

'Candidatus Neoehrlichia mikurensis' in Europe

A. Portillo, P. Santibáñez, A. M. Palomar, S. Santibáñez and J. A. Oteo

Center of Rickettsiosis and Arthropod-Borne Diseases, Infectious Diseases Department, Hospital San Pedro—Center of Biomedical Research from La Rioja (CIBIR), La Rioja, Spain

Abstract

'Candidatus Neoehrlichia mikurensis' is an uncultured emerging bacterium that is provisionally included in the family *Anaplasmataceae*. In Europe, it is transmitted by *Ixodes ricinus* ticks. Rodents are the reservoirs. It is widely distributed in mammals (both wild and domestic) and birds. It causes an inflammatory disease in humans with underlying diseases, but the microorganism also affects immunocompetent individuals in which asymptomatic infection has been recognized. A high degree of suspicion and the use of molecular tools are needed for the correct diagnosis. Efforts to cultivate it and to investigate its pathogenesis should be a priority.

© 2018 The Authors. Published by Elsevier Ltd.

Keywords: 'Candidatus Neoehrlichia mikurensis', coinfection, *Dermacentor reticulatus*, doxycycline, Europe, *Ixodes ricinus*, neohrlichiosis, reservoir, tick, tick-borne diseases

Original Submission: 15 November 2017; **Revised Submission:** 14 December 2017; **Accepted:** 20 December 2017

Article published online: 6 January 2018

Corresponding author: J. A. Oteo, Hospital San Pedro—Center of Biomedical Research from La Rioja (CIBIR), Infectious Diseases Department, Center of Rickettsiosis and Arthropod-Borne Diseases, C/ Piqueras, 98, 26006 Logroño, La Rioja, Spain

E-mail: jaoteo@riojasalud.es

microti, *Bartonella henselae*, Eyach virus, tick-borne encephalitis virus and Crimean-Congo haemorrhagic fever virus [2]. Recent reviews about 'Ca. N. mikurensis' have been published [3]; nevertheless, our aim was to compile and update, under a One Health scope, all available information about this microorganism that may be going unnoticed in our environment.

Introduction

'Candidatus Neoehrlichia mikurensis' was first identified as a human pathogen in 2010. This bacterium was molecularly detected (using PCR assays) from the blood of a Swedish patient with chronic lymphocytic leukaemia who developed prolonged fever, an erysipelas-like rash and thromboembolic complications. Subsequently the patient recovered with doxycycline [1]. The finding of 'Ca. N. mikurensis' contributed to increase the spectrum of human diseases transmitted by hard ticks in Europe, adding this microorganism to the list of *Borrelia burgdorferi* sensu lato, *Borrelia miyamotoi*, *Rickettsia* spp. (*R. conorii*, *R. helvetica*, *R. massiliiae*, *R. sibirica* subsp. *sibirica*, *R. sibirica* subsp. *mongolitimonae*, *R. monacensis*, *R. aeschlimannii*, *R. slovaca*, 'Candidatus *R. rioja'* and *R. raoultii*), *Anaplasma phagocytophilum*, *Francisella tularensis*, *Babesia divergens*, *Babesia*

Basic aspects of the bacterium

'Ca. N. mikurensis' was found for the first time in spleen samples from rats on Mikura Island (Japan), as well as in ticks [4]. Genetic variants of *Ehrlichia* (e.g. *Ehrlichia*-like variant Schotii or 'Candidatus *Ehrlichia walkeri*'), previously found in *Ixodes* spp. ticks or in rats, are synonyms of this species [4].

To date, it has not been possible to cultivate this microorganism. Nevertheless, on the basis of electron microscopic observations of the sinusoids of the spleen from infected rats by Kawahara et al. [4], it is assumed that 'Ca. N. mikurensis' are small (0.5–1.5 µm), Gram negative, pleomorphic cocci (belonging to the class α-proteobacteria) that live in an interacting cycle affecting rodents and ticks. 'Ca. N. mikurensis' grow in membrane-bound inclusions within the cytoplasm of endothelial cells, but the specific mechanism of infection remains

unknown. According to 16S rRNA and *groEL* sequences, the phylogenetic analysis revealed a new cluster in the family *Anaplasmataceae*, close to other known tick-borne agents such as *Anaplasma* spp. or *Ehrlichia* spp [4].

Apart from '*Ca. N. mikurensis*', the '*Candidatus Neoehrlichia*' genus includes at least four other members: '*Candidatus Neoehrlichia lotoris*' (detected in raccoons from North America) [5]; '*Candidatus Neoehrlichia*' sp. FU98 (closely related to '*Ca. N. lotoris*' and found in red foxes, a badger and one *Ixodes rugicollis* tick from Europe) [6,7]; and '*Candidatus Neoehrlichia australis*' and '*Candidatus Neoehrlichia arcane*', both described in *Ixodes holocyclus* ticks from Australia [8]. None of them, except '*Ca. N. mikurensis*', has been involved as a human pathogen.

Epidemiology

The main vector of '*Ca. N. mikurensis*' is *Ixodes ricinus*, a tick species that frequently bites humans in Europe. This tick is distributed throughout Europe and is involved in the transmission of a high number of human infectious diseases (Lyme borreliosis, human anaplasmosis, rickettsiosis, babesiosis, tick-borne encephalitis, *B. miyamotoi* infection and neoehrlichiosis) [2]. To date, '*Ca. N. mikurensis*' has been detected in *I. ricinus* ticks from 20 European countries (Table I). It has been found in *I. ricinus* from the vegetation in 18 European countries, with a prevalence of infection that ranges from 0.1% to 24.3% (Denmark and Hungary, respectively) and sometimes

TABLE I. Detection of *Candidatus Neoehrlichia mikurensis* in ticks from different sources in European countries

