinella* hemipopulations in the host population can be calculated by multiplying three factors, namely 1) the size of the local larvae hemipopulations, 2) *Trichinella* incidence index, and 3) the number of host individuals (Galaktionov & Dobrovolsky, 1998).

By parasitising on various hosts, *Trichinella* demonstrates morphological variations in larvae (organs and structures) and capsules (shape and size). We studied *Trichinella* larvae from six species of wild and domestic carnivores (*Vulpes vulpes, Canis lupus, Nyctereutes procyonoides, Martes martes, Meles meles*, and *Felis catus*). *Trichinella* capsules have a rounded or slightly elongated shape. The capsule walls are relatively thick with a larva inside staying in a helically coiled form. The capsule shape was evaluated using the shape index, i.e. the ratio of the capsule diametre to its length. A comparative morphological study of capsules and living *Trichinella* larvae was performed. For this purpose, temporary samples were prepared by placing the subjects on glass slides in normal saline. The diameter (D) and the length (L) of the capsules were measured. They were used to determine the shape index (V) of *Trichinella* capsules, calculated as a ratio of diameter to length (V = D/L). This parameter is used in zoological studies to assess the shape of rounded objects, e.g. bird and helminth eggs (Kostin, 1977; Romashov & Lomakin, 2000).

We carried out a comparative survey of *Trichinella* larvae morphometry from the Carnivora species of the study area. The measurement results are given in micrometers (µm). Comparative morphometry of the larvae was based on 11 parameters, including the already known (body length (µm), body width (µm), number of stichocytes (n), length of the muscular esophagus (µm), length of the mid-intestine (µm), length of the rectum (µm), length of gonads (µm) (Sokolova & Shaikenov, 1976), and the new (length of the trophic sensory part (µm), length of the trophic reproductive part (µm), length of the stichosome (µm), distance from the beginning of the gonads to the caudal end (µm)). The results of pairwise comparison of *Trichinella*
morphism were presented using the Student’s t-test. We determined that the host pairs significantly differed in a certain set of parameters (p < 0.050–0.001). Morphological and morphometric studies of Trichinella were carried out using a BIOMED-6 light microscope (40–1000 × magnification).

In addition, we investigated some aspects of the Trichinella dissemination involving invertebrate animals in the Voronezh State Nature Reserve. We disposed of the corpses of Nyctereutes procyonoides infected by T. nativa. The insects died in the Voronezh State Nature Reserve as a result of trichinellosis infestation (more than 700 larvae per 1 g of muscle tissue (lpg)). The experimental conditions were as close as possible to natural ones. We collected insects (imago and larvae) from the Nyctereutes procyonoides corpses and examined them on Trichinella larvae.

In order to assess the infection rate and distribution of Trichinella larvae in the hosts, we used the following indices: the abundance index, the infection intensity index, and the infection prevalence (incidence) index. When considering the structure of the helminth (Trichinella) population and the relationships within it, we took into account the current concepts and used the adopted terminology according to Galaktionov & Dobrovolsky (1998). Statistical data processing was performed using the software Statistica 5.5 (Statsoft, Tulsa, OK, USA) and Microsoft Excel 2013.

Results

In the VSNR, T. nativa was found in nine Carnivora species, including seven wild mammals (Vulpes vulpes, Nyctereutes procyonoides, Canis lupus, Meles meles, Martes martes, M. foina, and Erinaceus concolor) and two domestic carnivores (Felis catus and Canis lupus familiaris). The infection prevalence in wild mammals ranged from 8.3% to 70.0% (Fig. 2). These species have several levels of dominance. Typically, a small number of host species, one or two, play a key role in the Trichinella transmission. They are primarily the most numerous native Carnivora species. The highest infection prevalence was observed in four species (Martes martes, Meles meles, Nyctereutes procyonoides, and Vulpes vulpes). Other host animal species, including both native and alien species, occupy the other levels in the parasitic system of Trichinella, depending on the peculiarities of their ecology. Infection rates were much lower in Canis lupus and Martes foina. We should also note that Trichinella was found in Erinaceus concolor (Fig. 2).

We found a considerable difference between the relative number of Trichinella in the carnivores of the Voronezh State Nature Reserve and those on adjacent areas (Table 1). The number of larvae in Nyctereutes procyonoides was considerably higher than that in Vulpes vulpes, although both host species belong to the same family (Canidae) and have similar ecological preferences. Based on these data, we determine two population size variants depending on the host. In the first variant, the hosts are Vulpes vulpes and Martes martes with the average infection rates within 10 lpg, and in the other variant the hosts are Nyctereutes procyonoides and Felis catus with 600–700 lpg (Table 1).

