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The Taiga Tick *Ixodes Persulcatus* (Acari: Ixodidae), the Main Vector of *Borrelia Burgdorferi Sensu Lato* in Eurasia

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ABSTRACT

The taiga tick *Ixodes persulcatus* Schulze, 1930 is considered to be one of the most important vectors of human pathogens. The tick is a temporary ectoparasite characterized by exophilic (pasture) type of host-seeking behavior and prolonged feeding (blood-sucking). The tick is a typical three-host species, i.e., all three parasitic stages of the tick feed on different hosts. Its geographic range extends as a broad belt from the Baltic Sea to the Pacific Ocean mainly through the zone of boreal coniferous (taiga) forests. The essential role of the taiga tick in transmission of the Lyme disease pathogen Borrelia burgdorferi s.l. was recognized in the 1980s and the following years have been characterized by efforts to clarify the importance of this species in epidemic and epizootic components of the infection. The chapter consists of two parts. The first part is dedicated to the general characteristics of this tick species, its systematic position, geographic range and biological and ecological patterns, such as seasonal activity, questing behavior, hosts, specificity of blood feeding etc. Nearly all of these data were obtained in the 1950-1980s by Russian scientists. The data accumulated during the three last decades of research into the relationships between B. burgdorferi and I. persulcatus are reviewed and discussed in the second part of this chapter. The chapter concludes with a brief discussion of directions and challenges for future research in this field.

The taiga tick *lxodes persulcatus* was described by Schulze in 1930 and very soon thereafter this species became well known when its importance as the main vector of tick-borne encephalitis virus was revealed in the late 1930s [1,2]. Intensive and extensive studies of its biological characteristics as well as its role in epidemic and epizootic processes were carried out in the 1950-1980s by Russian scientists. In the 1980s the taiga tick was found to be a vector of human borreliosis (Lyme disease) and later its participation in the transmission of agents of human monocytic ehrlichiosis and human granulocytic anaplasmosis was also revealed [3]. Today *I. persulcatus* is considered to be one of the most important vectors of human pathogens. A special monograph dedicated to all aspects of taxonomy, biology, control and medical importance (concerning at that time only tick-borne encephalitis) of this species was published in the former USSR [4]. Various characteristics of this species were also analyzed in several reviews [3-6].

TAXONOMIC STATUS

The taiga tick is a member of the family Ixodidae Murray, 1877, the subfamily Ixodinae Murray, 1877, the genus *Ixodes* Latreille, 1795 and the subgenus *Ixodes* (s.str.) Latreille, 1795. The subfamily Ixodinae (Prostriata group) is more ancient and more primitive morphologically and ecologically than subfamilies from Metastriata group. The origin of the subfamily Ixodinae is attributed to the Cretaceous while that of the genus *Ixodes* to the Paleogene. The subgenus *Ixodes* is considered to be the most progressive among other subgenera of the genus whereas *Ixodes persulcatus* is evolutionarily one of the youngest species originated in the late Paleogene [4,7].

It was recognized that several *Ixodes* species in the Asian part of the former USSR had been previously erroneously identified as *I. persulcatus* because of their morphological similarity. It was suggested [8] to integrate these species in a *I. persulcatus* group. Later this group was increased to incorporate up to 14 species, including species from Nearctic and Indo-Malayan faunal regions [9,10]. Such important vectors as European *I. ricinus* and North-American *I. scapularis* and *I. pacificus* also belong to this group. The most accepted name of the group is *I. ricinus/I. persulcatus* complex though in accordance with the International Code of Zoological Nomenclature [11] the correct name of the group should be *persulcatus* [12]* (*Nomination of *I. ricinus* as the leading species of the group (complex) seems to be inconsistent with established practice. Besides, this species is the type species of the subgenus *Ixodes* and additional nomination of it as the leading species of the group might create an erroneous impression that all species in the subgenus belong to this group).