Country	Hard tick species (source)	Prevalence (%)	Sampling years	Coinfection	Reference
Austria	<i>Ixodes ricinus</i> (vegetation)	4.2–22.1	2002–13	<i>Babesia</i> spp., <i>Borrelia afzelii</i> , <i>Borrelia burgdorferi</i> , <i>Babesia divergens</i> , <i>Babesia venatorum</i> , <i>Rickettsia</i> spp., <i>Rickettsia helvetica</i> , <i>Rickettsia raoultii</i>	[9–11]
Belgium	<i>I. ricinus</i> (vegetation)	0.4	2012–14		[12]
	<i>I. ricinus</i> (hedgehogs)	2.7	2014–15		[13]
	<i>Ixodes hexagonus</i> (hedgehogs)	0.09	2014–15		[13]
Czech Republic	<i>I. ricinus</i> (vegetation)	0.4–10	2010–14		[10,14–16]
	<i>I. ricinus</i> (sheep)	30.7	2013–14		[16]
Denmark	<i>I. ricinus</i> (vegetation)	0.1–0.9	2008–12		[17,18]
Estonia	<i>I. ricinus</i> (vegetation)	1.3	2006–13		[19]
France	<i>I. ricinus</i> (vegetation)	0.2–1.7	2008–12		[14,18]
Germany	<i>I. ricinus</i> (vegetation)	2.2–24.2	2008–13	<i>Babesia</i> spp., <i>Anaplasma phagocytophilum</i> , <i>R. helvetica</i>	[14,20,21]
	<i>I. ricinus</i> (rodents)	3.8–6.4	2010–13		[20,21]
	<i>I. ricinus</i> (dogs)	4.1–4.3	2010–11		[22,23]
	<i>I. ricinus</i> (humans)	8.1	NA		[14]
	<i>I. ricinus</i> (wild boar)	6.25	2010–13		[24]
	<i>Dermacentor reticulatus</i> (vegetation)	0.08	2010–11		[22]
	<i>D. reticulatus</i> (rodents)	7.7	2010–11		[20]
	<i>Ixodes trianguliceps</i> (rodents)	2.5	2012–13		[21]
	<i>Ixodes</i> spp. (rodents)	100	2010–11		[20]
	Unidentified larva (rodents)	100	2010–11		[20]
	<i>I. hexagonus</i> (dogs)	5.9–6.6	2010–11		[22,23]
Hungary	<i>I. ricinus</i> (vegetation)	8.8–24.3	2007–12		[25,26]
Italy	<i>I. ricinus</i> (vegetation)	10.5	2006–8		[27]
	<i>I. ricinus</i> (rodents)	5.3	2011–13		[28]
	<i>I. ricinus</i> (humans)	0.5	1995–2011		[29]
Moldova	<i>I. ricinus</i> (vegetation)	0.8	1960		[30]
Norway	<i>I. ricinus</i> (vegetation)	5.9	1998–99	<i>B. afzelii</i> , <i>B. burgdorferi</i> sensu stricto	[31]
Poland	<i>I. ricinus</i> (vegetation)	0.3	2011		[32]
	<i>I. ricinus</i> (dogs)	8.1	2013–14		[33]
	<i>I. hexagonus</i> (dogs)	0.7	2013–14		[33]
Romania	<i>I. ricinus</i> (vegetation)	5.3–14.6	2013–14	<i>Borrelia</i> spp., <i>Rickettsia</i> spp.	[34,35]
	<i>I. ricinus</i> (humans)	100	2013	<i>B. afzelii</i>	[36]
Russia-Baltic region	<i>I. ricinus</i> (birds)	0.7	2009		[37]
	<i>Ixodes frontalis</i> (birds)	25	2009		[37]
Serbia	<i>I. ricinus</i> (vegetation)	4.2	NA		[38]
Slovakia	<i>I. ricinus</i> (vegetation)	1.1–11.6	2006–13	<i>A. phagocytophilum</i>	[10,39–42]
	<i>I. ricinus</i> (rodents)	0.3–1.3	2011–14	<i>Babesia microti</i>	[41,42]
	<i>I. trianguliceps</i> (rodents)	2.7	2011–13		[41]
Spain	<i>I. ricinus</i> (cows)	1	2013		[43]
Sweden	<i>I. ricinus</i> (vegetation)	6	2010–11	<i>B. afzelii</i>	[44]
	<i>I. ricinus</i> (birds)	2.1	2009		[45]
Switzerland	<i>I. ricinus</i> (vegetation)	3.5–8	2009–10	<i>B. afzelii</i>	[46,47]
	<i>I. ricinus</i> (rodents)	2.6	2011–12		[48]
	<i>I. ricinus</i> (birds)	3.3	2007–10		[49]
Netherlands	<i>I. ricinus</i> (vegetation)	2.4–11.7	2000–12	<i>B. afzelii</i> , <i>R. helvetica</i>	[18,50,51]
	<i>I. ricinus</i> (humans)	5.4	2007–8		[52]
Netherlands and Belgium	<i>I. ricinus</i> (vegetation)	7	2009–10		[53]
	<i>I. ricinus</i> (birds)	4.9	2012–14	<i>B. burgdorferi</i> sensu lato, <i>Borrelia miyamotoi</i> , <i>R. helvetica</i>	[54]
	<i>I. ricinus</i> (red deer)	6.3	2009–10		[53]
	<i>I. ricinus</i> (European mouflon)	4.3	2009–10		[53]
	<i>I. ricinus</i> (wild boar)	8.3	2009–10		[53]
	<i>I. ricinus</i> (sheep)	12.5	2009–10		[53]

NA, not available.

coinfecting with other tick-borne agents (Table 1). '*Ca. N. mikurensis*' has also been reported in *I. ricinus* collected on a variety of wild and domestic vertebrates, including rodents (0.3–6.4%), dogs (4.1–8.1%), birds (0.7–4.9%), hedgehogs (2.7%), sheep (12.5–30.7%), wild boar (6.25–8.3%), cattle (1%), red deer (6.3%) and European mouflon (4.3%) (Table 1). Moreover, studies about the presence of '*Ca. N. mikurensis*' in *I. ricinus* that had fed on humans have been reported from Germany, Italy, Romania and the Netherlands (Table 1). '*Ca. N. mikurensis*' has also been detected in other *Ixodes* spp., such as *I. hexagonus*, *I. trianguliceps* or *I. frontalis*, from various sources (vegetation, rodents, dogs, hedgehogs and birds) (Table 1). In addition, it is worth mentioning that '*Ca. N. mikurensis*' has been found in *Dermacentor reticulatus*, a tick species that frequently bites humans (Table 1). Transovarial transmission of '*Ca. N. mikurensis*' in ticks does not seem to occur [53].

