

Overview: Ticks as vectors of pathogens that cause disease in humans and animals

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1. ABSTRACT

Ticks (Acari: Ixodidae) transmit a wide variety of pathogens to vertebrates including viruses, bacteria, protozoa and helminthes. Tick-borne pathogens are believed to be responsible for more than 100,000 cases of illness in humans throughout the world. Ticks are considered to be second worldwide to mosquitoes as vectors of human diseases, but they are the most important vectors of disease-causing pathogens in domestic and wild animals. Infection and development of pathogens in both tick and vertebrate hosts are mediated by molecular mechanisms at the tick-pathogen interface. These mechanisms, involving traits of both ticks and pathogens, include the evolution of common and species-specific characteristics. The molecular characterization of the tick-pathogen interface is rapidly advancing and providing new avenues for the development of novel control strategies for both tick infestations and their associated pathogens.

2. INTRODUCTION: TICKS AND THE PATHOGENS THEY TRANSMIT

Molecular biology has made it possible to determine many of the genes that control the biological functions of numerous modern organisms, including blood feeding arthropods and the disease-causing pathogens that they transmit to humans and animals. Consequently, a substantial literature on these subjects has accumulated, including molecular studies of ticks and the disease agents that they transmit. This book compiles in a single comprehensive review the rapidly accumulating new knowledge of tick genomics and the molecular basis for many of the tick's biological processes, e.g., reproduction, salivation and innate immunity. In addition, it also reviews the latest findings on the molecular basis of the interactions between the tick vectors and their pathogens, e.g., tick-borne viruses, piroplasms, rickettsiae and borreliae. Finally, it brings together the new and often novel strategies for exploiting these findings to control ticks and tick-borne pathogens.

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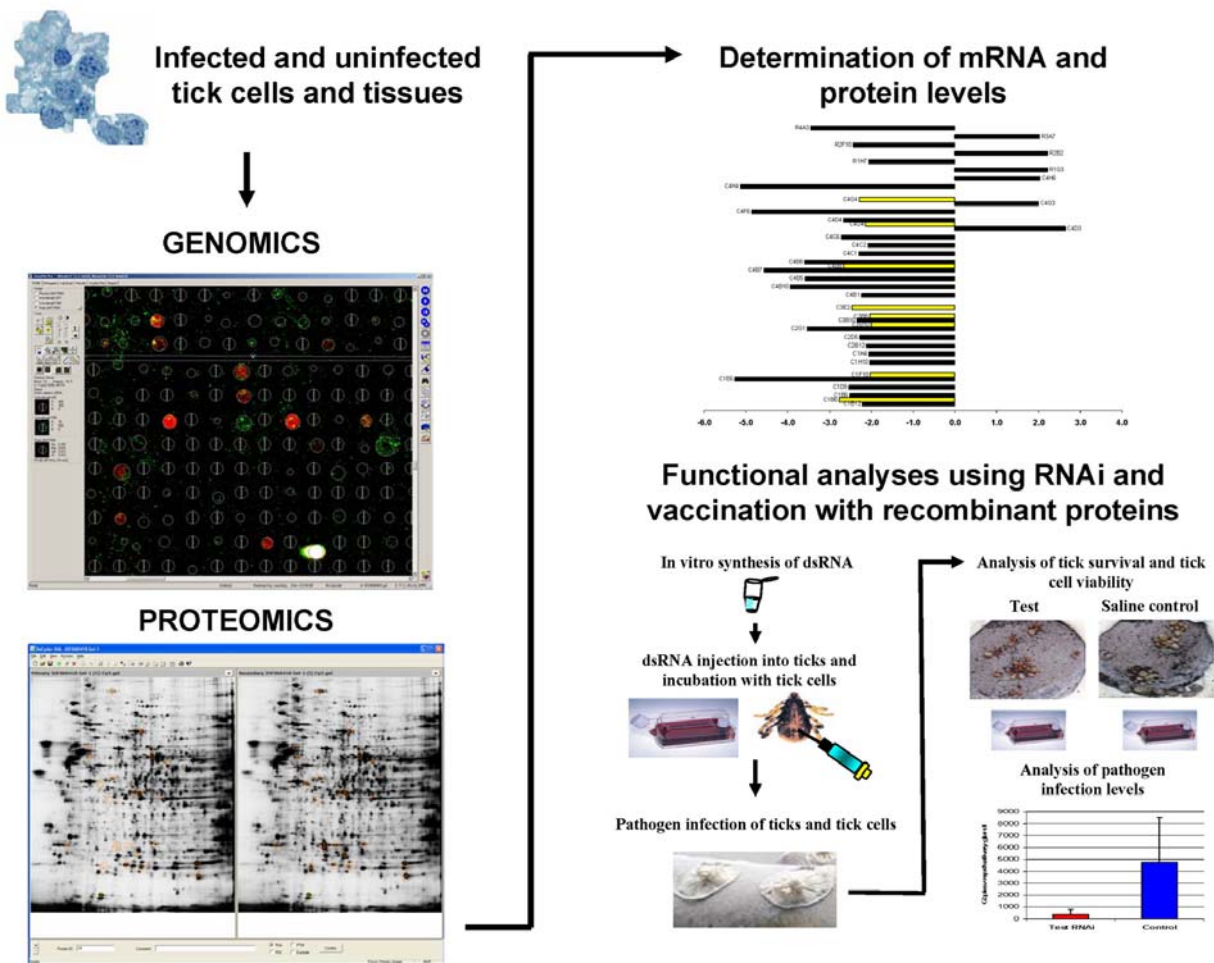