GEOGRAPHIC RANGE

The range of the taiga tick extends as a broad belt from the Baltic Sea to the Pacific Ocean mainly through the zone of boreal coniferous (taiga) forests. The northern border of the range is limited by the approximate value of the sum of temperatures for the period with the stable daily temperature above 5°C equal to 1,600°C. This roughly coincides with the southern border

of tundra and semi-tundra forests. Biotopes most suitable for *I. persulcatus* are located in midand southern-taiga forests whereas in the European part of the range such biotopes also exist in deciduous forests. Most often ticks exist in upland moraine and lowland glacial-lake landscapes. In the western part, the northern border of the range stretches along the latitudes of 62-63°N, in the Asian part it gradually moves southward and at the Far East the border lies at the latitudes of $52-54^{\circ}$ N. The southern border of the range is defined by soil humidity [3,4,13].

The main part of the range of *I. persulcatus* lies on the territory of Russia. In the western part of the range the taiga tick presents in the eastern parts of Latvia and Estonia and in some areas of Lithuania; some findings of the species have been reported in Finland as well as in Belorussia, Poland and Ukraine.

Isolated populations of the taiga tick persist in mountain areas of Kazakhstan, Uzbekistan and Kirgizstan [3,4], in Mongolia, in 13 of 28 provinces of China including Tibet [14], apparently in North Korea, in northern part of South Korea [15], in Japan (Hokkaido Island but also mountain areas of Honshū and Kyūshū Islands) [16,17], and in mountain areas of Taiwan [18]. Regular findings of single *I. persulcatus* adults have been documented along the routes of bird migrations, mainly along the valleys of big Siberian rivers, even within the Arctic Circle [19-21].

In recent years, many authors have insisted that the northern border of the taiga tick range moves northwards. Regrettably, these conclusions have been made, with rare exceptions, on the basis of single tick collections on the territories where no previous surveys had been carried out. The question whether such shifts are the result of the global climate warming and have a steady tendency [22-24] or they only reflect regular fluctuations of tick abundance or they are simply the result of surveying the areas, which had not been surveyed previously, [3,25] remains to be open.

BIOLOGICAL CHARACTERISTICS

The taiga tick *Ixodes persulcatus* is a temporary ectoparasite characterized by exophilic (pasture) type of host-seeking behavior and prolonged feeding (blood-sucking). The tick is a typical three-host species, i.e., all three parasitic stages of the tick feed on different hosts [4,7].

Ecological Patterns

As the majority of tick species living in the temperate zone, *I. persulcatus* has a well-defined seasonality. Over the main part of the range, unfed ticks of all parasitic stages demonstrate host-seeking activity during the spring-summer season of the year. The very first adults appear when snow still lies and only thawed patches appear in well-warmed parts of tick biotopes [26]. I was attacked by such ticks on the northern slope of a mountain ridge in the Western Sayan mountains in the end of April when the air temperature was near 0°C. Not all ticks become active at once in the beginning of the activity season but they do it gradually: the main part of adult ticks become active during the first 15-30 days after the beginning of host-seeking activity but the entire period of tick appearance can continue for up to 85 days [27-30]. Apparently the appearance of unfed nymphs and larvae follows a similar pattern although this has never been tested experimentally.

It is difficult if at all possible to collect ticks of these stages by flagging or dragging as adults are collected [4], while collection of preadults on the hosts excludes their repeated findings.

The seasonal host-seeking activity of unfed adult ticks continues for 60 to 120 days but sometimes even shorter or longer depending on the meteorological conditions of a particular year [4,31,32]. In a typical case, activity accelerates rapidly to a peak in April-May with a gradual decline until the end of host-seeking activity later in the summer or even in the autumn. Sometimes people can be attacked by an adult female much later than the last tick was found by flagging. The pattern of tick activity strongly depends on meteorological conditions, so that its dynamics in a particular year can be completely unpredictable. The dynamics of host-seeking activity of nymphal and larval ticks follows, in general, that of adult ticks with some increases in preadult occurrence during the period of activity and perhaps a longer duration of activity by nymphs [4].

