

Bacteria of the genus *Anaplasma* – characteristics of *Anaplasma* and their vectors: a review

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ABSTRACT: Over recent years, there has been a growing interest in bacteria from the genus *Anaplasma*, especially the species *A. marginale*, *A. ovis* and *A. phagocytophilum*. It is connected with the pathogenic activity of these bacteria in farm animals, and also, though to a lesser degree, in humans. Anaplasmosis, a disease caused by various species of anaplasma, is an especially important issue for animal breeders. The main vectors of the *Anaplasma* bacteria are ticks, common arachnida occurring everywhere in the world, especially the genera *Ixodes*, *Dermacentor*, *Rhipicephalus* and *Amblyomma*. The genus *Anaplasma* includes obligate intracellular bacteria, parasitizing in the vacuoles of cells in eukaryotic hosts. *A. marginale*, *A. centrale*, *A. ovis* and *A. bovis* are obligate intracellular bacteria parasitizing in erythrocytes and monocytes of higher vertebrates, mostly ruminants. *A. platys* is mainly a pathogen of canines (displaying tropism to thrombocytes) and the species *A. phagocytophilum* (displaying tropism to granulocytes) is pathogenic to people and domestic animals. In this paper we present characteristics and differentiation of six species of the genus *Anaplasma* and their vectors in the world.

Keywords: ticks; *Anaplasma* vectors; anaplasmosis; bacteria from the genus *Anaplasma*

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1. Introduction

Over recent years, there has been a growing interest in bacteria from the genus *Anaplasma*, especially in the species *A. marginale*, *A. ovis* and *A. phagocytophilum*. It is connected with the pathogenic activity of these bacteria in farm animals, and also, though to a lesser degree, in people. Anaplasmosis,

a disease caused by various species of anaplasma, is an especially important issue for animal breeders. Apart from the costs of the additional veterinary care, this disease reduces the animal's body weight, causes abortions, reduces milk production and frequently leads to death (Splitter et al., 1955; Sainz et al., 1999; Melendez, 2000; Stuenkel et al., 2002, 2003). When it comes to effects on people, only the spe-

cies *A. phagocytophilum* is pathogenic; it causes Human Granulocytic Anaplasmosis (HGA). The disease usually has mild symptoms, but in some cases, especially in individuals with a weakened immune system, it may be fatal (Fishbein et al., 1994; Dumler and Bakken, 1998; Dumler and Broqui, 2004).

Bacteria from the genus *Anaplasma* belong to the Procaryota kingdom, the family Anaplasmataceae. The present classification has been valid since 2001 (Table 1), when many independent researchers reorganized the families Rickettsiaceae and Ehrlichiaeae (Dumler et al., 2001).

Anaplasma belong to obligate intracellular microorganisms, Gram-negative bacteria, living in the blood cells of mammals. Vertebrates can be their reservoir, i.e. an environment where the pathogen can live and proliferate for many years. However, in many cases bacteria from the genus *Anaplasma* cause diseases in domestic animals and people.

An essential part in the life cycle of the bacteria is played by vectors, organisms which contribute to their circulation in the environment. The main vectors of the *Anaplasma* bacteria are ticks, common arachnida occurring everywhere in the world, especially the genera *Ixodes*, *Dermacentor*, *Rhipicephalus* and *Amblyomma*. Numerous research results have confirmed that a vector for anaplasma in Europe is the universally occurring tick *I. ricinus*, which can also be found in the African part of the Mediterranean Sea Basin (Ogden et al., 1998; Liz et al., 2000; Walker et al., 2001; Skotarczak et al., 2003). In the south of Europe, another vector of anaplasma is *R. sanguineus* (Sainz et al., 1999), and on the eastern borders of the continent and in Asia *I. persulcatus*. *R. sanguineus*, or the Brown dog tick, appears universally outside the European continent as well, i.e. in regions of Northern Africa, India, and in the territory of the United States (Stafford III, 2007).

Outside Europe, the most frequently mentioned vectors of the *Anaplasma* bacteria are the ticks *I. scapularis* (Black-legged tick) and *I. pacificus* (Western black-legged tick). These species occur mostly in the United States of America and Canada (Parola and Raoult, 2001; Stafford III, 2007). In North and Central America, bacteria are also transmitted by ticks from the species *A. americanum* – known as the Lone star tick (Parola and Raoult, 2001; Stafford III, 2007).