Figure 1. Systems biology approaches for the study tick-pathogen interface and development of vaccines for the control of tick infestations and the transmission of tick-borne pathogens.

A current and comprehensive listing of pathogens, their respective tick hosts and geographical distributions is presented in Table 1. This compilation is based on current literature, but readers are encouraged to consult the rapidly expanding literature which has been greatly impacted by molecular biology, especially concerning molecular methods for the identification of these pathogens and the definition of their relationship with ticks. The general name of the disease, which is often variable, has been included in the table when the name has been widely accepted. For viruses, only those with a demonstrated incidence in human and/or animal health have been included; disease names are rarely listed, except in limited cases where adequate clinical information was available. Proven vector species are included and complementary data, based on experimental evidence, is provided on the vector status of other tick species. Several *Rickettsia*, described recently in ticks or tick feeding lesions, are not included in the table, but rather are proposed as “*candidatus*” pending further information. In addition, changes in the taxonomic status of some pathogens are included that have been proposed or used in the scientific literature in recent years. For tick species, we

have incorporated the most widely accepted name following the review by Nava *et al.* (58). However, notes on the systematic status of selected pathogens have been provided. Finally, Table 1 should be considered a “work in progress” because new descriptions of tick-borne pathogens will undoubtedly continued to be reported and characterized.

3. MOLECULAR BIOLOGY OF TICKS AND THE TICK-PATHOGEN INTERFACE

The impact of molecular studies in conjunction with conventional taxonomy on tick systematics and evolution are discussed by Nava *et al.* (58). These studies are important toward understanding the evolution of ticks, tick-pathogen interactions, and the validity of tick nomenclature in relation to latest worldwide list of ticks.

Climatic and environmental conditions impact ticks and tick-pathogen interactions (59). Characterization of the climatic niche used by various tick species, in combination with the modeling of tick-host-pathogen interactions, will be fundamentally important toward our

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Table 1. Description of tick-borne pathogens, vectors, geographical distribution and host species affected