The activity period does not correspond to the life span of individual ticks. After hatching or molting, unfed preadults live about 14-15 months and unfed adults live about 12 months including post-molting development and diapause during hibernation when ticks do not seek hosts (see below). The life of host-seeking (questing) ticks is much shorter, from 1 day to about 75 days for unfed adults [27,33,34]. The adult ticks that become active earlier in the season live longer than those that become active later [29]; the latter begin their active life being physiologically older than the ticks which begin their activity earlier [33].

When seeking hosts, adult ticks use the so-called "ambush strategy": unfed ticks are positioned on the vegetation scanning nearby spaces with their chemoreceptors, which are parts of Haller's organs located on the dorsal surface of tarsus I [35,36]. They spend from several hours to several days in the questing position depending on the particular meteorological conditions. Adult ticks can sense the host stimuli from a distance of 5 to 10 m in still weather [37], and as far as 15 to 20 m from the windward direction under windy conditions [38]. They can also detect paths of regular host migrations, which explains their high concentration nearby. Questing ticks either catch onto a passing host or, sensing it from a distance, move in its direction. To maintain their water balance, ticks must regularly migrate down to the litter and soil where the temperature is lower and humidity higher. Unfed adult ticks have two peaks of activity during the day: in the morning (8-10 a.m.) and in the late afternoon (after 4 p.m.) [39,40]. Nymphal activity is maximal at dusk and the early hours of the night [41]. Cloudy weather, rain or high temperatures at nights can significantly alter this pattern. However, if a host appears close to an unfed tick, the latter becomes active and can attack the host at any time.

The capacity of *I. persulcatus* for active migrations is limited. The questing larvae usually can move up to 20 cm, nymphs up to 40-50 cm, and adults up to 1.0 m. The height of vertical movement is directly correlated with the air humidity being maximal in humid taiga forests of the Russian Far East [7,42]. Ticks also are unable to carry out long horizontal migrations; adults can migrate up to 10 m and larvae up to 1.5 m [43]. The rate of adult movement may be as high as 30 cm/min at 23°C but usually decreases with lower temperatures [42]. The taiga tick has no eyes though

it can distinguish between more and less illuminated areas, preferring the latter. Photosensitive cells were found in approximately the same area where eyes could be located [44]. The response of these photosensitive cells to light appears to be comparable to that in some species of ticks that have eyes [45].

Unfed host-seeking adults are characterized by a 1:1 sex ratio [4,30,46-48]. Only about 2% of active females are successful in finding hosts and engorgement [49-51]. The fraction of unfed larvae and nymphs that find hosts and engorge is higher because of greater number of host species and their higher abundance. It is estimated that more than 90% of larvae and more than 20 % of nymphs are successful in attaching to hosts and engorgement [49,50]. The fate of engorged ticks depends on the time of feeding. If the engorgement takes place in the first half of the activity season, larvae and nymphs molt into the next stage while females oviposit. Unfed nymphs and adults after ecdysis are able to enter behavioral diapause and overwinter. Unfed larvae emerging from eggs laid are also able to enter behavioral diapause. If the engorgement takes place in the second half of the activity season, engorged larvae and nymphs enter morphogenetic diapause and overwinter. Nymphal ticks start entering morphogenetic diapause in June whereas larval ticks do it in July [52,53]. In rare instances, engorged nymphs can survive for two winters being in diapause. Diapause is induced by declining photoperiod with an earlier beginning in more continental climate [4,54-56]. Eggs laid by engorged females are unable to overwinter and die. Thus, during the activity season there are simultaneously unfed ticks of several cohorts: those that were in behavioral and/or morphogenetic diapause and overwintered from the previous year and those that hatched (larvae) or emerged from engorged larvae or nymphs in the same year. Accordingly, adult ticks may be represented by specimens of up to 5 different cohorts.