Other vectors of anaplasma are the species from the genus *Haemaphysalis*, such as *H. leporispalustris*, *H. lagrangei* and *H. longicornis*, which are com-

mon in Japan, India, Russia, China, Korea, Thailand and in North America (Goethert and Telford III, 2003; Parola et al., 2003; Kawahara et al., 2006; Kim et al., 2006).

2. Taxonomical position of bacteria from the genus *Anaplasma*

In 2001, significant reorganizational changes were carried out within the order Rickettsiales. As a result of this reorganization, the family Ehrlichiaeae was replaced by the family Anaplasmataceae, and some changes were made in the classification of the bacteria within the genera (Table 1) (Dumler et al., 2001).

The genera *Anaplasma*, *Ehrlichia*, *Neorickettsia* and *Wolbachia* include obligate intracellular bacteria, parasitizing in the vacuoles of cells in eukaryotic hosts. Their systematic classification was performed on the basis of morphological, ecological, epidemiological and clinical characterization. However, the development of molecular methods over recent years has enabled the partial recognition of the genome of these bacteria and consequently the partial change in their systematic position. A comparison was performed for all sequences of conservative genes 16S rDNA (the gene coding 16S rRNA for the small subunit of the ribosome) and *groESL* (the gene coding the thermal shock protein), for those deposited in the GenBank up to the year 2000 (Dumler et al., 2001). A phylogenetic analysis was then carried out. The independent comparative analysis of the sequence of both genes gave similar results. Dendrograms of the sequences of the genes 16S rRNA and *groESL* have a similar arrangement, and bacteria are positioned in clades similarly on both independent phylogenetic trees, a further proof of the correctness of the research performed. Their accuracy is also shown by the high bootstrap obtained in both cases. Moreover, the new systematical division, emerging from the relationship analyses, is confirmed by the biological and antigenic characterization of each species (Dumler et al., 2001). Details concerning the changes in taxonomy in these groups of the bacteria are presented in Table 1.

Animal pathogens were attributed to the genus *Anaplasma*, such as *A. centrale*, *A. marginale*, *A. platys*, *A. ovis* and *A. bovis*, and also the aetiological factor of human anaplasmosis, *A. phagocytophilum* (Table 1).

Table 1. Taxonomy of the genera *Ehrlichia* and *Anaplasma* before and after 2001. Letters A, B, B2, B3, B4, B5, B6, B7, C indicate a change in the systematic position

Taxonomy before 2001		Taxonomy after 2001	
Class α-proteobacteria		Class α-proteobacteria	
Order II	Rickettsiales	Order II	Rickettsiales
Family I	Rickettsiaceae	Family I	Rickettsiaceae
Genera I	<i>Rickettsia</i>	Genera I	<i>Rickettsia</i>
Genera II	<i>Orientia</i>	Genera II	<i>Orientia</i>
Genera III	<i>Wolbachia</i> – A		
Family II	Ehrlichieae	Family II.	Anaplasmataceae
Genera I	<i>Ehrlichia</i> – B	Genera I	<i>Anaplasma</i>
	Species	Species	<i>A. marginale</i>
	<i>E. canis</i> – B2		<i>A. centrale</i>
	<i>E. chaffeensis</i> – B3		<i>A. phagocytophilum</i> – B4
	<i>E. equi</i> – B4		<i>A. platys</i> – B5
	<i>E. phagocytophila</i> – B4		<i>E. canis</i> – B2
	czynniki HGE – B4	Genera IV	<i>Ehrlichia</i> – B
	<i>E. platys</i> – B5		Species
	<i>E. risticii</i> – B6		<i>E. chaffeensis</i> – B3
	<i>E. sennetsu</i> – B7		<i>E. ruminantium</i> – C
Genera III	<i>Anaplasma</i>	Genera V	<i>Neorickettsia</i>
	Species		Species
	<i>A. marginale</i>		<i>N. helminthoeca</i>
	<i>A. centrale</i>		<i>N. risticii</i> – B6
Genera IV	<i>Cowdria</i>		<i>N. sennetsu</i> – B7
Genera V	<i>Neorickettsia</i>	Genera VI	<i>Wolbachia</i> – A

3. History of *Anaplasma* discovery

Cases of tick fever occurring in goats, sheep and calves have been reported in Europe since 1780. The aetiological factor of this disease was not known but the description of symptoms corresponds well to the modern characterizations of anaplasmosis. Later reports frequently described the occurrence of tick fever in farm animals, mostly in small ruminants, and in dogs as well. Sometimes the disease led to the death of the animals, however its aetiology was not known yet. 150 years later, the first report was released that the disease in question was caused by a small bacterium *E. phagocytophila* (presently known as *A. phagocytophilum*) displaying tropism to granulocytes (Jenkins et al., 2001).