Pathogen	Disease	Tick vectors	Geographical distribution	Host species affected	References
Protozoa: genus <i>Babesia</i>					
<i>B. beliceri</i>		<i>Hyalomma</i> spp.	Russia	Cattle	1
<i>B. bigemina</i>	Cattle babesiosis	<i>Boophilus</i> spp., <i>Rhipicephalus</i> spp.	Africa, America, Asia, Australia	Cattle, buffalo	1
<i>B. bovis</i>	Cattle babesiosis	<i>Boophilus</i> spp.	Africa, America, Asia, Australia	Cattle, buffalo	1
<i>B. major</i>	Cattle babesiosis	<i>Haemaphysalis</i> spp.	Europe	Cattle	1
<i>B. ovata</i>		<i>Haemaphysalis</i> spp.	Asia	Cattle	1
<i>B. occultans</i> ¹		<i>Hyalomma</i> spp.	Africa	Cattle	1
<i>B. divergens</i>	Cattle babesiosis	<i>Ixodes</i> spp.	Europe	Cattle, Human ²	1
<i>B. microti</i> ³		<i>Ixodes scapularis</i>	USA, Canada	Human, rodents	1
<i>B. canis</i>	Dog babesiosis	<i>Rhipicephalus sanguineus</i> , <i>Dermacentor reticulatus</i> , <i>D. marginatus</i>	Tropical and semitropical worldwide	Dogs	1, 2, 3
<i>B. vogeli</i>	Dog babesiosis	<i>Rhipicephalus sanguineus</i>	Tropical and semitropical worldwide	Dogs	1
<i>B. rossi</i>	Dog babesiosis	<i>Haemaphysalis leachi</i>	Southern Africa	Dogs	1, 2
<i>B. gibsoni</i>	Dog babesiosis	<i>Haemaphysalis bispinosa</i> , <i>H. longicornis</i> , <i>Rhipicephalus sanguineus</i>	Africa, Asia, USA, southern Europe	Dogs	1, 2, 3
<i>B. ovis</i>	Sheep babesiosis	<i>Rhipicephalus bursa</i> , <i>R. turanicus</i>	Africa, Asia, Europe	Sheep	1
<i>B. motasi</i>	Sheep babesiosis	<i>Haemaphysalis</i> spp.	Africa, Asia, Europe	Sheep	1
<i>B. caballi</i>	Horse babesiosis	<i>Dermacentor</i> spp., <i>Rhipicephalus evertsi evertsi</i>	Africa, America, Asia, Europe	Horses, mules, donkeys	1, 3
<i>B. felis</i>		Unknown	Africa	Cats	1
<i>B. bicornis</i>		Unknown	Southern Africa	Black rhinoceros (<i>Diceros bicornis</i>)	5
<i>B. odocoilei</i>		<i>Ixodes scapularis</i>	America	Cervidae and wild Bovidae	6
Protozoa: genus <i>Theileria</i>					
<i>T. annulata</i>	Tropical theileriosis	<i>Hyalomma</i> spp.	Eurasia, Africa, Central Asia	Cattle, Camels	7
<i>T. orientalis</i>		<i>Haemaphysalis</i> spp.	Asia	Cattle, Asian buffalo	8
<i>T. parva</i>	East Coast Fever	<i>Rhipicephalus appendiculatus</i>	Africa	Cattle	9
<i>T. lawrencei</i>	Corridor disease	<i>Rhipicephalus zambeziensis</i>	Africa	Cattle	10
<i>T. velifera</i>	Apathogenic	<i>Amblyomma</i> spp.	Africa	Cattle, African buffalo	11
<i>T. buffeli</i>	Apathogenic	<i>Haemaphysalis</i> spp.	Asia	Cattle, Asian buffalo	11
<i>T. mutans</i>	Benign Theileriosis	<i>Amblyomma hebraeum</i> , <i>A. lepidum</i> , <i>A. variegatum</i> , <i>A. cohaerens</i> , <i>A. gemma</i>	Africa	Cattle	3, 10
<i>T. taurotragi</i>	Benign Theileriosis	<i>Rhipicephalus appendiculatus</i> , <i>R. pulchellus</i> , <i>R. zambeziensis</i>	Africa	Cattle	8
<i>T. ovis</i>	Sheep theileriosis	<i>Hyalomma</i> spp., <i>Rhipicephalus bursa</i>	Africa, Asia	Sheep	12
<i>T. lestoquardi</i> ⁴	Sheep theileriosis	<i>Hyalomma</i> spp.	Mediterranean region, Asia, Middle East, India	Sheep, Goats	11
<i>T. separata</i>	Non-pathogenic	<i>Rhipicephalus evertsi</i>	Sheep, goats	Africa	11
<i>T. bicornis</i>	Not named	Unknown	Black rhinoceros (<i>Diceros bicornis</i>)	Southern Africa	5
<i>T. equi</i>	Equine biliary fever	<i>Dermacentor</i> spp., <i>Rhipicephalus</i> spp., <i>Hyalomma</i> spp., <i>Boophilus</i> spp.	Horses, mules, donkeys	Southern Europe, Africa, Asia	3, 11
<i>T. cervi</i>	Non-pathogenic	<i>Amblyomma americanum</i>	White-tailed deer (<i>Odocoileus virginianus</i>)	Nearctic	76
Protozoa: genus <i>Hepatozoon</i>					
<i>H. canis</i>	Hepatozoonosis	<i>Rhipicephalus sanguineus</i> , <i>Haemaphysalis longicornis</i>	Southern Europe, Middle East, Far East, Africa	Dogs	13
<i>H. americanum</i>	Hepatozoonosis	<i>Amblyomma maculatum</i>	Southern USA	Dogs	13
Protozoa: genus <i>Cytauxzoon</i>					
<i>Cytauxzoon felis</i>	Cytauxzoonosis ⁵	<i>Dermacentor variabilis</i>	USA, Brazil	Domestic cats and wild felids	15, 16
Bacteria: genus <i>Aegyptianella</i>					
<i>A. pullorum</i> ⁶	Aegyptianellosis	<i>Argas walkerae</i> , <i>A. persicus</i> , <i>A. reflexus</i>	Africa, southern Europe, Middle Asia, Indian subcontinent	Domestic poultry	3, 17
Bacteria: genus <i>Rickettsia</i>					
<i>R. rickettsii</i>	Rocky Mountain spotted fever	<i>Dermacentor andersoni</i> , <i>D. variabilis</i> , <i>Amblyomma cajennense</i> , <i>A. aureolatum</i> , <i>Rhipicephalus sanguineus</i>	Americas	Human, dog	18, 19, 20