![Fig. 2. Prevalence of Trichinella nativa infection in wild mammals in the Voronezh State Nature Reserve and its adjacent areas (Russia).](https://dx.doi.org/10.24189/ncr.2021.023)

<table>
<thead>
<tr>
<th>Groups of muscles</th>
<th>N</th>
<th>Average lpg in the muscle (minimum and maximum number)</th>
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</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Vulpes vulpes</td>
</tr>
<tr>
<td>Muscles of the head (I), lpg</td>
<td>6</td>
<td>4.3 (2–7)</td>
</tr>
<tr>
<td>Muscles of the neck (II), lpg</td>
<td>4</td>
<td>6.0 (3–12)</td>
</tr>
<tr>
<td>Muscles of the shoulder and foreleg (III), lpg</td>
<td>6</td>
<td>12.6 (7–19)</td>
</tr>
<tr>
<td>Trunk muscles (IV), lpg</td>
<td>11</td>
<td>11.0 (5–20)</td>
</tr>
<tr>
<td>Muscles of the pelvic arch and hind leg (V), lpg</td>
<td>12</td>
<td>10.9 (3–23)</td>
</tr>
<tr>
<td>Muscles of the tail (VI), lpg</td>
<td>1</td>
<td>1.8 (1–3)</td>
</tr>
<tr>
<td>Average number of larvae in the studied groups of muscles, lpg</td>
<td>40</td>
<td>9.4</td>
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*Note: n – number of examined muscles in each group (I–VI).*
We assume that such a great difference can be explained by the host specificity of *Trichinella* manifested in the host-parasite relationship system. We believe that carnivores are adapted to «local» *Trichinella*. In this regard they have developed congruent host-parasite relationships. On the contrary, predator species (*Nyctereutes procyonoides* and *Felis catus*), alien to the VSNR, do not have the necessary adaptive immune response to the «local» *Trichinella*. In these hosts, a significant increase in the size of local *Trichinella* hemipopulations (intensity of infection) was observed.

The distribution of *T. nativa* larvae in different groups of carnivore muscles is of aggregated nature, considering their absolute and relative numbers. A pairwise comparison of individual muscle groups showed significant (p < 0.05–0.01) differences between these values. The highest relative number was observed in the muscle groups of the forelegs and hind legs (Table 1).

Similar ratio tendencies were observed for the larvae distributed in the muscles of Canidae (*Vulpes vulpes* and *Nyctereutes procyonoides*), Mustelidae (*Martes martes*) and Felidae (*Felis catus*). The ratios of the relative larvae number by muscle groups were calculated for these carnivores. The highest number of *Trichinella* was concentrated in the muscles of the shoulder girdle, foreleg, pelvis, and hind leg, followed by the head, neck, and trunk. The minimum number was registered in the tail muscles (Fig. 3).

The *Trichinella* circulation is accounted by the diversity and the number of host animals. In the VSNR, the number of carnivores determines the size of larval *Trichinella* hemipopulations. On the other hand, estimating the size of larval *Trichinella* hemipopulations in carnivore populations makes it possible to consider the role of certain carnivore species in the transmission of natural foci of *T. nativa* on a case-by-case basis.

We performed a special measurement of the local *Trichinella* hemipopulation in *Vulpes vulpes*. The average relative number of *Trichinella* in *V. vulpes* muscles was 35.5 lpg. The average muscle mass of the sample was 2500 g. By multiplying these values (35.5 × 2500 = 88 750), we determined the size of the local *Trichinella* hemipopulation in *V. vulpes* (about 88 750 larvae). Then, we calculated the size of *Trichinella* hemipopulations in the *V. vulpes* population in the whole Voronezh State Nature Reserve. In this Protected Area, the *V. vulpes* population currently consists of 104 individuals (according to the 2015–2019 data), while the *Trichinella* infection rate is 35.7% (index of prevalence is 0.357). By multiplying these values, the number of individuals per local hemipopulation (88 750 × 104 × 0.357 = 3 295 110), we obtained the size of *Trichinella* larvae hemipopulation in the *V. vulpes* population in the Voronezh State Nature Reserve. Such studies have not yet been conducted for other carnivores.