The first reports on *A. marginale* appeared as early as in 1894, when Salmon and Smith detected the presence of inclusion in the cells of calf erythrocytes. The first full description came from Sir Arnold Theiler, who observed bacteria in erythrocytes of South African cattle in 1910 (Kocan et al., 2003). He discriminated two subspecies: the one more frequently creating “marginal points” in erythrocytes of calves (*A. marginale*) and the other more frequently creating concentrations in the central part of blood cells, hence the name *A. centrale*. The latter of the subspecies was found to be less pathogenic to domestic animals (Kocan et al., 2003). The following years saw the discoveries of other species of anaplasmosis, pathogenic to animals (Table 2), such as *A. ovis*, *A. platys* (formerly

known as *E. platys*) and *A. bovis* (Harvey et al., 1978; Kuttler, 1984).

After some research, it was shown that anaplasmosis may affect humans; the first case of human HGA was observed in 1994 in the United States. It was caused by *A. phagocytophilum* (an agent of human granulocytic anaplasmosis, at first described as an agent of granulocytic ehrlichiosis) (Chen et al., 1994; Dumler et al., 2001). In 1996 this disease was also diagnosed in Europe, in Slovenia (Petrovec et al., 1997) and reported in Poland in 2001 (Tylewska-Wierzbanowska et al., 2001).

4. Characterization of bacterium from the genus *Anaplasma*, pathogenic to farm animals and people

4.1. *Anaplasma marginale*

A. marginale are obligate intracellular bacteria parasitizing in erythrocytes of higher vertebrates, mostly ruminants (Table 2). It causes so called bovine anaplasmosis which has been described in domestic and wild animals, i.e. in calves, water buffalo, bison, African antelopes and the mule deer (Kocan et al., 2003). In infected animals, red blood cells were observed to have from 4 to 8 bacterial inclusions (Barbet et al., 1999; Kocan et al., 2004). The time of incubation of the disease depends on the degree of infection and can last from 7 to 60 days, on average approximately 28 days. During

Table 2. The characteristic of pathogens of genus *Anaplasma*

Aetiological agent		Disease	Vector	Infected organism or host	Infected cell
before 2001	after 2001				
<i>Ehrlichia bovis</i>	<i>Anaplasma bovis</i>	bovine anaplasmosis	<i>Haemaphysalis</i> sp. <i>Rhipicephalus</i> sp. <i>Amblyomma</i> sp.	ruminants farming, small mammals	monocytes
<i>Anaplasma ovis</i>	<i>Anaplasma ovis</i>	bovine anaplasmosis	<i>Dermacentor</i> sp.	small ruminants (sheep, goats)	erythrocytes
<i>Anaplasma marginale</i>	<i>Anaplasma marginale</i>	bovine anaplasmosis	<i>Ixodes</i> sp. <i>Dermacentor</i> sp.	ruminants farming	erythrocytes
<i>Anaplasma centrale</i>	<i>Anaplasma centrale</i>	bovine anaplasmosis	<i>Ixodes</i> sp. <i>Haemaphysalis</i> sp.	ruminants farming	erythrocytes
<i>E. equi</i> <i>E. phagocytophila</i> Czynnik HGE	<i>Anaplasma phagocytophilum</i> (HGA agent)	human and animal granulocytic anaplasmosis	<i>Ixodes</i> sp. <i>Dermacentor</i> sp.	small ruminants forming and wild, horses, dogs, humans	granulocyte
<i>E. platys</i>	<i>Anaplasma platys</i>	canine cyclic thrombocytopenia	<i>Rhipicephalus sanguineus</i>	dogs	platelets

that time, the number of infected blood cells increases geometrically and in the acute phase of the disease 70% or more erythrocytes can be infected. During infection, cattle erythrocytes are phagocytosed by reticuloendothelial cells, which leads to anaemia and icterus without haemoglobinaemia and haemoglobinuria. Moreover, a clinical picture of the disease also includes fever, weight loss, abortions and lethargy. Animals older than two years may even die due to the infection. Calves that experience the acute phase of the disease are permanently infected with the cyclic low-level rickettsemia or become carriers of *A. marginale*, and consequently in both cases they are reservoirs for the bacteria (Barbet et al., 1999; Kocan et al., 2003).

A. marginale is a tick-borne pathogen whose presence is mainly connected with tropical or subtropical regions, although these bacteria are detected more and more frequently in other regions of the world, e.g. Europe (Sicily, Hungary and Spain), Australia, South Africa (De Waal, 2000; de la Fuente et al., 2004, 2005; Kocan et al., 2004; Naranjo et al., 2006; Hornok et al., 2007).