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<i>R. amblyommii</i>	Spotted fever rickettsiae group; no disease name	<i>Amblyomma americanum</i> , <i>A. neumanni</i> , <i>A. cajennense</i> , <i>A. coelebs</i>	Americas	Human	77-79
<i>R. conorii conorii</i>	Mediterranean spotted fever	<i>Rhipicephalus sanguineus</i>	Europe, Africa, Asia	Human, dog	18, 19, 20
<i>R. conorii israelensis</i>	Israeli spotted fever	<i>Rhipicephalus sanguineus</i>	Israel	Human	19, 20
<i>R. conorii caspia</i>	Astrakhan fever	<i>Rhipicephalus sanguineus</i> , <i>R. pumilio</i>	Africa, Asia	Human	19, 20
<i>R. conorii indica</i>	Indian tick typhus	<i>Rhipicephalus sanguineus</i>	India	Human	23
<i>R. sibirica sibirica</i>	Siberian or North Asian tick typhus	<i>Dermacentor nuttalli</i> , <i>D. marginatus</i> , <i>D. silvarum</i> , <i>D. sinicus</i> , <i>Haemaphysalis concinna</i>	Asia	Human	18, 19, 20
<i>R. sibirica mongolotimonae</i>	Unnamed	<i>Hyalomma asiaticum</i> , <i>H. truncatum</i> , <i>H. anatolicum excavatum</i>	Africa, China, France	Human	18, 19, 20
<i>R. australis</i>	Queensland tick typhus	<i>Ixodes holocyclus</i> , <i>I. tasmani</i>	Australia	Human	18, 19, 20
<i>R. japonica</i>	Oriental or Japanese spotted fever	<i>Ixodes ovatus</i> , <i>Dermacentor taiwanensis</i> , <i>Haemaphysalis longicornis</i> , <i>H. flava</i>	Japan	Human	18, 19, 20
<i>R. africae</i>	African tick-bite fever	<i>Amblyomma hebraeum</i> , <i>A. variegatum</i>	Africa, Reunion Island, West Indies	Human	18, 19, 20
<i>R. honei</i>	Flinders island spotted fever	<i>Bothriocroton hydrosauri</i> , <i>Amblyomma cajennense</i> , <i>Ixodes granulatus</i>	Australia, USA, Thailand	Human	18, 19, 20
<i>R. slovacae</i>	TIBOLA, DEBONEL ⁷	<i>Dermacentor marginatus</i> , <i>D. reticulatus</i>	Europe, Asia	Human	18, 19, 20
<i>R. helvetica</i>	Pathogenicity suspected in humans	<i>Ixodes ricinus</i>	Europe	Human	24
<i>R. heilongjiangensis</i>	Unnamed	<i>Dermacentor silvarum</i>	China	Human	23
<i>R. aeschlimannii</i>	Unnamed	<i>Hyalomma marginatum marginatum</i> , <i>H. m. rufipes</i> , <i>Rhipicephalus appendiculatus</i>	Europe, Africa	Human	18, 19, 20
<i>R. parkeri</i>	Unnamed	<i>Amblyomma maculatum</i> , <i>A. triste</i> , <i>A. dubitatum</i>	USA, Uruguay, Brazil	Human	19, 23, 25
<i>R. massiliae</i>	Unnamed	<i>Rhipicephalus sanguineus</i> , <i>R. turanicus</i> , <i>R. muhsamae</i> , <i>R. lunulatus</i> , <i>R. sulcatus</i>	Europe, Asia, Argentina, USA	Human	19, 26, 27, 28
<i>R. marmionii</i>	Australian spotted fever	<i>Haemaphysalis novaeguineae</i> , <i>Ixodes holocyclus</i>	Australia	Human	19, 23
<i>R. monacensis</i>	Unnamed	<i>Ixodes ricinus</i>	Europe	Human	19, 29
Bacteria, <i>Ehrlichia</i> genus					
<i>E. chaffeensis</i>	Human monocytic ehrlichiosis	<i>Amblyomma americanum</i> , <i>Dermacentor variabilis</i>	USA	Human and various mammals	2, 20
<i>E. ewingii</i>	Canine granulocytic ehrlichiosis, Human ehrlichiosis	<i>Amblyomma americanum</i>	USA	Human, dogs	2, 18
<i>E. ruminantium</i>	Heartwater	<i>Amblyomma hebraeum</i> , <i>A. astrion</i> , <i>A. cohaerens</i> , <i>A. gemma</i> , <i>A. marmoreum</i> , <i>A. lepidum</i> , <i>A. pomposum</i> , <i>A. variegatum</i> , <i>A. americanum</i> ⁸	Africa, Caribbean	Mainly cattle ⁹	3, 30
<i>E. canis</i>	Canine ehrlichiosis	<i>Rhipicephalus sanguineus</i>	Southern USA, southern Europe, Africa, Middle East, eastern Asia	Dogs	2
Bacteria, <i>Anaplasma</i> , <i>Francisella</i> and <i>Coxiella</i> genera					
<i>Anaplasma phagocytophilum</i>	Human granulocytic anaplasmosis	<i>Ixodes scapularis</i> , <i>I. pacificus</i> , <i>I. ricinus</i> , <i>I. hexagonus</i>	USA, Europe	Human and various mammals	20, 33, 34
<i>A. marginale</i>	Bovine Anaplasmosis	Various	Worldwide	Cattle	35
<i>A. centrale</i>	Bovine Anaplasmosis	Various	Worldwide	Cattle	35
<i>A. ovis</i>	Ovine Anaplasmosis	Various	Worldwide	Sheep	35
<i>A. platys</i>	Canine ehrlichiosis	<i>Rhipicephalus sanguineus</i>		Dog	36
<i>F. tularensis</i>	Tularemia	Various	Eurasia, Nearctic	Human and various mammals	18
<i>C. burnetii</i>	Q fever	Various ¹⁰	Worldwide	Human and various mammals	18
Bacteria, <i>Borrelia</i> ¹¹ genus					
<i>B. burgdorferi</i>	Lyme disease	<i>Ixodes pacificus</i> , <i>I. persulcatus</i> , <i>I. ricinus</i> , <i>I. scapularis</i>	USA, Canada, Europe, Asia, northern Africa	Human	18, 33
<i>B. garinii</i>	Lyme disease	<i>Ixodes persulcatus</i> , <i>I. ricinus</i>	Europe, Asia, northern Africa	Human	33
<i>B. afzelii</i>	Lyme disease	<i>Ixodes persulcatus</i> , <i>I. ricinus</i>	Europe, Asia, northern Africa	Human	33
<i>B. valaisiana</i>	Lyme disease	<i>Ixodes ricinus</i>	Europe, Asia	Human	18, 33
<i>B. lusitaniae</i>		<i>Ixodes ricinus</i>	Europe		33
<i>B. spielmani</i>	Lyme disease	<i>Ixodes ricinus</i>	Europe	Human	80