Therefore, based on the size of the local *Trichinella* hemipopulation, the incidence in host populations and their number, we can assess the role of a certain carnivore species in the trichinellosis transmission. *Vulpes vulpes* is the main driver in the *Trichinella* preservation and maintaining the stable trichinellosis transmission in the VSNR. First, *V. vulpes* demonstrated a relatively high prevalence of infection with *Trichinella* larvae (35.7%). Second, *V. vulpes* is the most numerous carnivore species. Its abundance is currently several times higher than the abundance of other carnivores. The role of other Carnivora species in this process is low, taking into account the parameters of the infection rate (Fig. 2) and their number (according to the 2015–2019 data).

The data on the form index of *Trichinella* capsules in the studied Carnivora species are listed in Table 2. The largest capsules were detected in *Felis catus* and *Nyctereutes procyonoides*. In other studied hosts, the capsules were considerably smaller (Table 2). Based on the data obtained, the capsule shape index was calculated. It varied from 0.63 to 0.97. Rounded capsules were observed in *Felis catus*, *Nyctereutes procyonoides*, and *Vulpes vulpes*. Elongated capsules were registered in *Martes martes*, *Canis lupus*, and *Meles meles* (Fig. 4).
The pairwise comparison of the capsules from different hosts demonstrated that in most of the compared pairs of carnivore species, the shape indices differed significantly (p < 0.001). The only exceptions were the pairs Meles meles – Canis lupus and Vulpes vulpes – Nyctereutes procyonoides. For them, no significant differences in the shape index (p > 0.05) were observed (Table 3). These results indirectly indicate close trophic relationships in these pairs. Therefore, we assume that Nyctereutes procyonoides is being infected with Trichinella via Vulpes vulpes, while Canis lupus via Meles meles. The comparative morphometry of Trichinella larvae by 11 parameters from the studied mammal species is presented in Table 4.

Significant differences were revealed among the larvae of the studied mammal species in pairwise comparison. The maximum differences (by 11 parameters) were recorded in Trichinella nativa larvae in the host pair Vulpes vulpes – Felis catus. Three pairs of hosts differ in 10 parameters: Martes martes – Felis catus, Nyctereutes procyonoides – Felis catus, and Canis lupus – Felis catus. Four pairs of carnivores differ in 9 parameters: Vulpes vulpes – Martes martes, Meles meles – Felis catus, Meles meles – Vulpes vulpes, and Martes martes – Nyctereutes procyonoides. There are pairs that differ in 8, 7, and 6 parameters: Canis lupus – Martes martes, Meles meles – Nyctereutes procyonoides, and Meles meles – Canis lupus, respectively. A difference in five parameters was revealed in two pairs: Vulpes vulpes – Nyctereutes procyonoides and Vulpes vulpes – Canis lupus. A difference in four parameters was observed in Meles meles – Martes martes and in two parameters in Canis lupus – Nyctereutes procyonoides.

We believe that the degree of variability of a particular parameter depends on the host species. The most significant difference, both in the number of parameters and in the variation degree, was noted in the pair Felis catus – Nyctereutes procyonoides. Thus, we can say that T. nativa is more adapted to the native host species (Vulpes vulpes, Canis lupus, Meles meles, and Martes martes) than to the alien species (Nyctereutes procyonoides and Felis catus). Therefore, Trichinella larvae should exhibit higher pathogenicity for alien host species, Nyctereutes procyonoides and Felis catus.

The complex joining method was used to analyse T. nativa larvae by 12 parameters (including the capsule shape index). The analysis identified six Trichinella morphs representing the individual characteristics for each carnivore host species, and two relatively independent clusters of host eco-forms. The first cluster included two Mustelidae species (Martes martes and Meles meles) and one Canidae species (Canis lupus). The second cluster was formed by two Canidae species (Vulpes vulpes and Nyctereutes procyonoides) and one Felidae species (Felis catus) (Fig. 5).
Table 3. The results of pairwise comparison of *Trichinella nativa* capsules by the shape index from six carnivore species in the Voronezh State Nature Reserve and its adjacent areas

<table>
<thead>
<tr>
<th>Host types and species</th>
<th>Host 1 V (M ± m (lim))</th>
<th>Host 2</th>
<th>Criteria for the pairwise comparison by the shape index</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Martes foina</em></td>
<td>0.70 ± 0.02 (0.54–0.87)</td>
<td></td>
<td>-</td>
</tr>
<tr>
<td><em>Meles foina</em></td>
<td>0.63 ± 0.01 (0.56–0.74)</td>
<td></td>
<td>-</td>
</tr>
<tr>
<td><em>Vulpes foina</em></td>
<td>0.88 ± 0.02 (0.74–1.00)</td>
<td></td>
<td>-</td>
</tr>
<tr>
<td><em>Canis lupus</em></td>
<td>0.64 ± 0.02 (0.42–0.79)</td>
<td></td>
<td>-</td>
</tr>
<tr>
<td><em>Nyctereutes procyonoides</em></td>
<td>0.90 ± 0.02 (0.82–1.00)</td>
<td></td>
<td>-</td>
</tr>
<tr>
<td><em>Felis catus</em></td>
<td>0.97 ± 0.01 (0.86–1.00)</td>
<td></td>
<td>-</td>
</tr>
</tbody>
</table>

Note: V – the shape index of *Trichinella* capsules; M – mean value; m – error of the mean; lim – limits of values; t – Student’s t-test; p – significance level; significantly different pairs are marked in bold.