From an economic point of view, *A. marginale* plays an important role in the world economy. De Waal (2000) revealed that more than 99% of the total cattle population in South Africa was at risk. Rajput et al. (2005) informed us that in Pakistan, *Anaplasma* parasites in buffalo and cattle *Bubalus bubalis* were 30% and 52%, respectively. It is estimated that in the United States alone, losses incurred by *A. marginale* may reach approx. 300 million dollars a year (McCallon, 1973). Losses in Latin America reach over 800 million dollars a year (Lonibardo, 1976). Such great losses are also recorded in Africa, where cattle are often the only means of income. This has bred the idea of genetically modifying the cattle to achieve an inborn resistance to *A. marginale* (Melendez, 2000).

4.2. *Anaplasma centrale*

A. centrale is a parasite in the erythrocytes of ruminants, mainly cattle (Table 2). As opposed to *A. marginale*, it creates concentrations in the central part of the cell. This bacterium is widely-spread throughout the world. In spite of morphological differences, occurrence and virulence, *A. centrale* is closely related to *A. marginale*. *A. centrale* is the cause of mild anaemia in most cases

of cattle infection (Kuttler, 1984; Kocan et al., 2003; Rajput et al., 2005). According to Theiler, *A. centrale* is less pathogenic to cattle than *A. marginale* but, most importantly, occasionally gives resistance against the latter. Hence it is used for the preparation of live vaccine strains, assuring immunological protection against bovine anaplasmosis. Such vaccines are produced in Africa, Australia, Latin America and Israel (Kocan et al., 2003).

4.3. *Anaplasma bovis*

A. bovis is a bacterium detected mainly in cattle, but also observed in small mammals which are probably a reservoir of this bacterium (Table 2) (Goethert and Telford III, 2003). It occurs in monocytes, and the disease it causes is called monocytic anaplasmosis. The symptoms of the disease are most visible in calves, but also in adult animals; symptoms include weakening of the body, marked reduction in weight, elevated temperature, enlargement of prescapular lymph nodes, paling of the mucous membranes, and in many cases an elevated amount of secreted mucus (Uilenberg, 1997; Santos and Carvalho, 2006).

A. bovis has been detected in Brazil, North America, Africa and Japan (Goethert and Telford III, 2003; Kawahara et al., 2006; Santos and Carvalho, 2006).

4.4. *Anaplasma ovis*

A. ovis mainly parasitizes small ruminants, i.e. sheep and goats, and its presence has been confirmed in most regions of the world both in farm and wild animals (Kuttler, 1984). Similarly to *A. marginale* and *A. centrale*, these bacteria live in erythrocytes, hence the anaemia of the infected animals (Splitter et al., 1956). In the case of *A. ovis*, bacterial inclusions are found 35–40% of the time in the central or submarginal part of the erythrocyte of the host, and the remaining 60–65% of the time in the marginal part (Shompole et al., 1989).

Experimental research shows that animals experiencing an acute phase of the disease display depression, debility, marked decline in body weight, fever and progressive anaemia whose consequence is among other things a reduction in milk production. The infection is also potentially lethal. The symptoms of the disease depend on age, on the general condition of the animals and their breed (Splitter et al., 1955; Shompole et al., 1989).

Clinical symptoms occurring in animals resulting from the presence of *A. ovis* in blood cause huge losses in farming stock. This has considerable significance for poorer countries in tropical and subtropical regions where animal stocks consist mainly of sheep and goats. They are the basic and frequently only producers of milk and meat, and also provide farms with natural manure used in agriculture (Jensen, 1955). Hence the losses *A. ovis* causes in the economy, especially in those regions, are significant and create a need for further research concerning the spreading and pathogenicity of *A. ovis* and possibilities for prophylaxis (Kuttler, 1984).

Contemporary research conducted by various scientific centres shows that *A. ovis* is a problem not only in poorer countries. These bacteria have been detected in the U.S.A., Italy and Hungary (de la Fuente et al., 2002, 2005; Hornok et al., 2007). The occurrence of *A. ovis* is not widespread there, but one ought to allow for the fact that a return to traditional methods of farming (pasturage on meadows) can facilitate the expansion of this pathogen. The spread of this bacterium may also be enhanced by modern climatic changes.