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<i>B. japonica</i>	Lyme disease	<i>Ixodes ovatus</i>	Japan	Human	33
<i>B. lonestari</i>		<i>Amblyomma americanum</i>	USA	Human	37
<i>B. theileri</i>	Bovine borreliosis	<i>Boophilus</i> spp., <i>Rhipicephalus evertsi</i>	Africa, Central and South America, Australia	Cattle	38
<i>B. turcica</i>		<i>Hyalomma aegyptium</i>	Parts of Turkey ¹²		39
<i>B. miyamotoi</i>		<i>Ixodes persulcatus</i>	Asia		40
<i>B. hermsii</i>	New World tick-borne relapsing fever	<i>Ornithodoros hermsi</i>	USA, Canada	Human	18, 33, 38
<i>B. turicatae</i>	New World tick-borne relapsing fever	<i>Ornithodoros turicata</i>	USA, Mexico	Human	18, 33, 38
<i>B. parkeri</i>	New World tick-borne relapsing fever	<i>Ornithodoros parkeri</i>	USA	Human	18, 33, 38
<i>B. mazzottii</i>	New World tick-borne relapsing fever	<i>Ornithodoros talaje</i>	USA, Mexico		18, 38
<i>B. venezuelensis</i>	New World tick-borne relapsing fever	<i>Ornithodoros rudis</i>	Central and South America	Human	18, 33, 38
<i>B. duttonii</i>	Old World tick-borne relapsing fever	<i>Ornithodoros moubata</i>	Africa	Human	18, 33, 38
<i>B. crocidurae</i>	Old World tick-borne relapsing fever	<i>Ornithodoros erraticus</i> ¹³	Europe, Africa	Human	18, 33, 38
<i>B. persica</i>	Persian relapsing fever	<i>Ornithodoros tholozani</i>	Asia	Human	18, 33, 38
<i>B. hispanica</i>	Old World tick-borne relapsing fever	<i>Ornithodoros erraticus</i>	Spain, Portugal	Human	18, 33, 38
<i>B. latyschevii</i>	Old World tick-borne relapsing fever	<i>Ornithodoros tartakovskyi</i>	Iran, Central Asia	Human	18, 33, 38
<i>B. caucasica</i>	Old World tick-borne relapsing fever	<i>Ornithodoros aspersus</i>	Asia (Caucasus and Iraq)	Human	18, 33, 38
<i>B. graingeri</i>		<i>Ornithodoros graingeri</i>	Africa	Human	18
<i>B. anserina</i>	Avian borreliosis	<i>Argas</i> spp.	Worldwide	Birds	38
<i>B. tillae</i>		<i>Ornithodoros zumpti</i>	Africa	Human	42
<i>B. coraciae</i>	Bovine epizootic abortion	<i>Ornithodoros coriaceus</i>	USA	Cattle	33, 38
<i>B. parkeri</i>		<i>Ornithodoros parkeri</i>	USA	Human	42
Bacteria, genus <i>Dermatophilus</i>					
<i>D. congolensis</i>	Dermatophilosis	<i>Amblyomma variegatum</i> ¹⁴	Africa	Ruminants	3
Viruses					
Bunyaviridae, Nairovirus	Crimean-Congo Hemorrhagic Fever virus	<i>Hyalomma marginatum</i> , <i>Hy. a. anaticum</i> , <i>Hy. truncatum</i> , <i>Amblyomma variegatum</i> , <i>Haemaphysalis punctata</i> , <i>Ixodes ricinus</i> , <i>Dermacentor</i> spp., <i>Rhipicephalus</i> spp.	Africa, Asia, Europe	Human	3, 33, 43, 44