Table 4. Morphometric study of *Trichinella nativa* larvae from six carnivore species in the Voronezh State Nature Reserve and its adjacent areas

<table>
<thead>
<tr>
<th></th>
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<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Meles meles</td>
<td>967.7 ± 30.3</td>
<td>1124.0 ± 22.6</td>
<td>1164.9 ± 32.4</td>
<td>982.1 ± 27.9</td>
<td>1137.0 ± 19.3</td>
<td>889.6 ± 5.98</td>
</tr>
<tr>
<td>Vulpes foina</td>
<td>38.8 ± 0.87</td>
<td>41.9 ± 0.95</td>
<td>41.5 ± 0.8</td>
<td>39.3 ± 0.4</td>
<td>43.0 ± 0.69</td>
<td>44.5 ± 0.27</td>
</tr>
<tr>
<td>Canis lupus</td>
<td>636.6 ± 22.7</td>
<td>735.5 ± 17.7</td>
<td>808.4 ± 23.1</td>
<td>662.4 ± 20.1</td>
<td>775.1 ± 16.5</td>
<td>592.2 ± 3.56</td>
</tr>
<tr>
<td>Nyctereutes procyonoides</td>
<td>339.6 ± 14.5</td>
<td>388.0 ± 17.7</td>
<td>362.5 ± 20.0</td>
<td>319.7 ± 16.0</td>
<td>361.6 ± 9.1</td>
<td>297.4 ± 7.79</td>
</tr>
<tr>
<td>Felis catus</td>
<td>479.2 ± 21.3</td>
<td>535.3 ± 15.0</td>
<td>648.2 ± 25.5</td>
<td>553.0 ± 19.2</td>
<td>624.9 ± 16.5</td>
<td>481.6 ± 3.6</td>
</tr>
<tr>
<td>Meles meles</td>
<td>49 ± 1</td>
<td>48 ± 1</td>
<td>51 ± 1</td>
<td>51 ± 1</td>
<td>51 ± 1&lt;br&gt;(50–52)</td>
<td>51 ± 1&lt;br&gt;(50–52)</td>
</tr>
<tr>
<td>Vulpes foina</td>
<td>153.4 ± 4.53</td>
<td>204.0 ± 8.93</td>
<td>160.2 ± 9.2</td>
<td>109.4 ± 6.4</td>
<td>150.2 ± 7.9</td>
<td>110.5 ± 1.9</td>
</tr>
<tr>
<td>Canis lupus</td>
<td>326.3 ± 16.7</td>
<td>369.3 ± 17.5</td>
<td>340.7 ± 19.0</td>
<td>294.2 ± 15.8</td>
<td>339.5 ± 9.1</td>
<td>265.8 ± 7.7</td>
</tr>
<tr>
<td>Nyctereutes procyonoides</td>
<td>20.8 ± 0.55</td>
<td>18.7 ± 17.2</td>
<td>21.8 ± 1.5</td>
<td>25.5 ± 2.4</td>
<td>22.1 ± 1.6</td>
<td>31.6 ± 0.91</td>
</tr>
<tr>
<td>Felis catus</td>
<td>316.5 ± 14.3</td>
<td>365.7 ± 17.6</td>
<td>331.5 ± 20.2</td>
<td>294.8 ± 16.0</td>
<td>312.1 ± 8.9</td>
<td>259.4 ± 7.8</td>
</tr>
<tr>
<td>Meles meles</td>
<td>21.8 ± 1.85</td>
<td>22.3 ± 1.62</td>
<td>31.1 ± 1.8</td>
<td>24.9 ± 1.3</td>
<td>49.5 ± 1.9</td>
<td>38.1 ± 0.92</td>
</tr>
</tbody>
</table>