4.5. *Anaplasma platys*

For the first time (1978), *A. platys* was detected in the U.S.A. (Florida; Harvey et al., 1978). It was also detected in Central America (Venezuela), Asia (Korea, Japan, China, Taiwan and Thailand), Australia and Africa. In Europe, the presence of *A. platys* was confirmed around the Mediterranean Basin, in France, Greece, Spain and Italy (Sainz et al., 1999; Suksawat et al., 2001; Sparagano et al., 2003; de la Fuente et al., 2006; Kawahara et al., 2006; Kim et al., 2006).

A. platys is mainly a pathogen of canines, usually dogs. It lives in blood cells, causing Canine Cyclic Thrombocytopenia (CCT) (Table 2). Clinical symptoms of this disease vary depending on the species of the dog, but fundamentally CCT causes weakening of the animal, lethargy, anorexia, respiratory distress, fever, increased mucous secretion, purulent ocular discharge, splenomegaly and muzzle hyperkeratosis (Sainz et al., 1999). These bacteria do not cause death, and treatment with suitable antibiotics is usually successful after four weeks. Doxycycline is one of such effective antibiotic (Sainz et al., 1999).

4.6. *Anaplasma phagocytophilum*

A. phagocytophilum is a frequently described bacterium from the genus *Anaplasma*. It has been detected in domestic animals, mostly small ruminants, but also in people (Table 2). These bacteria are an aetiological factor of granulocytic anaplasmosis which in the case of human infections is defined as Human Granulocytic Anaplasmosis – HGA (Human Anaplasmosis – HA).

Human anaplasmosis is a multisystemic disease, difficult to diagnose due to unspecific symptoms which are similar to the common flu. It may have mild or heavy symptoms, depending on the patient's condition. One estimate states that about 40% of those infected require hospitalization. Early detected HGA is curable. In Europe no deadly cases have been reported, and in the U.S.A. the mortality rate was 7–10% (Fishbein et al., 1994; Dumler and Bakken, 1998; Blanco and Oteo, 2002; Dumler and Broqui, 2004). Although anaplasmosis alone is not perhaps a life-threatening disease, frequent complications such as renal failure, intravascular coagulation, cardiomegaly, seizures or coma, may all be fatal. These involved mostly the elderly, who became ill after receiving immunosuppressive medicines, and children (Fishbein et al., 1994; Dumler and Bakken, 1998).

In the case of farm animals anaplasmosis caused by *A. phagocytophilum* is a disease with subclinical or heavy symptoms. It has unspecific symptoms, such as fever, drowsiness, anorexia, abortions, drop in body weight and reduction in milk production. Left unattended, especially in weaker individuals, it can lead to death (Stuenkel et al., 2002; Garcia-Perez et al., 2003).

5. Vectors of bacteria from the genus *Anaplasma*

5.1. Morphology and behaviour of questing ticks

In epidemiology, a vector is an organism which contributes to the spreading of a pathogen in a population of hosts. Among saprophytes, it is ticks that play the greatest part in the epidemiology of disease transmission. Anaplasmosis is a typical tick-borne zoonosis, in which ticks are the vectors, i.e. they transfer bacteria to the blood of the higher vertebrates on which they feed.

The world fauna of ticks, representative of arachnida (phylum *Arachnida*), includes about 860 species divided into four families. All ticks are obligate, blood-sucking arthropods (*Arthropoda*), belonging to the subdivision Acari (mites) (Siuda, 1991). All of them parasitize vertebrate animals. Their contact with hosts is usually temporary, restricted in most species to only receiving nourishment, but their medical, veterinary and economic impact is immense. Two families of proper ticks especially – Ixodidae and Amblyomidae – have an impact on the economy and medicine. Parasitizing ticks cause immediate results, such as damage to the host's skin or local inflammations; and more importantly being reservoirs of viruses, bacteria or protozoans, ticks may transfer pathogens to people and animals (Siuda, 1991).

Ticks do not have a segmentation characteristic of most other arachnids. Their bodies are oval, narrowing towards the front, with the gnathosoma embedded on their idiosoma. The gnathosoma made from mouth organs is also the main adhesive organ. The idiosoma, from the dorsal side, is coated with so-called scutum, which in males covers almost all the dorsum, and in females only the forepart. The remaining part of the dorsal surface of the idiosoma is covered by an extensible epidermis, so called alloscutum, which enables the extraction of large amounts of blood by females, nymphs and larvae. In males, the epidermis covers only the side surfaces of the idiosoma. The size of the scutum is a means of differentiation between male and female ticks. On the abdominal side, there are four pairs of limbs. On the feet of the first pair, on the dorsal side, the Haller's organ can be noticed. It is the most important sensory organ of ticks, enabling the search for a host (Siuda, 1991; Buczek and Magdon, 1999). Some species, e.g. from the genera *Amblyomma* and *Rhipicephalus*, have eyes, while other ticks have photosensitive cells which react to light intensity changes (Buczek and Magdon, 1999).