Bunyaviridae, Nairovirus	Nairobi Sheep Disease	<i>Rhipicephalus appendiculatus</i> , <i>R. pulchelus</i> , <i>Amblyomma variegatum</i>	Kenya, Uganda, Rwanda, Tanzania, Somalia	Sheep, goats	45
Bunyaviridae, Nairovirus	Soldado virus	<i>Ornithodoros maritimus</i> , <i>O. capensis</i> , <i>O. denmarki</i>	Trinidad, Hawaii, Seychelles, Ethiopia, South Africa, Morocco, France, United Kingdom	Human	33, 44
Bunyaviridae, ungrouped	Issyk-Kul fever virus	<i>Argas vespertilionis</i> , <i>A. pusillus</i> , <i>Ixodes vespertilionis</i> ¹⁵	Tadzhikistan, Kyrgyzstan, Turkmenistan	Bats, human	46
Reoviridae, Orbivirus	Eyach virus	<i>Ixodes ricinus</i> , <i>I. ventalloi</i>	Europe	Human	44
Reoviridae, Orbivirus	Colorado tick fever	<i>Dermacentor andersoni</i> , <i>D. occidentalis</i> , <i>D. albipictus</i>	Nearctic	Human	44
Reoviridae, Orbivirus	Mono Lake virus	<i>Argas monolakensis</i> , <i>A. cooleyi</i>	USA	Human	33, 47, 48, 49
Orthomyxoviridae, Thogotovirus		<i>Amblyomma</i> spp., <i>Boophilus</i> spp., <i>Hyalomma</i> spp., <i>Rhipicephalus</i> spp.	Central and East Africa, southern Europe	Sheep, human	44, 50
Flaviviridae, Flavivirus	Omsk Hemorrhagic fever	<i>Dermacentor reticulatus</i>	Omsk and Novosibirsk regions of western Siberia (Russia)	Human	33, 44
Flaviviridae, Flavivirus	Tick Borne Encephalitis	<i>Ixodes ricinus</i> , <i>I. persulcatus</i> , <i>Haemaphysalis concinna</i> , <i>H. punctata</i>	Europe, Asia	Human	33, 44
Flaviviridae, Flavivirus	Langat virus	<i>Ixodes granulatus</i> , <i>Haemaphysalis papuana</i>	Malaysia	Wild rodents	51
Flaviviridae, Flavivirus	Louping ill	<i>Ixodes ricinus</i>	United Kingdom	Sheep	52
Flaviviridae, Flavivirus	Powassan Encephalitis	<i>Ixodes cookie</i> , <i>I. marxi</i> , <i>I. scapularis</i> , <i>Dermacentor andersoni</i>	Canada, USA, Russia	Human, rodents	3, 44, 53
Flaviviridae, Flavivirus	Kyasanur Forest disease	<i>Haemaphysalis</i> spp. (mainly <i>H. spinigera</i> and <i>H. turturis</i>)	Forested areas of the Kyasanur district in India	Human, monkeys	54
Asfaviridae, Asfavirus	African Swine Fever	<i>Ornithodoros moubata</i> , <i>O. erraticus</i> , <i>O. turicata</i> , <i>O. coriaceus</i> , <i>O. puertoricensis</i> ¹⁶	Africa, southern Europe, Caribbean (outbreaks in Brazil, likely imported)	Domestic pigs	44, 55
Nematoda:					