The duration of life cycle (from eggs laid by an engorged female until the unfed adult ticks) can reach 3, 4 or 5 years (rarely 6 years) depending on the number of diapauses in a particular tick population. The maximal duration of life cycle has been observed in Siberian region and the western part of Far East while in the southern Far East or in the European part of the range this period is much shorter (sometimes only 2 years) [4,53,56,57]. The ratio between the parts of tick population developing with different duration depends on climatic conditions of particular years.

Hosts and Feeding Specificity

The taiga tick is one of the most opportunistic (generalist) tick species, being able to use nearly 300 species of different vertebrates (about 100 species of mammals, more than 175 species of birds and a few species of reptiles) as hosts [4]. As a rule, each stage has hosts of preferential sizes: larvae feed on small mammals such as mice, voles and shrews; nymphs feed on small and medium-sized mammals such as chipmunks, squirrels, hedgehogs as well as on ground-dwelling birds; adults use as hosts medium-sized and large animals such as hares, deer, elk but also cattle and dogs. The questing height of ticks of each stage coincides with the size and location of their main hosts. Although larvae and nymphs can use the same hosts, nymphs demonstrate clear preference for medium-sized mammals and birds [49]. Apparently, the taiga tick employs

the strategy of partitioning resources (a case of classical niche partitioning [58,59]) in order to diminish interstage competition. All stages of *I. persulcatus* are active at the same time, so they should diversify their hosts (partitioning in space). In two other closely related vectors from the same group, *I. ricinus* and *I. scapularis*, the time of host-seeking activity of adults does not usually coincide with that of nymphal ticks, i.e. these species utilize partitioning in time. Perhaps, this phenomenon explains why only adult *I. persulcatus* attack humans (large animals) being more aggressive than adult *I. ricinus* [60] (nymphal *I. persulcatus* are also more aggressive towards their hosts than *I. ricinus* nymphs [61]). In this respect, *I. persulcatus* differ from *I. ricinus* and *I. scapularis*, which seek hosts at the time of maximal human outdoor activity, are the main attackers and vectors of human pathogens. Preadult ticks can switch from their typical hosts for unusual ones under specific conditions such as depression in the abundance of small mammals [62,63] or on animal farms with a dominant host species [64].

Each parasitic stage feeds on a vertebrate host only once: the larva feeds for 3-5 days, the nymph for 3-6 days and an adult female for 6-10 days. The duration of feeding and the weight of engorged specimens strongly depend on the host species [65]. Males feed several times during their active life for 15-30 min each time. The obligatory condition for normal female engorgement is its insemination. The taiga tick females can be inseminated on the hosts as well as in nature, before finding a host. In the middle of the activity season up to 50% of unfed females collected from vegetation were inseminated [47,66,67]. Adult ticks produce an assembly pheromone which increases the probability of mating off hosts [68].

Larval and nymphal ticks increase their weight during feeding 15- to 30-fold, while engorged females are heavier than unfed ones 100- to 150-fold. The mass of an unfed larva is 0.03-0.045 mg, whereas the mass of fully engorged females reaches 250-470 mg [69]. Fully engorged females lay from 2,000 to 4,000 eggs. The minimal weight of engorgement, after which females are capable of laying single eggs, is about 20-35 mg. The heavier the engorged female becomes, the greater the ratio of the number of laid eggs/mg of the female's weight. When the female weight reaches 140-200 mg, the ratio reaches a plateau at approximately 8.5-9.5 eggs/mg [69]. The values of all of these parameters may differ between tick populations in different parts of the taiga tick range and strongly depend on the host species.

THE TAIGA TICK AS VECTOR OF *BORRELIA BURGDORFERI* S.L., THE CAUSATIVE AGENT OF TICK BORRELIOSIS

Lyme disease was identified and described in the USA where the spirochete, *Borrelia burgdorferi* s.str., is the only causative agent of the disease. In Europe and Asia there is a set of closely related infections caused by several genospecies of *B. burgdorferi* s.l. At present, Lyme disease is the most common tick-borne zoonosis in temperate regions of the Northern Hemisphere. The pathogen is transmitted to humans by several species of hard ticks from the genus *Ixodes*, the taiga tick *I. persulcatus* being the main vector through its entire range in Eurasia [3,70].