Some species of ticks actively seek hosts (e.g. *I. scapularis*, *I. pacificus* or *A. americanum*), others assume the wait-and-see attitude (e.g. *I. ricinus*). They cling to plants, and when a potential host approaches, they raise up their first pair of legs and hook onto the host (hair, clothes, etc.) with small claws (Buczek and Magdon, 1999).

The tick's reaction to a host depends on several factors. Research has shown that a major factor influencing the behaviour of questing individuals is an increased concentration of CO₂ in the air, i.e.

exhalations from a host. Changes in the concentration of carbon dioxide in the air, registered by ticks, make them adopt an expectant position (in passive species) or move towards the source of the gas (in active species). The scent of a host and its body temperature can also significantly influence the tick's choice. Other impulses that activate ticks are pheromones, urine, sounds given off by animals or an environmental factor – light intensity (Buczek and Magdon, 1999). Stimulating factors vary and characterize different species of ticks.

The behaviour of ticks may also be influenced by seasonal and diurnal rhythms characteristic of local populations. Seasonal activity is conditioned by ambient temperature, and diurnal activity by potential host activity, ambient temperature and humidity.

5.2. Vectors of bacteria from the genus *Anaplasma* in Europe

The most widespread European tick is the common tick *I. ricinus*, a vector for bacteria from the genus *Anaplasma*. Sometimes *I. ricinus* is also called the European sheep tick (Parola and Raoult, 2001). This common tick is a blood-sucking arachnid, belonging to the order Ixodida, the genus *Ixodes* (Siuda, 1991). Its parasitizing activity on vertebrates, including humans, causes many pathological alterations, sometimes even life-threatening for its hosts (swelling, inflammatory disorders of skin, allergy, paralysis). *I. ricinus* belongs to extranidamental ectoparasites (which feed on trans migrant animals in the open environment), with a three-host life cycle (each active developmental phase of the tick has an independent host) (Siuda 1991).

The life cycle of the tick *I. ricinus* includes the following phases: larva, nymph and adult (imago). This species is characterized by seasonality, or in other words dependence of the activity and course of each phase of the life cycle on the seasons, characteristic of a temperate climate. The life cycle of the common tick in temperate climates lasts several years, most frequently approximately three years, but sometimes under favourable environmental conditions even longer (Siuda, 1991).

In the annual cycle, ticks show seasonal activity which depends on their geographical location. In temperate climates it starts from mid-April (sometimes earlier, even in March) and lasts to the beginning of November with two peaks – the first

from May to mid-June, the second in September. After hot summers, it is possible that the autumnal peak will not occur. It is accepted that mature forms can be active already at temperatures of 16°C, with nymphs active at temperatures of 25°C. The embryonic period lasts, depending on the ambient temperature, from 24 to 50 days (Siuda, 1991; Skotarczak et al., 1999).

Besides seasonal activity, ticks are also observed by diurnal activity. It is connected, among other things, with local environmental conditions, such as temperature and humidity, and with the host activity. Species of ticks parasitizing on animals are characterized by diurnal activity, and will feed most intensely in these hours. *I. ricinus* is the most active in morning hours (about 8) and in the afternoon and at night (15–24), but can attack at other times of the day as well (Siuda, 1991; Buczek and Magdon, 1999).

Among European ticks from the genus *Ixodes*, the taiga tick – *I. persulcatus*, is also known as a vector for anaplasma (Figure 1). The geographical range of this species covers north-eastern Europe, areas of the taiga and forested steppes. Similarly to *I. ricinus*, the taiga tick is an extranidamental parasite, i.e. it has three hosts in its lifetime. Also like the common tick, it is characterized by a lack of host specificity (Siuda, 1991). The life cycle of *I. persulcatus* can last from two to five years.

R. sanguineus, the dog tick (Brown dog tick), belongs to cosmopolitan ticks (the detailed character-

istics of *R. sanguineus* in chapter 5.3). Its presence has been established in all regions of Africa, India, in the United States and around the Mediterranean Basin (Parola and Raoult, 2001; Stafford III, 2007). In Europe, the occurrence of another tick from the genus *Rhipicephalus* was observed in Spain, namely *R. bursa*, which is also probably a vector for pathogens from the genus *Anaplasma* (de la Fuente et al., 2004).