The role of rodents as reservoir hosts of '*Ca. N. mikurensis*' was suggested as a result of the higher prevalence rates of the bacterium found in ticks from rodents than in questing ticks [20,21]. Xenodiagnoses on rodents confirmed their capacity to transmit '*Ca. N. mikurensis*' to ticks, and therefore their role as

reservoir [48]. Transplacental transmission in rodents was also observed [21]. In Europe, '*Ca. N. mikurensis*' has been detected in at least seven rodent species belonging to the genera *Apodemus* (*A. flavicollis*, *A. agrarius*, *A. sylvaticus*), *Myodes* (*M. glareolus*) and *Microtus* (*M. arvalis*, *M. agrestis*, *M. minutus*) [20–22,42,55]. It has been also detected in other mammals such as wild boar, bears, badgers, chamois and mouflons [56], hedgehogs [57] and dogs [58–61] (Fig. 1). To our knowledge, despite having been investigated, '*Ca. N. mikurensis*' has never been found in cats, wild cervids, martens, jackals or foxes [22,56,60,61], perhaps as a result of low or intermittent bacteraemia or the low tropism of the microorganism in tested tissues.

Neohrlichiosis/'*Ca. N. mikurensis*' infection in humans

Clinical manifestations

Since the first European case of '*Ca. N. mikurensis*' infection in a Swedish patient with a chronic lymphocytic leukaemia with prolonged fever, erysipelas-like rash and thromboembolic

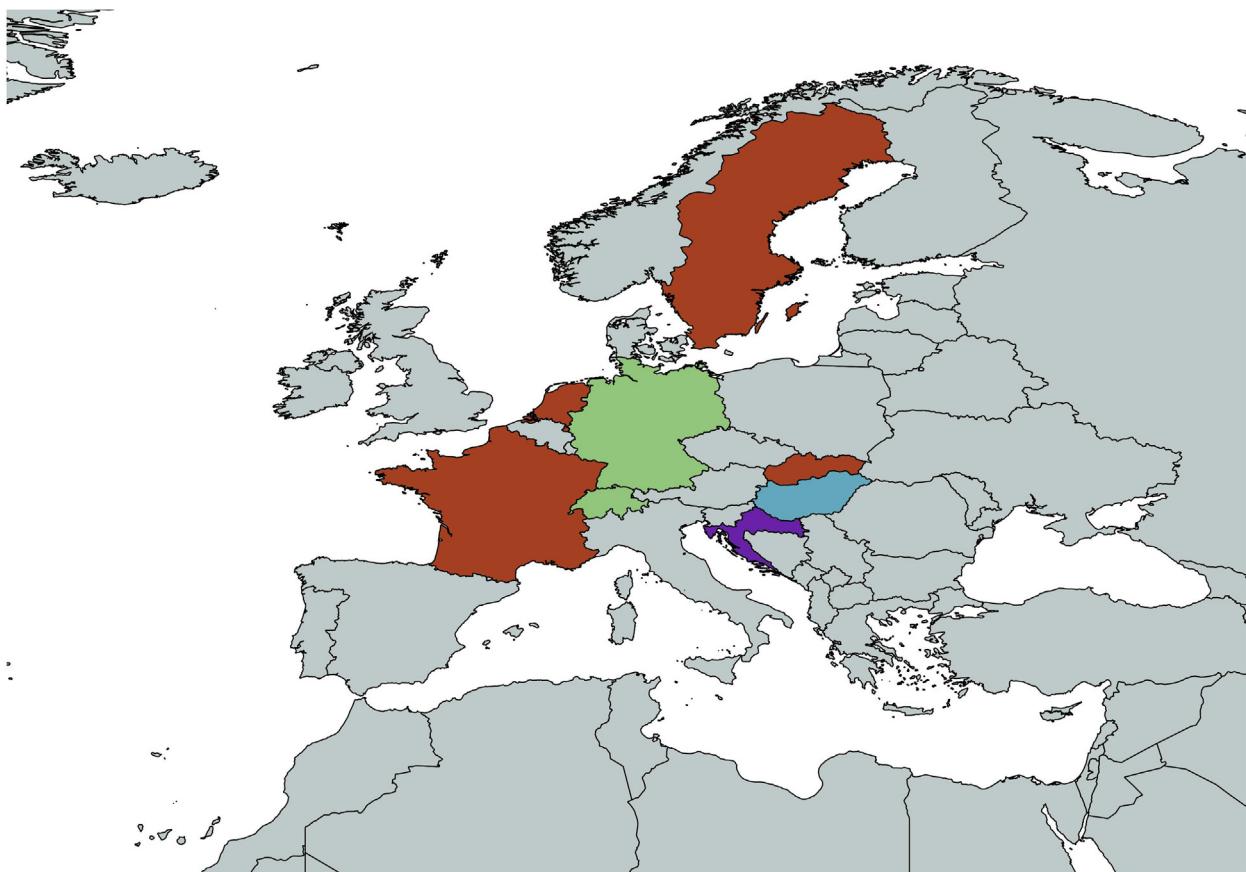


FIG. 1. European mammals in which '*Candidatus Neoehrlichia mikurensis*' has been molecularly detected. Red, rodents; green, rodents and dogs; blue, rodents and hedgehogs; purple, rodents, dogs, wild boars, bears, badgers, chamois and mouflons.