B. burgdorferi in infected ticks

The vector competence of *Ixodes persulcatus* for the causative agent of the disease was recognized in Russia in the middle of the 1980s [71]. Two genospecies of *Borrelia burgdorferi* s.l., which are absent in the USA, *B. garinii* and *B. afzelii*, are the most common pathogens of *I. persulcatus*, while the American genospecies, *B. burgdorferi* s.str., occurs only infrequently. *B. garinii* and *B. afzelii* are widespread over the entire range of the taiga tick [15,72,73]. The prevalence of ticks infected with *B. burgdorferi* s.l. varies by geographic areas and particular years being usually rather high.

The *Borrelia* spirochetes inhabit the gut of unfed ticks. A subset of unfed infected ticks demonstrates systemic (generalized) infection when *B. burgdorferi* penetrates into the body cavity, gonads and salivary glands. Such ticks are able to infect hosts soon after attachment [74]. It was shown that the number of *I. persulcatus* females with systemic infection positively correlates with the spirochete concentration in the tick gut [75]. The intensity of infection in physiologically older ticks is lower compared to physiologically younger specimens [76]. Intensive migration of spirochetes from the gut into salivary glands is initiated after the tick has been attached to the host for 2.5 days [3].

There is a very significant difference between *I. persulcatus* and the main vector of this pathogen in the USA and Canada, *I. scapularis*, with regard to the prevalence of ticks with systemic infection. The proportion of unfed *I. scapularis* with systemic infection was found to be much lower than that of *I. persulcatus* (2.4% and 12.9%, respectively) [77]. The small numbers of *I. scapularis* with systemic infection lead to a conclusion that attachment by this tick for less than 48 hours is not sufficient for human infection with *B. burgdorferi* [78,79]. In contrast, the taiga tick is able to infect humans within the first 24 hours after attachment [80]. The proportion of unfed *I. ricinus* with systemic infection is lower than that of *I. persulcatus* but higher than in *I. scapularis*, which leads to a higher rate of host infection during the first 24 hours after tick attachment [3,81,82]. However, even a small number of unfed ticks with systemic infection can be a source of human infection within a very short time after tick attachment. In some studies, the samples were too small to detect the ticks with systemic infection. It appears that host infection during the very first hours after tick attachment has not been studied [83].

B. burgdorferi spirochetes infect a number of eggs laid by engorged females but only a negligibly small number of larval ticks become infected [84,85]. (Some authors observed nearly 100% transovarial transmission by *I. persulcatus* [86], but this finding is likely attributable to methodological deficiencies). It is generally accepted that there is negligible if any transovarial transmission of *B. burgdorferi* in the main vectors from the genus *Ixodes*. Ticks mainly acquire the spirochetes during their larval or nymphal feeding on small mammals that are reservoir hosts for *B. burgdorferi*. The importance of reservoir hosts for infecting ticks is very high since they are able to preserve the spirochetes for a very long, perhaps lifelong, periods [3,87]. Larvae

and nymphs effectively transmit spirochetes to the next developmental stages (transstadial transmission) [88,89]. The infection rate of each subsequent stage with *B. burgdorferi* is higher than that of the preceding stage. An alternative route of tick infection by spirochete is the so-called co-feeding (distant) transmission, when uninfected larvae receive pathogen from infected nymphs feeding simultaneously on the same host [90,91]. Such transmission in *I. persulcatus* was found in only a single study with this species [92], while in *I. ricinus* it was demonstrated multiple times [93,94]. The significance of this route of infection is presently under discussion [87,95]. Sexual transmission by spirochetes between copulating adult partners [96] might be also of some, though probably minor, importance.