5.3. Vectors of bacteria from the genus *Anaplasma* outside Europe

The most frequently mentioned pathogen vector outside Europe is the tick *I. scapularis* (Black-legged tick), which is also called a deer tick in some regions. *I. scapularis* can be found in the eastern and south-eastern United States and Canada (Figure 1). The biotope of this species is forests with a rich forest floor. *I. scapularis* hosts include mainly deer, dogs, small birds and rodents, but also people. Males of this tick reach a length of 2–2.7 mm, females are larger, approximately 3–3.7 mm. The imago has a red-brown colour, with younger ones being brighter. *I. scapularis* prefers a deciduous forest and a habitat containing litter (Stafford III, 2007).

I. scapularis is a species with the two-host development cycle. Juvenile phases, larvae and nymphs feed on one host and adult forms on the other. A female feeds for about 3–7 days, then falls off and

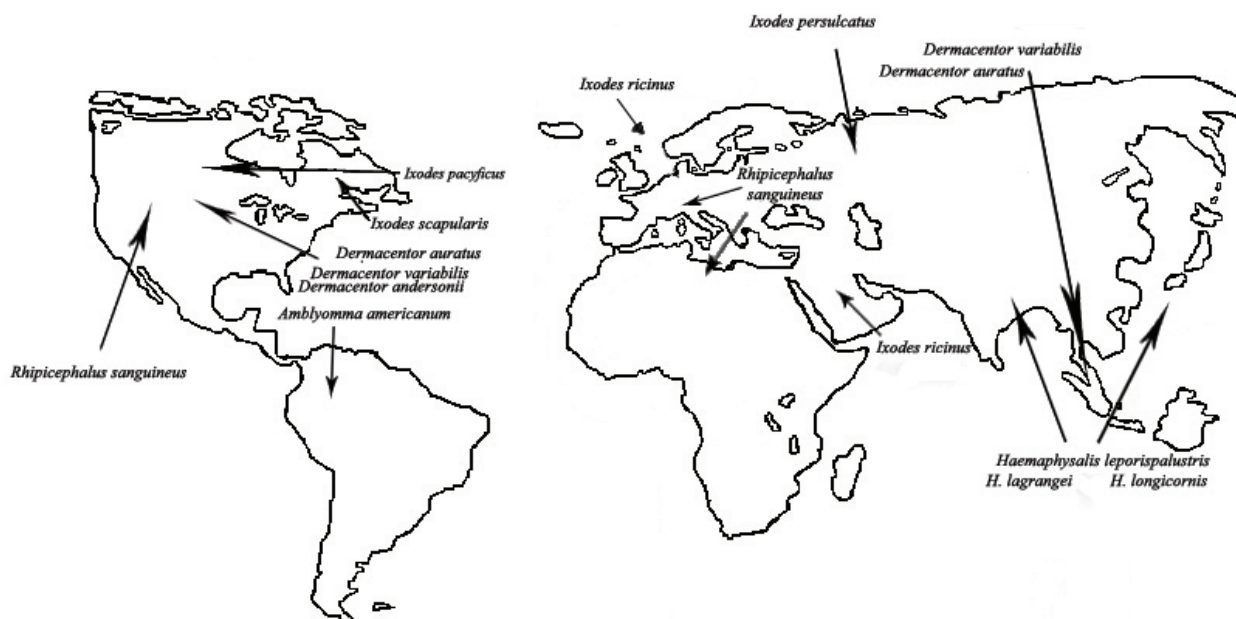


Figure 1. The world distribution of vectors of bacteria from the genus *Anaplasma*

oviposits in the litter, laying 2 000 or more eggs. Larvae hatch from eggs on the ground, and then attach to the first host. After feeding, the larva moults into a nymph which feeds on the same host. Only after filling itself with blood does the nymph fall from the host and transform into an adult on the ground. The imago actively seeks a second host, where it feeds (Stafford III, 2007). Another species of ticks which is physically similar to *I. scapularis* is *I. pacificus*, occurring mostly in the west of the United States of America (Pacific US and British Columbia). *I. pacificus*, also called the Western black-legged tick, is found mostly on the edges of forests, coppices or in open spaces, such as meadows and pastures. Its hosts are people, deer, dogs, coyotes, small rodents, wild hogs, horses and lizards (Stafford III, 2007).