TABLE 2. Clinical background of 18 patients with neoehrlichiosis reported in European countries

Country	Year	Gender	Age	Medical condition	Reference
Czech Republic	2008	F	55	Mantle cell lymphoma, asplenic	[62]
	2009	M	58	Liver transplantation, sclerosing cholangitis, splenectomy	[62]
Germany	2007	M	69	Chronic inflammatory demyelinating polyneuropathy	[63]
	2008	M	57	Previously healthy	[63]
Sweden	2009	M	77	B cell chronic lymphocytic leukaemia, asplenic	[1]
	2011	M	75	B cell chronic lymphocytic leukaemia, splenectomy	[64]
	2011	F	67	Follicular lymphoma, systemic lupus erythematosus, (inborn) asplenic	[64]
	2013	F	67	T cell large granular lymphoma, psoriasis arthropathy, splenectomy	[64]
	2013	M	54	Psoriasis, immunosuppressive therapy	[64]
	2013	M	59	Diffuse large cell B cell lymphoma, rheumatoid arthritis splenectomy	[64]
	2014	F	71	Rheumatoid arthritis, recurrent fever, immunosuppressive therapy	[65]
	2015	M	78	Rheumatoid arthritis	[66]
	2015	M	55	Granulomatosis with polyangiitis	[66]
	2015	M	57	Pre-B cell acute lymphocytic leukaemia	[66]
Switzerland	N.A.	F	65	Autoimmune haemolytic anaemia	[67]
	2009	M	61	Coronary artery bypass grafting, septicaemia	[68]
	2011	M	68	Chronic lymphocytic leukaemia, asplenic	[47]
	2012	M	58	Follicular lymphoma	[47]

F, female; M, male; N.A., Not available.

complications that recovered with doxycycline [1], a total of 18 cases have been reported from European countries, more often in patients with immunocompromised conditions associated with hematologic neoplasias (Table 2). All patients were older than 54 years (mean, 64 years). Seventy-two percent were men and 28% were women. Immunosuppressive conditions were present for 88.9% (16/18), and 50% (9/18) had hematologic neoplasia. Half (9/18) were splenectomized or asplenic, and 78% (14/18) had received immunosuppressive therapy (e.g. 8/14 with rituximab) for the treatment of lymphomas or rheumatic diseases. Fever was present in the 14 cases with available data, and nine of them experienced myalgia and arthralgia. Vascular events such as deep vein thrombosis, thromboembolic events, aneurysm and transitory ischemic accidents were reported for 56% of patients (10/18). At least 22% of the patients (4/18) had skin manifestations, such as erythema nodosum or erysipelas-like rashes. A tick bite was reported by 39% (7/18) patients. Nevertheless, in our opinion, only data from four patients were valid because clinical manifestations had begun between 6 and 16 weeks after the tick bites. Although some

patients had recalled previous tick bites, who was going to think about a tick in a patient who had developed deep thrombosis and fever of unknown origin 2 years after a tick bite? When available (14/18 patients), C-reactive protein levels were elevated in all patients (100%), and most ($n = 10$) had leukocytosis (neutrophilia) and anaemia. Whenever it was analysed, the delay in diagnosis resulted in very high values. Thus, for 11 studied patients, the median number of days from the onset of symptoms to diagnosis was 60 [64].

Moreover, 'Ca. N. mikurensis' has been found in the blood of asymptomatic people from Poland. The study was carried out in healthy foresters bitten by ticks, and 'Ca. N. mikurensis' was amplified from blood in five (1.6%) of 316 subjects [32]. With the purpose of studying possible coinfections in immunocompetent patients with tick-borne diseases, some recent studies have investigated the presence of 'Ca. N. mikurensis' in patients with erythema migrans (Table 3). Thus, in Sweden, the bacterium was detected in two (1.9%) of 102 immunocompetent people bitten by ticks. Both had an erythematous rash; one was infected with *B. burgdorferi*, and the other one seroconverted to

TABLE 3. Clinical picture of European patients with 'Candidatus Neoehrlichia mikurensis' infection, with or without coinfecting pathogens, after receipt of tick bite

Country, no. infections/no. patients (prevalence)	Year	Gender	Age (years)	Medical condition (no. patients)	Reference
Poland, 5/316 (1.6%)	2012	4 M, 1 F	44.1 (mean)	Asymptomatic, previously healthy (foresters with high risk of tick bites)	[32]
Norway, 7/70 (10%)	2014–15	1.6F:1M (ratio)	55 (mean)	EM (7), fatigue (1)	[69]
Sweden, 2/102 (1.9%)	2015	F	68	EM, tick bite, <i>Borrelia</i> -specific IgM and IgG antibodies	[70]
Netherlands, 7/626 (1.1%)	2007–8	F	57	EM, tick bite, <i>Anaplasma phagocytophilum</i> -specific IgG antibodies	[52]
		M	63	Tick bite, arthralgia	
		M	79	Tick bite	
		M	40	EM	
		F	60	EM	
		F	61	EM, headache, myalgia, pain in limbs	
		M	48	EM, tingling in limbs	
		M	71	Tick bite	

EM, erythema migrans; M, male; F, female.

A. phagocytophilum [70]. In the Netherlands, '*Ca. N. mikurensis*' was detected in EDTA blood samples from four (1.4%) of 291 patients with erythema migrans. Two of them did not report any additional symptom, whereas the remaining two reported pain or tingling in the limbs, in one case associated with headache and myalgia [52]. The same authors also found '*Ca. N. mikurensis*' prevalences of 0.9% (3/335) in humans bitten by ticks and higher than 5% (17/314) in ticks removed from participants [52]. In Norway, '*Ca. N. mikurensis*' was found in seven (10%) of 70 blood specimens (whole blood and/or plasma/pellet fraction) from recently tick-bitten persons with erythema migrans. The authors assumed that all of them were immunocompetent, and only one patient showed fatigue as a symptom [69]. It seems that coinfections with *B. burgdorferi* s.l. and '*Ca. N. mikurensis*' do not alter the clinical picture of Lyme disease patients, although more studies are necessary.