Note: n – number of the studied larvae from carnivore hosts; M – mean value; m – error of the mean; lim – limits of values. Studied parameters: 1 – body length, μm; 2 – body width, μm; 3 – length of the trophic sensory part, μm; 4 – length of the trophic reproductive part, μm; 5 – length of the stichosome, μm; 6 – number of stichocytes, n; 7 – length of the muscular esophagus, μm; 8 – length of the mid-intestine, μm; 9 – length of the rectum, μm; 10 – length of gonads, μm; 11 – distance from the gonads’ beginning to caudal end of larvae, μm.
The similarity of *T. nativa* larvae morphs indicates a stable and successful infection of predators in each cluster by trichinellosis. This indicates trophic relationships between host animals (predation, necrophagia, cannibalism) participating in the *Trichinella* transmission in the VSNR. We determined two groups of trophically similar carnivore species, which exchange *Trichinella* the most actively: *Meles meles* – *Canis lupus* and *Vulpes vulpes* – *Felis catus domestica* (Fig. 5). *Vulpes vulpes* is also more likely to contribute to the sustainable infestation of *Nyctereutes procyonoides* with *T. nativa* than other hosts.

Thus, there is much carrion, i.e. dead carnivores, on the diet of *Nyctereutes procyonoides* in the Voronezh State Nature Reserve (Ivanova, 1962). The results of the clustering of the hostal forms of *Trichinella* reflect the specifics of trophic relationships between individual groups of predators, including the native and alien species) (Fig. 5).

We have confirmed the participation of insects in the *Trichinella* larvae transmission. Predatory beetles, namely Carabidae species, were identified as disseminators of larvae. *Trichinella* larvae were found in 28% of the studied Carabidae beetles. A negative result was obtained for other insects, namely necrophagous beetles and Calliphoridae larvae. The number of invertebrates increased manifold on the decomposing animal body. This is an important trophic factor in the trichinellosis dissemination. Consequently, *Trichinella* larvae can be transmitted to carnivores and insectivores (i.e. *Erinaceus concolor*) from Carabidae species. Ivanova (1962) studied the eating habits of *Nyctereutes procyonoides*, *Vulpes vulpes*, and *Meles meles* in the Voronezh State Nature Reserve. The obtained results confirm that in natural environment in the trophic chain «mammals ↔ insects», *Trichinella* is transmitted by insects.

Below, we present the ecological model of the *T. nativa* transmission in the VSNR. In the diagram, the host animals are arranged according to their contribution to the preservation of trichinellosis, taking into account the factors of the trichinellosis dissemination (trophic chorological interactions and the number of host animals) (Fig. 6).

*Vulpes vulpes* is the dominant host which forms the core of the *T. nativa* parasitic system. This is indicated by the high infection rate and the size of the local *Trichinella* hemipopulation, as well as by the highest number of *Vulpes vulpes* compared to other carnivores. The next level is formed by *Nyctereutes procyonoides*, *Meles meles*, and *Martes martes*. The remaining two carnivore species (*Canis lupus* and *Martes foina*) are on the periphery (Fig. 6).

On the one hand, the mechanism of the trichinellosis circulation is based on the interactions between the parasite and host populations, and, on the other hand, on the inter-population and intrapopulation relationships of the hosts, which occur among their food chains. In the VSNR, the main ecological forms of these relationships and *Trichinella* transmission routes are predation, necrophagy and cannibalism. In addition, *Erinaceus concolor* and insects (Carabidae) can serve as an important additional source of *Trichinella* infection (Fig. 6).
Discussion

The obtained results characterise the features of the biology and ecology of *Trichinella* in the VSNR. *Trichinella nativa* was found in mammals, primarily carnivores. This corresponds to the current geographical distribution of this species (Britov, 1982; Pozio & Murrell, 2006; Gajadhar et al., 2009; Bukina & Odoevskaya, 2013; Odoevskaya & Spiridonov, 2016). The study area is located in the centre of European Russia (Black Soil Region) in the forest-steppe zone (Milkov, 1961). Therefore, the obtained results concerning the biology and ecology of *T. nativa* can be extrapolated to the natural conditions of this whole region.

*Trichinella nativa* is found under natural conditions where carnivores are obligatory (principal) hosts. In the VSNR, *T. nativa* is found in populations of eight carnivore species. The highest infection prevalence (35.7–70.0%) was observed in *Vulpes vulpes*, *Nyctereutes procyonoides*, *Meles meles*, and *Martes martes*.

Our study of over 2000 *Sus scrofa* specimens from the Voronezh State Nature Reserve for *T. nativa* gave negative results. The material was obtained during the population control activities in 1980–1990s. Therefore, *S. scrofa* is not infected with *T. nativa* and does not transmit this parasite in the Voronezh State Nature Reserve. A number of studies indicate the *T. nativa* phenotypic variations caused by parasitising in non-specific hosts. These variations include morphological changes, lower virulence, fecundity, and resistance and shorter time of parasitism (Odoevskaya et al., 2009; Bukina, 2013).