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genus					
<i>Acanthocheilonema</i>					
<i>Acanthocheilonema viteae</i> ¹⁷		<i>Ornithodoros tartakovskyi</i>	Asia	Rodents	17
<i>Acanthocheilonema dracunculoides</i> ¹⁸		<i>Rhipicephalus sanguineus</i>		Dogs	57

No one has yet demonstrated that *B. occultans* is a different species from *B. beliceri* (1).² While *B. divergens* is primarily a species associated with cattle, it may cause disease in humans (4).³ Most probably this is not a *Babesia* sp. (1).⁴ A yet unnamed species has been reported from China transmitted by *Haemaphysalis qinghaiensis* and different from *T. lestoquardi* and *T. buffeli* (14).⁵ *Cytauxzoon manul* and another probably new species have been described in wild felids in Mongolia and Spain (21,22).⁶ An *Aegyptianella* found in wild turkeys (*Meleagris gallopavo*) in the USA and described as *A. pullorum* appears to be a different species, specific to turkeys.⁷ There is not a total agreement on the name of the disease. While some authors prefer to use Tick-Borne-Lymphadenitis, others prefer to use Dermacentor-Borne-Necrosis-Erythema-Lymphadenopathy.⁸ Other tick species endemic in the Nearctic and Neotropics have been involved in the laboratory transmission of *E. ruminatum*. Even in the absence of the main vectors in these regions, it has been claimed that local ticks could spread the infection (31).⁹ Many wild ruminants may act as hosts of *E. ruminatum* (32).¹⁰ Transmission of *C. burnetti* by tick bite is very rare.¹¹ Species of *Borrelia* are herein arranged as transmitted by hard ticks or soft ticks.¹² This species has not been identified in humans but only in the tick *Hyalomma aegyptium*. However, Vatansver et al. (41) reported the high prevalence of *H. aegyptium* immature on humans in Istanbul, Turkey. Therefore, the possibility of a transmission to humans should be considered when analyzing cases of Borreliosis in this area.¹³ It is not yet clear whether *O. erraticus* is the same species as *O. sonrai* and *O. marocanus*. The current point of view is that *O. erraticus* and *O. sonrai* are valid species, while *O. marocanus* would be a junior synonym of *O. erraticus*.¹⁴ Associated with severe dermatophilosis of ruminants.¹⁵ This is a primary disease of bats. However, several strains of the virus have been isolated from bat-biting ticks and these ticks have the ability to feed on humans.¹⁶ The role in transmission by *O. turicata*, *O. coriaceus* and *O. puertoricensis* has been demonstrated under laboratory conditions (56)¹⁷ Some authors prefer to place these species under the genus *Dipetalonema*.¹⁸ Idem to "17"