Diagnosis

To date, the diagnosis of '*Ca. N. mikurensis*' is based on molecular techniques, specifically PCR assays and sequence analysis of 16S rRNA (pan-bacterial, *Anaplastmataceae* specific and/or '*Ca. N. mikurensis*' specific) and *groEL* genes [1,47,62,63,65,68,70]. Positive PCR results for '*Ca. N. mikurensis*' were obtained using EDTA blood (plasma, serum, whole blood, and blood culture flask contents) and bone marrow [47,64–66] as human specimens. The microorganism has never been amplified from cerebrospinal fluid samples [64].

Serologic tests are not good tools. No cross reactions with other *Anaplastmataceae* (*A. phagocytophilum* or *Ehrlichia chaffeensis*) have been reported. The absence of cross reactions due to the immunosuppressive condition of patients or due to the low level of bacteraemia in immunocompetent people is plausible [63,69]. Only one immunocompetent patient seroconverted to *A. phagocytophilum*, but the tick was not studied [70]. No serologic assay to detect specific antibodies against '*Ca. N. mikurensis*' is available to date.

Other molecular techniques such as multilocus sequence analysis and PCR–reverse line blot hybridization have occasionally been used for epidemiologic studies in Europe [11,66]. Multilocus sequence analysis was applied to investigate the genetic diversity of '*Ca. N. mikurensis*' DNA obtained from immunocompromised patients. The data were based on the study of five housekeeping genes in addition to the 16S rRNA gene, although the technique was limited by the amount of available DNA [66]. Moreover, PCR–reverse line blot hybridization allowed the detection of simultaneous bacteria in *I. ricinus* ticks from Austria, suggesting a significant co-occurrence of '*Ca. N. mikurensis*' and *Babesia* spp. [11].

Treatment

Doxycycline (100 mg twice a day) has been used in nearly all published cases (Table 2). One patient with a suspected allergy to doxycycline was successfully treated with rifampin (300 mg twice a day) [71]. A combination of rifampin (450 mg twice a day) plus doxycycline (100 mg twice a day) was also prescribed in another case [68]. The optimal duration of treatment still remains unknown, but in most cases it lasted for 3 weeks [62,63]. The mean time to resolution of symptoms was 5 days. All patients recovered, and PCR analysis after treatment yielded negative results for '*Ca. N. mikurensis*'.

Future key questions

Studies about the distribution of '*Ca. N. mikurensis*' in nature, as well as reports about the clinical manifestations, diagnosis and treatment of patients infected with this bacterium, have increased in recent years. Nevertheless, there are still several key questions about this pathogen that need to be answered: What are the human target cells of '*Ca. N. mikurensis*'? It seems that leukocytes and endothelial cells are the human targets. Structures with coccoid forms were observed in circulating leukocytes from patients [62]. In addition, apart from humans, is '*Ca. N. mikurensis*' pathogenic for other mammals, such as dogs? Is there a possibility of outbreaks by '*Ca. N. mikurensis*'? And finally, how will climate change affect the distribution of this tick-borne disease?

Conflict of interest

None declared.

References

- [1] Welinder-Olsson C, Kjellin E, Vaht K, Jacobsson S, Wenneras C. First case of human '*Candidatus Neoehrlichia mikurensis*' infection in a febrile patient with chronic lymphocytic leukemia. *J Clin Microbiol* 2010;48:1956–9.
- [2] García-Álvarez L, Palomar AM, Oteo JA. Prevention and prophylaxis of tick-bites and tick-borne related diseases. *Am J Infect Dis* 2013;9: 104–16.
- [3] Silaghi C, Beck R, Oteo JA, Pfeffer M, Sprong H. Neoehrlichiosis: an emerging tick-borne zoonosis caused by '*Candidatus Neoehrlichia mikurensis*'. *Exp Appl Acarol* 2016;68:279–97.
- [4] Kawahara M, Rikihisa Y, Isogai E, Takahashi M, Misumi H, Suto C, et al. Ultrastructure and phylogenetic analysis of '*Candidatus Neoehrlichia mikurensis*' in the family *Anaplastmataceae*, isolated from wild rats and found in *Ixodes ovatus* ticks. *Int J Syst Evol Microbiol* 2004;54:1837–43.
- [5] Yabsley MJ, Murphy SM, Luttrell MP, Wilcox BR, Howerth EW, Munderloh UG. Characterization of '*Candidatus Neoehrlichia lotoris*'