A. americanum (Lone Star Tick) is also considered a vector for bacteria from the genus *Anaplasma*. It appears in regions of the southeastern United States, and also in some regions of Central America. Its biotope is wooded areas with brush, especially hickory-oak forests, along creeks and rivers (Parola and Raoult, 2001). The colour of the *A. americanum* imago varies from brown to tan, and in females the dorsum has a characteristic silvery white spot, so-called “lone star”. Males of the *A. americanum* have some small spots or streaks around the perimeter of the body. The average size of an unfed imago ranges from 3 to 4 mm. All developmental phases can feed on humans, and both small and large mammals as well. Juvenile phases have been observed mainly on small birds and rodents. A female of *A. americanum* lays up to 3 000 eggs (Parola and Raoult, 2001; Stafford III, 2007).

An essential role in the transmission of *Anaplasma* is played by ticks from the genus *Dermacentor* sp., most of all *D. variabilis* (American dog tick). This tick, also called the wood tick, occurs in grassy meadows, wooded areas, near barns, kennels, and anywhere the animals are kept. The imago of *D. variabilis* is dark brown. Adult and juvenile forms feed mostly on small rodents, opossums, raccoons, horses, deer and cattle. The female of this tick attains 13 mm in length after feeding. It feeds for about 10–12 days, and after falling off it can lay from 3 000 to 7 000 eggs. The life cycle lasts about two years. Among ticks from the genus *Dermacentor*, *D. andersoni* (Rocky Mountain wood tick) and *D. auratus* are included. *D. andersoni* is also a vector of anaplasma, living outdoors in low vegetation. This tick is brown, and the scutum is

grey. *D. andersoni* hosts are rodents, sheep, cattle, deer and humans. The ticks *D. andersoni* and *D. auratus* occur in the United States and Asia (Parola and Raoult, 2001; Parola et al., 2003; Stafford III, 2007).

Both within and outside Europe, one of the common ticks as a vector for *Anaplasma* is the dog tick *R. sanguineus* (Brown dog tick, or interchangeably Red dog tick). These ticks are characterized by their elongated body without ornamentation, with a reddish-brown colour. These ticks occur most frequently in houses, kennels and dog runs (Parola and Raoult, 2001). *R. sanguineus* has a three-host developmental cycle, but contrary to the aforementioned species from the genus *Ixodes*, each stage of *R. sanguineus* has a high specificity to one host – the dog, while humans are only an accidental host. The life cycle of *R. sanguineus* may last 4–5 months. The female lays a minimum of 4 000 eggs. Humans are parasitized by *R. sanguineus* extremely seldom, and only when there is no other host in its environment. This tick occurs most of all in Northern Africa, India, United States and Europe, mainly around the Mediterranean Basin (Figure 1) (Stafford III, 2007).

Over recent years, some reports have shown that ticks from the genus *Haemaphysalis* also play an important part in the transmission of anaplasma. Most frequently mentioned are the species *H. leporispalustris* (*A. bovis*) and *H. longicornis* (*A. bovis*, *A. centrale*, *A. phagocytophilum*). These ticks have two-host life cycles, and the length of the life cycle does not exceed one year. Ticks from the genus *Haemaphysalis* occur mostly in Japan, Russia (Far East), China, Koreas, Northern America and southeastern Australia (Parola and Raoult, 2001; Goethert and Telford III, 2003; Kawahara et al., 2006; Kim et al., 2003, 2006).

The tick *I. ricinus*, a vector for *Anaplasma*, has been reported in Northern Africa, in around Morocco and Tunisia (Parola and Raoult, 2001; Sarih et al., 2005).

The recent increasing popularity of recreation in forested areas or in the vicinity of forests raises the risk of contact with ectoparasites, for example with various species of ticks. Such grounds are typical biotopes of hard ticks. The end of the 20th century also brought about changes in the manner of animal breeding. In many industrialized countries the number of large farms is decreasing, where animals used to be raised in a closed system. Traditional methods of breeding animals are more popular and

natural pastures are more and more popular for both large ruminants (cattle) and small ones (sheep, goats). Apart from changes in people's lifestyles and farming techniques, we can also observe environmental changes. Environmental factors such as climate changes, deforestation, an increase in the roe deer population, or the introduction of new animal species which are potential pathogen reservoirs and hosts of ticks, favour an increase in the number of various tick populations (Wielinga et al., 2006).

It is also important to remember that in the present world both the humans and the animals bred by them are subject not only to indigenous domestic pathogens but also to imported ones. Hence the diagnosis becomes more and more difficult and laborious. It seems that the development of molecular techniques can facilitate the identification of these diseases, and information on the occurrence of ticks and pathogens transmitted by them will help veterinary services diagnose diseased farm animals correctly.

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