It was found that *I. persulcatus* is a more effective vector than *I. ricinus* not only because of its greater aggressiveness towards humans [60,61] but also because of a greater proportion of unfed specimens with systemic infection (see above) [3,70]. The suggestion of the positive influence of *I. persulcatus* on the infection rate of *I. ricinus* in their sympatry zones [97] was found to be incorrect [87,98,99].

Influence of B. burgdorferi Infection on the Tick Hosts

It was postulated [100] that although the relationships of *Borrelia* with ixodid ticks are evolutionary younger and more primitive than its relationships with argasid ticks, the symbiotic interrelations *Borrelia*/ixodid ticks are well balanced. At the same time, it is widely accepted that microbial pathogens harbored by the tick vectors are parasites and as such are expected to have some negative effect on the ticks [101-103]. Only a small number of studies were carried out to elucidate any evidence of parasitic influence of *Borrelia burgdorferi* on the tick hosts.

In one study [104] no differences in tick development were found between *I. persulcatus* specimens infected and uninfected with *B. burgdorferi*. One study [105] found lower motor activity in infected ticks as compared to uninfected ones, although a more comprehensive study [106] of infected *I. persulcatus* demonstrated a greater questing activity than that of uninfected ticks, and this activity was positively correlated with the intensity of tick infection. It was also reported that there is a trend towards an increase in the longevity of infected adult *I. persulcatus* (and of *I. ricinus* too) [34]. The infection rate of adult ticks that were able to climb up the experimental rod to its maximal height of 30 cm was twice as high as that of the ticks that could not reach this height [107]. The above abnormalities in the behavior of infected ticks can be considered an example of the "parasite manipulation of the host" [108].

Coinfection of B. burgdorferi-infected ticks with other pathogens

Until recently, the pathogen-vector relationships of the taiga tick *Ixodes persulcatus* were studied only with regard to the tick-borne encephalitis virus (TBEV). The discovery of *B. burgdorferi* and consequent detection of many other pathogens harbored by *I. persulcatus* (as well as by other ticks from *persulcatus* group) was followed by studies of coinfection of *B. burgdorferi*-

infected ticks with other pathogens. The development of molecular techniques for detection of pathogenic DNA and RNA gave an additional impetus to these studies. It was shown that one and the same adult *I. persulcatus* could be simultaneously infected with *B. garinii* of different genome groups (NT29 and 20047^T), with *B. afzelii* and each of the groups of *B. garinii*, and with *B. afzelii* and the both groups of *B. garinii* [3,72,109]. Coinfection of *I. persulcatus* with both genospecies of *B. burgdorferi* was observed in several other studies [110,111]. Coexistence of *B. afzelii* and *B. garinii* with *B. miyamotoi*, a relapsing fever spirochete, which earlier was misidentified with *B. burgdorferi*, has also been reported [112, 113].

The most intriguing question for Russian scientists concerning *I. persulcatus* was whether *B. burgdorferi* can coexist with TBEV, the most intensively studied pathogen during the 1960s-1980s. It was suggested that the relationships between the virus and the spirochete function as a "one-sided antagonism": the presence of spirochetes in tick body limits the virus circulation while the virus does not prevent, and possibly even stimulates, the transmission of spirochetes [105,114]. However, long-term observations carried out in different areas using a variety of techniques decisively demonstrated the absence of such "antagonism" between pathogens, or of any influence of pathogens on each other [87,115,116]. Coinfection of adult *I. persulcatus* by *B. burgdorferi* and TBEV was found in 18% of ticks examined in western Siberia [117].