- (family Anaplasmataceae) from raccoons (*Procyon lotor*). Int J Syst Evol Microbiol 2008;58:2794–8.
- [6] Hodžík A, Cézanne R, Duscher GG, Harl J, Glawischnig W, Fuehrer HP. 'Candidatus Neoehrlichia sp.' in an Austrian fox is distinct from 'Candidatus Neoehrlichia mikurensis,' but closer related to 'Candidatus Neoehrlichia lotoris'. Parasit Vectors 2015;8:539.
- [7] Hornok S, Trauttwein K, Takács N, Hodžík A, Duscher GG, Kontschán J. Molecular analysis of *Ixodes ruficollis*, 'Candidatus Neoehrlichia' sp. (FU98) and a novel *Babesia* genotype from a European badger (*Meles meles*). Ticks Tick Borne Dis 2017;8:41–4.
- [8] Gofton AW, Doggett S, Ratchford A, Ryan U, Irwin P. Phylogenetic characterisation of two novel Anaplasmataceae from Australian *Ixodes holocyclus* ticks: 'Candidatus Neoehrlichia australis' and 'Candidatus Neoehrlichia arcana'. Int J Syst Evol Microbiol 2016;66:4256–61.
- [9] Glatz M, Müllegger RR, Maurer F, Fingerle V, Achermann Y, Wilske B, et al. Detection of 'Candidatus Neoehrlichia mikurensis,' *Borrelia burgdorferi* sensu lato genospecies and *Anaplasma phagocytophilum* in a tick population from Austria. Ticks Tick Borne Dis 2014;5:139–44.
- [10] Derdáková M, Václav R, Pangrácová-Blaňárová L, Selyemová D, Kočí J, Walder G, et al. 'Candidatus Neoehrlichia mikurensis' and its co-circulation with *Anaplasma phagocytophilum* in *Ixodes ricinus* ticks across ecologically different habitats of Central Europe. Parasit Vectors 2014;7:160.
- [11] Schötta AM, Wijnveld M, Stockinger H, Stanek G. Approaches for reverse line blot-based detection of microbial pathogens in *Ixodes ricinus* ticks collected in Austria and impact of the chosen method. Appl Environ Microbiol 2017;83. e00489–17.
- [12] Heylen D, Fonville M, van Leeuwen AD, Sprong H. Co-infections and transmission dynamics in a tick-borne bacterium community exposed to songbirds. Environ Microbiol 2016;18:988–96.
- [13] Jahfari S, Ruyts SC, Frazer-Mendelewska E, Jaarsma R, Verheyen K, Sprong H. Melting pot of tick-borne zoonoses: the European hedgehog contributes to the maintenance of various tick-borne diseases in natural cycles urban and suburban areas. Parasit Vectors 2017;10:134.
- [14] Richter D, Matuschka FR. 'Candidatus Neoehrlichia mikurensis,' *Anaplasma phagocytophilum*, and Lyme disease spirochetes in questing European vector ticks and in feeding ticks removed from people. J Clin Microbiol 2012;50:943–7.
- [15] Venclíková K, Rudolf I, Mendel J, Betášová L, Hubálek Z. Rickettsiae in questing *Ixodes ricinus* ticks in the Czech Republic. Ticks Tick Borne Dis 2014;5:135–8.
- [16] Venclíková K, Mendel J, Betášová L, Blažejová H, Jedlicková P, Straková P, et al. Neglected tick-borne pathogens in the Czech Republic, 2011–2014. Ticks Tick Borne Dis 2016;7:107–12.
- [17] Fertner ME, Mølbak L, Boye Pihl TP, Fomsgaard A, Bøcker R. First detection of tick-borne 'Candidatus Neoehrlichia mikurensis' in Denmark 2011. Euro Surveill 2012;17. 20096.
- [18] Michelet L, Delanoy S, Devillers E, Umhang G, Aspan A, Juremalm M, et al. High-throughput screening of tick-borne pathogens in Europe. Front Cell Infect Microbiol 2014;4:103.
- [19] Ivanova A, Geller J, Katargina O, Värv K, Lundkvist Å, Golovljova I. Detection of 'Candidatus Neoehrlichia mikurensis' and *Ehrlichia muris* in Estonian ticks. Ticks Tick Borne Dis 2017;8:13–7.
- [20] Silaghi C, Woll D, Mahling M, Pfister K, Pfeffer M. 'Candidatus Neoehrlichia mikurensis' in rodents in an area with sympatric existence of the hard ticks *Ixodes ricinus* and *Dermacentor reticulatus*, Germany. Parasit Vectors 2012;5:285.
- [21] Obiegala A, Pfeffer M, Pfister K, Tiedemann T, Thiel C, Balling A, et al. 'Neoehrlichia mikurensis' and *Anaplasma phagocytophilum*: prevalences and investigations on a new transmission path in small mammals and ixodid ticks. Parasit Vectors 2014;7:563.
- [22] Krücke J, Schreiber C, Maaz D, Kohn M, Demeler J, Beck S, et al. A novel high-resolution melt PCR assay discriminates *Anaplasma phagocytophilum* and 'Candidatus Neoehrlichia mikurensis'. J Clin Microbiol 2013;51:1958–61.