We examined the relative number (lpg) and distribution of *T. nativa* larvae in various muscles of carnivores. Consequently, we determined two patterns for the relative number of larvae. In the native carnivore species (*Vulpes vulpes* and *Martes martes*), the number of larvae was about 10 lpg, while in alien carnivores (*Nyctereutes procyonoides* and *Canis lupus familiaris*) there were more than 700 lpg. This leads to a higher pathogenicity of trichinellosis. The obtained results allow us to consider *Trichinella* as a considerable factor in controlling the number of hosts that affects the naturalisation of alien carnivore species in natural conditions.

The results of similar studies revealed similar rates for the relative intensity of *T. nativa* infection in muscle tissue. In the arctic regions, *Trichinella* was found in *Vulpes vulpes* (48.5 lpg), *Vulpes lagopus* (11.8 lpg), and *Canis lupus familiaris* (90 lpg) (Bukina, 2012b). A considerably lower number of larvae (1.6 lpg) was found in *Ursus maritimus* Phipps, 1774 (Moorhead et al., 1999). In marine mammals, the number of larvae varied from 0.2 lpg to more 40 lpg (Proulx et al., 2002; Isomursu & Kunnasrantta, 2011; Bukina, 2012a).

An overview of the existing literature showed that similar numbers were registered for predators in European countries. On average, the level of 2–8 lpg is common for Canidae (*Vulpes vulpes* and *Canis lupus*) (Krois et al., 2005; Malakauskas et al., 2007; Beck et al., 2009), although in some cases infection rates of 50–200 lpg were registered (Krois et al., 2005; Beck et al., 2009).

Material from carnivores demonstrated an uneven distribution of *Trichinella* in the muscles and a relatively pronounced «selectivity» in regard to certain muscle groups. Our study showed that the highest relative number of *Trichinella* (lpg) was observed in the muscle groups of foreleg and hind leg. According to other authors, the muscles of the foreleg and hind leg, masseter, muscles of the tongue and diaphragm are the most actively infested by larvae (Kapel et al., 1995; Belozerov & Zhdanova, 2000; Kapel, 2000; Zhdanova & Uspensky, 2002; Bukina et al., 2008; Gajadhar & Forbes, 2010). Kapel (2000) noted that the parasite distribution in the host’s muscles does not depend on the genotype, but it is rather determined by the host type, the period and intensity of infestation by adult *Trichinella*.

During the comparative morphological study, we noted the host-depending variability of *T. nativa* capsules and larvae. The highest level of variability in *Trichinella* larvae was found between groups, on the one hand, from local host species, on the other, from alien host species. According to the results of larvae morphs’ clustering, trophically close predator groups were identified. Similar studies were conducted in other regions and also demonstrated the adaptive variability of the *Trichinella* larvae (Britov, 1982; Bukina et al., 2013). A considerable difference was observed in *Trichinella* between certain carnivore groups, for example, between Canidae and Felidae (Odoevskaya et al., 2009).

Invertebrates can actively contribute to the trichinellosis transmission under natural conditions. Our field experiment, conducted in the Voronezh State Nature Reserve, demonstrated that predatory Carabidae species can disseminate *T. nativa* larvae. For other insects (necrophagous beetles and Calliphoridae larvae), we obtained negative results. We suggest that further studies are necessary to assess the role of insects as paratenic hosts of *Trichinella* in the Voronezh State Nature Reserve.

Ecological diversity in certain regions is a determining factor for the species diversity of definitive (intermediate) hosts of *Trichinella*. Our study
showed that in the VSNR, the *Trichinella nativa* transmission is being performed by the carnivore communities represented by six species. The key role belongs to *Vulpes vulpes* as the dominant host. This demonstrates high rates of the *Trichinella* infection prevalence and a high index of abundance. In addition, *Vulpes vulpes* has the largest population in the carnivore community of the VSNR.

Similar results were obtained by other researchers. *Vulpes vulpes* is the main carrier and disseminator of trichinellosis in the central part of European Russia (Moskvitin et al., 2006; Vagin et al., 2011; Andreyanov, 2012). In Western and Eastern Europe, where *Vulpes vulpes* also has the largest population among carnivores, it is the dominant host for *Trichinella britovi*, too (Poilio, 1991; Krois et al., 2005; Szell et al., 2008; Blaga et al., 2009; Frey et al., 2009; Hurníková & Dubinský, 2009; Oivanen & Oksanen, 2009).