Babesia microti infection was found in taiga tick adults near St. Petersburg in combination with one or both genospecies of B. burgdorferi or with B. garinii and TBEV (18 combinations in total) [118]. Coexistence of *B. burgdorferi* with rickettsiae of several genera in *I. persulcatus* adults was demonstrated in several studies. Since the prevalence of rickettsial agents in I. persulcatus is much lower than that of *B. burgdorferi*, coinfection has usually been registered only in a few ticks. Six of 1,146 adults and 1 of 199 nymphs (totally 0.5%) collected in northeastern China were infected with *B. burgdorferi* and *Anaplasma phagocytophilum* [119], whereas in western Siberia the prevalence of ticks coinfected by this combination of pathogens reached 6% [117]. Several combinations of 2 to 4 pathogens were revealed in adult ticks in north-western regions of Russia including B. burgdorferi, Ehrlichia muris, Rickettsia tarasevichiae and Rickettsia-like endosymbiont Montezuma [120]. Coexistence of *B. burgdorferi* with the West Nile virus (WNV) or with *R. tarasevichiae* as well as with 2 other pathogens (TBEV+WNV, TBEV+ *R. tarasevichiae*, WNV+ *R. tarasevichiae*) was found in adult taiga ticks near Tomsk (West Siberia) [121]. Moreover, 17 combinations of DNA of different pathogens were revealed in adult taiga ticks in the Cis-Ural region (both genospecies of B. burgdorferi, Ehrlichia muris, Anaplasma phagocitophilum and Rickettsia spp.), and DNA of 4 or 5 agents being identified in 1.3% of ticks (5 out of 400) [122]. In fact, the larger the tick sample and the greater the number of PCR primers, specific for different pathogenic agents, the higher the chance to find multiple pathogens in a single tick.

Ultrastructural analysis confirmed the possibbility of tick coinfection with borreliae (*B. burgdorferi* s.l.), rickettsiae (*E. muris* and *R. tarasevichiae*) and a flavivirus (apparently TBEV), and showed that different microorganisms are harbored by different organs of the same tick [123].

It appears that different pathogenic agents tend to localize to distinct cells, organs or tissues of the tick body thus avoiding antagonistic relationships with each other [3,109,124]. It is clear that the research on pathogen coexistence in ticks is still in its early stages, so it can be expected that the data reviewed here will be further expanded, and the conclusions may be refined or revised.

The phenomenon of coinfection of ticks with multiple pathogens necessitates that human protection be focused on prevention of tick bites rather than on elimination of tick-borne pathogens. The territorial principle of human protection through total tick control, as it was applied in the former Soviet Union [125,126], is unrealistic in the modern era. The modern strategy of human protection relies on individual protection of each person in the area where they can be attacked by infected ticks [127,128]. Protective clothes impregnated with acaricides with low toxicity for mammals were developed and successfully used in different areas of Russia [129].

CONCLUSION

The discovery of Lyme disease and its causative agent, and the identification of ixodid ticks as its vectors ushered a new era in research on ticks and tick-borne diseases. In the three decades that followed, at least 10 microbial pathogens and a number of potential pathogens of humans and animals were identified in ticks of the *persulcatus* group, and in *I. persulcatus* in particular. This wealth of new information poses of multitude of questions concerning the relationships between ticks and pathogens, and especially between ticks and various combinations of different pathogens. In the absence of answers to these questions, it is impossible to reliably estimate the danger presented by particular areas populated by infected ticks (i.e., natural foci of tick-borne diseases), nor it is possible to forecast the rate of tick infection by a particular pathogen and, most importantly, the expected human morbidity. The answers for these questions are reasonably connected with in-depth investigation of ticks and pathogens using new molecular techniques. Likewise, numerous questions arise in connection with the appearance of tick populations at anthropogenically modified territories and in urban environment [130-132]. At the same time, it is important to keep in mind that in spite of decades of intensive research on the taiga tick at the population and organismal levels, some important questions are still awaiting answers. As one example, little is known about the biological mechanisms underlying the differences in aggressiveness of closely related tick species towards humans. Finding the answers to these questions would require the deployment of the whole arsenal of available approaches and methodologies to in-depth analysis of various aspects of tick and pathogen biology. Laboratory experiments and molecular methods are invaluable in addressing these issues; however, a thorough understanding of such highly complex problem cannot be attained without long-term observational studies of tick populations in nature.

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