- [23] Schreiber C, Krücke J, Beck S, Maaz D, Pachnicke S, Krieger K, et al. Pathogens in ticks collected from dogs in Berlin/Brandenburg, Germany. Parasit Vectors 2014;7:535.
- [24] Silaghi C, Pfister K, Overzier E. Molecular investigation for bacterial and protozoan tick-borne pathogens in wild boars (*Sus scrofa*) from southern Germany. Vector Borne Zoonotic Dis 2014;14:371–3.
- [25] Hornok S, Meli ML, Gönczi E, Hofmann-Lehmann R. First evidence of 'Candidatus Neoehrlichia mikurensis' in Hungary. Parasit Vectors 2013;6:267.
- [26] Szekeres S, Claudia Coipan E, Rigo K, Majoros G, Jahfari S, Sprong H, et al. 'Candidatus Neoehrlichia mikurensis' and *Anaplasma phagocytophilum* in natural rodent and tick communities in Southern Hungary. Ticks Tick Borne Dis 2015;6:111–6.
- [27] Capelli G, Ravagnan S, Montarsi F, Ciocchetta S, Cazzin S, Porcellato E, et al. Occurrence and identification of risk areas of *Ixodes ricinus*-borne pathogens: a cost-effectiveness analysis in north-eastern Italy. Parasit Vectors 2012;5:61.
- [28] Baráková I, Derdáková M, Selyemová D, Chrostáć M, Špitalská E, Rosso F, et al. Tick-borne pathogens and their reservoir hosts in northern Italy. Ticks Tick Borne Dis 2017. pii: S1877-959X(17)30014-6. <https://doi.org/10.1016/j.ttbdis.2017.08.012>.
- [29] Otranto D, Dantas-Torres F, Giannelli A, Latrofa MS, Cascio A, Cazzin S, et al. Ticks infesting humans in Italy and associated pathogens. Parasit Vectors 2014;7:328.
- [30] Movila A, Toderas I, Uspenskaja I, Conovalov J. Molecular detection of tick-borne pathogens in *Ixodes ricinus* from Moldova collected in 1960. Ticks Tick Borne Dis 2013;4:359–61.
- [31] Jenkins A, Kristiansen BE, Allum AG, Akare RK, Strand L, Kleveland Ej, et al. *Borrelia burgdorferi* sensu lato and *Ehrlichia* spp. in *Ixodes* ticks from southern Norway. J Clin Microbiol 2001;39:3666–71.
- [32] Welc-Falęciak R, Kowalec M, Karbowiak G, Bajer A, Behnke JM, Siński E. *Rickettsiaceae* and Anaplasmataceae infections in *Ixodes ricinus* ticks from urban and natural forested areas of Poland. Parasit Vectors 2014;7:121.
- [33] Król N, Obiegala A, Pfeffer M, Lonc E, Kiewra D. Detection of selected pathogens in ticks collected from cats and dogs in the Wrocław Agglomeration, south-west Poland. Parasit Vectors 2016;9:351.
- [34] Kalmár Z, Sprong H, Mihalca AD, Gherman CM, Dumitrache MO, Coipan EC, et al. *Borrelia miyamotoi* and 'Candidatus Neoehrlichia mikurensis' in *Ixodes ricinus* ticks, Romania. Emerg Infect Dis 2016;22: 550–1.
- [35] Raileanu C, Moutailler S, Pavel I, Porea D, Mihalca AD, Savuta G, et al. *Borrelia* diversity and co-infection with other tick borne pathogens in ticks. Front Cell Infect Microbiol 2017;7:36.
- [36] Andersson M, Zaghdoudi-Allan N, Tambo P, Stefanache M, Chitimia L. Co-infection with 'Candidatus Neoehrlichia mikurensis' and *Borrelia afzelii* in an *Ixodes ricinus* tick that has bitten a human in Romania. Ticks Tick Borne Dis 2014;5:706–8.
- [37] Movila A, Alekseev AN, Dubinin HV, Toderas I. Detection of tick-borne pathogens in ticks from migratory birds in the Baltic region of Russia. Med Vet Entomol 2013;27:113–7.
- [38] Potkonjak A, Gutierrez R, Savić S, Vračar V, Nachum-Biala Y, Jurisić A, et al. Molecular detection of emerging tick-borne pathogens in Vojvodina, Serbia. Ticks Tick Borne Dis 2016;7:199–203.
- [39] Špitalská E, Boldis V, Kostanová Z, Kociánová E, Stefanidesová K. Incidence of various tick-borne microorganisms in rodents and ticks of central Slovakia. Acta Virol 2008;52:175–9.
- [40] Pangrácová L, Derdáková M, Pekárik L, Hviščová I, Víchová B, Stanko M, et al. *Ixodes ricinus* abundance and its infection with the tick-borne pathogens in urban and suburban areas of eastern Slovakia. Parasit Vectors 2013;6:238.
- [41] Blaňárová L, Stanko M, Miklisová D, Víchová B, Mošanský L, Kraljik J, et al. Presence of 'Candidatus Neoehrlichia mikurensis' and *Babesia*