Poizio & Murrell (2006) suggested that carnivores should be considered as indicators of trichinellosis-contaminated areas. Earlier, Romashov et al. (2006) suggested *Vulpes vulpes* as a biological object for the trichinellosis monitoring in the Central Black Soil Region. Currently, among wild mammals, *Vulpes vulpes* is the most widespread and numerous hunting prey in this region. We used *V. vulpes* as an indicator species when we monitored the trichinellosis in the study area.

A high infestation potential of *Trichinella* (*T. nativa* and *T. britovi*) is formed in the wild carnivore community under natural conditions in Europe. As a result, we observe the creation of conditions for transferring causative agents of trichinellosis from natural to anthropogenic conditions. There are two cycles of trichinellosis transmission, including natural and synanthropic cycles (Britov, 1975; Boev, 1978; Poilio, 2001; Poilio & Zarlanga, 2005; Poilio & Murrell, 2006). *Trichinella spiralis* is being transmitted steadily through the synanthropic cycle involving *S. scrofa domesticus* (Rosenthal, 2008). However, some authors (Poilio & Zarlanga, 2005; Zarlanga et al., 2006) believe that *T. spiralis* survives to a certain extent due to the natural cycle involving *Sus scrofa* and carnivores.

*Trichinella nativa* and *T. britovi* are being transmitted through the natural cycle involving mainly wild carnivores (Poilio, 2001; Poilio & Murrell, 2006). The mechanism of the trichinellosis transmission is based on the interactions between the parasite and host populations, and the interpopulation and intrapopulation relationships of the hosts, which occur among their food chains. In the wild, *Trichinella* is transmitted via nutritional chains of animal hosts, including predation, necrophagy and cannibalism.

**Conclusions**

We found that *Trichinella nativa*, registered in the VSNR, circulates as a natural focal infection. Its obligatory hosts are represented by six species of wild carnivores (*Vulpes vulpes*, *Nyctereutes procyonoides*, *Canis lupus*, *Meles meles*, *Martes martes*, and *M. foina*). *Trichinella* was also found in *Eriaceus concolor* and two domestic carnivores (*Felis catus* and *Canis lupus familiaris*). The highest infection prevalence was observed in *Vulpes vulpes*, *Nyctereutes procyonoides*, *Meles meles*, and *Martes martes* (35.7–70.0%). These hosts play a leading role in the natural trichinellosis transmission.

The carnivores can be divided into two groups according to the infection intensity by *Trichinella* larvae in the muscles (lpg). The first group included native species (*Vulpes vulpes*, *Canis lupus*, *Meles meles*, *Martes martes*, and *M. foina*), with an average of 10 lpg. The second group included alien wild (*Nyctereutes procyonoides*) and domestic (*Felis catus*) species, with an average of 700 lpg. The lpg of native carnivores is considerably lower than that of alien species. Due to its high infestation rate, *Trichinella* can be considered as a considerable factor in the regulation of the number of alien carnivores. In the Voronezh State Nature Reserve, *Trichinella* contributes highly to the *Nyctereutes procyonoides* naturalisation. Therefore, we can say that under natural conditions, *T. nativa* protects the ecosystem from alien species invasion.

*Trichinella* is characterised by an aggregated distribution in the muscles. In the Voronezh State Nature Reserve, carnivores had the highest lpg values in the muscles of the front and rear limbs with over 50% of the local hemipopulation of *Trichinella*. We therefore recommend taking samples from the muscles of the front and rear limbs to examine carnivores for trichinellosis.

Evaluation of the morphological characteristics of *Trichinella* capsules and larvae from carnivores allowed us to identify two clusters of hosts. These clusters comprised carnivore species with closer trophic connections. The variability of the quantitative and qualitative parameters of *Trichinella* indirectly reflects the specificity of its relations in the parasite-host system by showing the food preferences of the studied carnivores in the Voronezh State Nature Reserve.

Under natural conditions, invertebrates can also transmit trichinellosis. In the Voronezh State Nature Reserve, Carabidae beetles disseminate *T. nativa*. *Trichinella* can also be transmitted from insects to mammals.

In the VSNR, *Vulpes vulpes* is the main driver of the stable trichinellosis transmission. *Vulpes vulpes*
showed high rates of infection prevalence (35.7%) and large *Trichinella* larvae hemipopulations (3.3 million specimens). *Vulpes vulpes* is the most abundant species among the carnivores in the Voronezh State Nature Reserve. In addition, this carnivore is the most common hunting prey and can serve as an object for the trichinellosis monitoring in European Russia.