ability to understand the impact of climate change on tick range and the epidemiology of tick-borne diseases (59). The impact of global climate change on the distribution of vector-borne diseases is increasingly acknowledged, and this emerging area of research will be receiving considerable attention in the future (60-62).

Developing an understanding of the molecular basis for tick reproduction (63) and immunity (64) is critically important for defining tick-host and tick-pathogen interactions. Recent advances in these areas have demonstrated the complexity of these systems and are providing novel approaches for the manipulation and study of tick biology. The diversity of molecular mechanisms that govern tick-pathogen and tick-host interactions are illustrated in recent research on the interface between ticks and selected pathogens, including *Anaplasma* (65), *Ehrlichia* (66), *Borrelia* (67), viruses (68), and piroplasms (69), as well as at the interface between ticks and vertebrate hosts (70). These reviews cover current knowledge on the most important groups of pathogens transmitted by ticks and emphasize the important role of tick salivary glands in tick feeding, pathogen infection and transmission.

The sequence and annotation of the first tick genome, *Ixodes scapularis*, is rapidly progressing (71). The entire *I. scapularis* genome shotgun sequence data have now been generated and assembled, providing 3.8 fold coverage of the genome (GenBank project accession number ABJB010000000; VectorBase designation IscaW1). The knowledge gained from tick genomics (72), molecular biology of tick feeding, reproduction and immune response (63, 64, 70) and tick-pathogen (65-69) and tick-host (70) interactions will markedly contribute to our understanding of the tick-pathogen interface and may ultimately be expected to lead to development of new strategies for control of ticks and tick-borne pathogens (73, 74).

4. SUMMARY AND PERSPECTIVE

According to Dennis and Peisman (75), tick-borne pathogens are believed to be responsible for more than 100,000 cases of illness in humans throughout the world and are the most important vectors of disease-causing pathogens in domestic and wild animals. The control of tick infestations and the pathogens they transmit constitute a priority to improve human and animal health worldwide. Research on the characterization of the molecular events that mediate vital tick functions such as feeding, reproduction, immunity and the tick-pathogen interface, is still in its infancy. However, as summarized in the following chapters (58, 59, 63-70, 72-74), the recent application of molecular and systems biology approaches will contribute to our understanding of the biology of the tick-pathogen interface. Collectively, this information may be useful for the development of new and novel control methods for both tick infestations and the prevention of transmission of tick-borne pathogens (Figure 1).

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