- microti* in rodents and two tick species (*Ixodes ricinus* and *Ixodes trianguliceps*) in Slovakia. *Ticks Tick Borne Dis* 2016;7:319–26.
- [42] Svitálková ZH, Haruštiaková D, Mahríková L, Mojšová M, Berthová L, Slovák M, et al. *Candidatus Neoehrlichia mikurensis* in ticks and rodents from urban and natural habitats of south-western Slovakia. *Parasit Vectors* 2016;9:2.
- [43] Palomar AM, García-Álvarez L, Santibáñez S, Portillo A, Oteo JA. Detection of tick-borne '*Candidatus Neoehrlichia mikurensis*' and *Anaplasma phagocytophilum* in Spain in 2013. *Parasit Vectors* 2014;7:57.
- [44] Andersson M, Bartkova S, Lindestad O, Råberg L. Co-infection with '*Candidatus Neoehrlichia mikurensis*' and *Borrelia afzelii* in *Ixodes ricinus* ticks in southern Sweden. *Vector Borne Zoonotic Dis* 2013;13:438–42.
- [45] Labbé Sandelin L, Tolf C, Larsson S, Wilhelmsson P, Salaneck E, Jaenson TG, et al. '*Candidatus Neoehrlichia mikurensis*' in ticks from migrating birds in Sweden. *PLoS One* 2015;10, e0133250.
- [46] Lommano E, Bertaiola L, Dupasquier C, Gern L. Infections and coinfections of questing *Ixodes ricinus* ticks by emerging zoonotic pathogens in Western Switzerland. *Appl Environ Microbiol* 2012;78:4606–12.
- [47] Maurer FP, Keller PM, Beuret C, Joha C, Achermann Y, Gubler J, et al. Close geographic association of human neoehrlichiosis and tick populations carrying '*Candidatus Neoehrlichia mikurensis*' in eastern Switzerland. *J Clin Microbiol* 2013;51:169–76.
- [48] Burri C, Schumann O, Schumann C, Gern L. Are *Apodemus* spp. mice and *Myodes glareolus* reservoirs for *Borrelia miyamotoi*, '*Candidatus Neoehrlichia mikurensis*', *Rickettsia helvetica*, *R. monacensis* and *Anaplasma phagocytophilum*? *Ticks Tick Borne Dis* 2014;5:245–51.
- [49] Lommano E, Dvořák C, Vallotton L, Jenni L, Gern L. Tick-borne pathogens in ticks collected from breeding and migratory birds in Switzerland. *Ticks Tick Borne Dis* 2014;5:871–82.
- [50] van Overbeek L, Gassner F, van der Plas CL, Kastelein P, Nunes-da Rocha U, Takken W. Diversity of *Ixodes ricinus* tick-associated bacterial communities from different forests. *FEMS Microbiol Ecol* 2008;66:72–84.
- [51] Coipan EC, Jahfari S, Fonville M, Maassen CB, van der Giessen J, Takken W, et al. Spatiotemporal dynamics of emerging pathogens in questing *Ixodes ricinus*. *Front Cell Infect Microbiol* 2013;3:36.
- [52] Jahfari S, Hofhuis A, Fonville M, van der Giessen J, van Pelt W, Sprong H. Molecular detection of tick-borne pathogens in humans with tick bites and erythema migrans, in The Netherlands. *PLoS Negl Trop Dis* 2016;10, e0005042.
- [53] Jahfari S, Fonville M, Hengeveld P, Reusken C, Scholte Ej, Takken W, et al. Prevalence of *Neoehrlichia mikurensis* in ticks and rodents from north-west Europe. *Parasit Vectors* 2012;5:74.
- [54] Heylen D, Fonville M, Docters van Leeuwen A, Stroo A, Duisterwinkel M, van Wieren S, et al. Pathogen communities of songbird-derived ticks in Europe's low countries. *Parasit Vectors* 2017;10:497.
- [55] Vayssier-Taussat M, Le Rhun D, Buffet JP, Maaoui N, Galan M, Guivier E, et al. *Candidatus Neoehrlichia mikurensis* in bank voles, France. *Emerg Infect Dis* 2012;18:2063–5.
- [56] Beck R, Čurik VC, Račić I, Šprem N, Vujičić A. Identification of '*Candidatus Neoehrlichia mikurensis*' and *Anaplasma* species in wildlife from Croatia. *Parasit Vectors* 2014;7(Suppl. 1):O28.
- [57] Földvári G, Jahfari S, Rigó K, Jablonczky M, Szekeres S, Majoros G, et al. '*Candidatus Neoehrlichia mikurensis*' and *Anaplasma phagocytophilum* in urban hedgehogs. *Emerg Infect Dis* 2014;20:496–8.
- [58] Diniz PP, Schulz BS, Hartmann K, Breitschwerdt EB. '*Candidatus Neoehrlichia mikurensis*' infection in a dog from Germany. *J Clin Microbiol* 2011;49:2059–62.
- [59] Beck A, Huber D, Antolić M, Anzulović Z, Reil I, Beck R. Retrospective molecular study of canine infectious haemolytic anaemias. *J Comp Pathol* 2015;152:53.
- [60] Hofmann-Lehmann R, Wagmann N, Meli ML, Riond B, Novacco M, Joeckel D, et al. Detection of '*Candidatus Neoehrlichia mikurensis*' and other *Anaplasmataceae* and *Rickettsiaceae* in *Canidae* in Switzerland and Mediterranean countries. *Schweiz Arch Tierheilkd* 2016;158:691–700.
- [61] Liesner JM, Krücke J, Schaper R, Pachnicke S, Kohn B, Müller E, et al. Vector-borne pathogens in dogs and red foxes from the federal state of Brandenburg, Germany. *Vet Parasitol* 2016;224:44–51.
- [62] Pekova S, Vydra J, Kabickova H, Frankova S, Haugvicova R, Mazal O, et al. *Candidatus Neoehrlichia mikurensis* infection identified in 2 hematooncologic patients: benefit of molecular techniques for rare pathogen detection. *Diagn Microbiol Infect Dis* 2011;69:266–70.
- [63] von Loewenich FD, Geissdörfer W, Disque C, Matten J, Schett G, Sakka SG, et al. Detection of '*Candidatus Neoehrlichia mikurensis*' in two patients with severe febrile illnesses: evidence for a European sequence variant. *J Clin Microbiol* 2010;48:2630–5.
- [64] Grankvist A, Andersson PO, Mattsson M, Sender M, Vaht K, Höper L, et al. Infections with the tick-borne bacterium '*Candidatus Neoehrlichia mikurensis*' mimic noninfectious conditions in patients with B cell malignancies or autoimmune diseases. *Clin Infect Dis* 2014;58:1716–22.
- [65] Andréasson K, Jönsson G, Lindell P, Gülfé A, Ingvarsson R, Lindqvist E, et al. Recurrent fever caused by '*Candidatus Neoehrlichia mikurensis*' in a rheumatoid arthritis patient treated with rituximab. *Rheumatology (Oxford)* 2015;54:369–71.
- [66] Grankvist A, Moore ERB, Svensson Stadler L, Pekova S, Bogdan C, Geißdörfer W, et al. Multilocus sequence analysis of clinical '*Candidatus Neoehrlichia mikurensis*' strains from Europe. *J Clin Microbiol* 2015;53:3126–32.
- [67] Wennerås C, Goldblatt D, Zancolli M, Mattsson M, Wass L, Hörrkö S, et al. Natural IgM antibodies in the immune defence against neoehrlichiosis. *Infect Dis (Lond)* 2017;49:809–16.
- [68] Fehr JS, Bloomberg GV, Ritter C, Hombach M, Lüscher TF, Weber R, et al. Septicemia caused by tick-borne bacterial pathogen *Candidatus Neoehrlichia mikurensis*. *Emerg Infect Dis* 2010;16:1127–9.
- [69] Quarsten H, Granqvist A, Høyvoll L, Myre IB, Skarpaas T, Kjelland V, et al. '*Candidatus Neoehrlichia mikurensis*' and *Borrelia burgdorferi* sensu lato detected in the blood of Norwegian patients with erythema migrans. *Ticks Tick Borne Dis* 2017;8:715–20.
- [70] Granqvist A, Sandelin LL, Andersson J, Fryland L, Wilhelmsson P, Lindgren PE, et al. Infections with *Candidatus Neoehrlichia mikurensis* and cytokine responses in 2 persons bitten by ticks, Sweden. *Emerg Infect Dis* 2015;21:1462–5.
- [71] Wennerås C. Infections with the tick-borne bacterium '*Candidatus Neoehrlichia mikurensis*'. *Clin Microbiol Infect* 2015;21:621–30.