At present, in the VSNR, trichinellosis is being transmitted within the community of wild carnivores. The dominant position in the parasitic system of *T. nativa* is occupied by *Vulpes vulpes*, while the other species of carnivorous are subdominant. *Erinaceus concolor* and Carabidae species can also disseminate *Trichinella*.

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ЭКОЛОГИЧЕСКИЕ ОСОБЕННОСТИ ЦИРКУЛЯЦИИ ТРИХИНЕЛЛЕЗА В ВОРОНЕЖСКОМ ЗАПОВЕДНИКЕ И НА СМЕЖНЫХ ТЕРРИТОРИЯХ (РОССИЯ)

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В статье рассмотрены отдельные аспекты морфологии и биологии трихинелл, экологии и эпизоотологии трихинеллеза в Воронежском заповеднике и на смежных территориях (Центро-Черноземный регион России). Оригинальные материалы собраны в течение 30 лет (1990–2019 гг.) от потенциальных хозяев трихинелл, преимущественно хищных млекопитающих. За указанный период исследовано свыше 200 особей диких и домашних хищников (Carnivora) трех семейств (Canidae, Mustelidae, Felidae). Обитателями хищников являются шесть видов диких хищных млекопитающих: Canis lupus, Vulpes vulpes, Nyctereutes procyonoides, Meles meles, Martes martes, Martes foina. Также трихинеллы обнаружены у Erinaceus concolor и двух видов домашних хищников: Canis lupus familiaris, Felis catus. Наиболее высокая экстенсивность инвазии отмечена у Vulpes vulpes, Nyctereutes procyonoides, Meles meles, Martes martes (35.7–70.0%). Эти хозяева играют ведущую роль в циркуляции природного трихинеллеза. Среди хищников выделены две группы по интенсивности инфекции личинок трихинелл в мышцах (число личинок на 1 г – lpg). В первую группу входят местные виды хищников (Vulpes vulpes, Canis lupus, Meles meles, Martes martes, M. foina), выявлено в среднем 10 lpg. Во вторую группу входят виды интродуценты (Nyctereutes procyonoides) и виды вселенцы (иммигранты) из антропогенных территорий (Felis catus), у них выявлено в среднем 700 lpg. При высокой патогенности трихинелл мы рассматриваем этих паразитов как весомый фактор регуляции численности чужеродных видов хищников. Для трихинелл отмечено агрегированное распределение в мышцах. В Воронежском заповеднике у хищных выявлены самые высокие показатели lpg в мышцах передней и задней конечностей. Здесь сосредоточено свыше 50% численности локальной гемипопуляции трихинелл. По оценке морфологических признаков капсул и личинок трихинелл от хищников выделены два кластера хозяев T. nativa. В этих кластерах объединены виды хищников, которые имеют более близкие трофические связи. Гостальная изменчивость количественных и качественных признаков косвенно отражает специфику отношений в системе паразит-хозяин и показывает пищевые предпочтения хищников. В Воронежском заповеднике и на смежных территориях Vulpes vulpes является основным звеном в устойчивой циркуляции природных очагов трихинеллеза. У лисицы выявлены высокие показатели экстенсивности инвазии (35.7%) и большая численность гемипопуляции личинок трихинелл (3 300 000 экз.). Vulpes vulpes – самый многочисленный вид среди хищников в Воронежском заповеднике и наиболее распространенный объект охоты на сопредельных территориях. Vulpes vulpes может служить объектом мониторинга трихинеллеза в европейской части России. В Воронежском заповеднике циркуляцию трихинеллеза могут поддерживать беспозвоночные животные, в качестве диссеминаторов T. nativa зарегистрированы жуки-карабиды (Carabidae). В настоящее время в Воронежском заповеднике и на смежных территориях трихинеллез циркулирует в сообществе диких хищников. Доминирующее положение в паразитарной системе трихинелл занимает лисица, субдоминантами являются остальные виды хищников. Особенно циркуляции трихинеллеза и факторы передачи трихинелл обусловлены трофическими связями животных-хищников. Основными формами трофических связей среди хищников являются хищничество, некрозфагия и каннибализм. Erinaceus concolor и насекомые (Carabidae) выявлены в качестве дополнительных источников в резервировании и диссеминации трихинелл.

Ключевые слова: Trichinella nativa, Vulpes vulpes, естественная экосистема, индекс формы, морфометрия, популяционная структура, хищные млекопитающие, экстенсивность инвазии