# **Tick-Borne Infectious Diseases in China, 2003–2023**

Yinshan Zhu<sup>1,2</sup>; Tianxiang Jiang<sup>1,2</sup>; Zhishan Sun<sup>1,2</sup>; Jingxian Yin<sup>1,2</sup>; Hanqing Zhao<sup>1,2</sup>; Xiaonong Zhou<sup>1,#</sup>; Kokouvi Kassegne<sup>1,#</sup>

### **ABSTRACT**

Tick-borne diseases (TBDs) pose a substantial threat to global public health, including China. This article systematically reviews the epidemiology of TBDs in Chinese mainland between 2003 and 2023. We retrieved and analyzed the incidence and distribution of notable bacterial, viral, and parasitic TBDs in China, and discussed their clinical characteristics, diagnostic approaches, and vectors. Severe fever with thrombocytopenia syndrome is the leading TBD in China, with cases rapidly increasing since its first report in 2009, and a high case fatality rate of 5%-33% in patients with severe disease. Other TBDs reported in China include spotted fever group rickettsiosis, human granulocytic anaplasmosis, Lyme borreliosis, babesiosis, and tick-borne encephalitis. Infections with emerging tick-borne pathogens, such as Yezo virus and Beiji nairovirus, have appeared very recently. Although evidence of human infection with Crimean-Congo hemorrhagic fever virus, Francisella tularensis, and Bartonella is currently unavailable, continuous monitoring of these pathogens remains necessary. This study enhances our understanding of TBDs, provides guidance for public health decisionmaking regarding important bacterial, viral, and parasitic TBDs, and offers new opportunities for TBD research in China.

Tick-borne diseases (hereafter referred to as TBDs), which are transmitted through ticks as vectors, are characterized by their high infectivity and wide transmission range. Recent years have witnessed a notable increase in the global incidence of these diseases (1–2), along with rising cases of coinfections with multiple tick-borne pathogens (3–4). These diseases pose serious health risks and economic burdens to both humans and animals worldwide, underscoring the critical importance of research in this field. Ongoing processes of globalization, climate change,

and increased human-animal interactions have further accelerated the global spread of TBDs, leading to the emergence of new diseases (5–6). China's complex terrain, combined with diverse tick species and pathogens, presents a major public health challenge. Urbanization and afforestation projects complicate the distribution of ticks and pathogens, creating significant obstacles for disease prevention and control efforts in the country (7). Despite the diversity and complex distribution of tick species in China, the health threats posed by TBDs are often underestimated, hampering current prevention and control initiatives. By analyzing research literature, we can develop a better understanding of the characteristics of various TBDs in China to improve preventive measures.

### **METHODS**

# **Search Strategy**

A systematic literature review was conducted using Embase, PubMed, Scopus, and the Web of Science Core Collection. The search terms included 'Tickborne diseases' OR 'Lyme borreliosis' OR 'Tick-borne encephalitis' OR 'Q fever' OR 'Babesiosis' OR 'Crimean-Congo haemorrhagic fever' OR 'African swine fever' OR 'Tularemia' OR 'Kyasanur forest disease' OR 'Colorado tick fever' OR 'Ehrlichiosis' OR 'Anaplasmosis' OR 'Rocky Mountain spotted fever' OR 'Southern tick-associated rash illness' OR 'Tickborne relapsing fever' OR 'Pacific coast tick fever' OR 'Omsk haemorrhagic fever' OR 'Rickettsiosis' OR 'Spotted fever group rickettsiosis' OR 'Novel bunyavirus infection' OR 'Piroplasmosis' OR 'Oriental spotted fever' OR 'Relapsing fever borrelioses' OR 'Powassan disease' OR 'Powassan encephalitis' OR 'Imported tick-borne spotted fever' OR 'North-Asia spotted fever' OR 'African tick bite fever'. As China CDC Weekly is an international peer-reviewed journal that ensures research reproducibility, this study included only English-language publications. The review was limited to research articles published between 2003 and 2023, excluding reviews, case

reports, commentaries, interviews, letters, and editorials. Mite-, flea-, and louse-borne diseases, such as Tsutsugamushi, murine rickettsiosis, and infections caused by *Rickettsia felis*, *Rickettsia akari*, and *Rickettsia prowazekii*, were excluded. After removing duplicate articles using EndNote and Excel, only studies related to TBDs were included.

# **Data Processing**

We systematically processed data on TBDs from the articles that met the inclusion criteria. These data include geographical origins of cases, incidence and distribution, clinical characteristics, diagnostic methods, and tick vectors.

#### **RESULTS**

# **Literature Screening**

A total of 3,483 articles were initially retrieved from Embase, PubMed, the Web of Science Core Collection, and Scopus. After screening, 67 relevant articles specifically related to TBDs were retained for

inclusion in the study. The detailed screening process is illustrated in Figure 1.

Analysis of these 67 articles revealed temporal trends in TBD cases (Figure 2A). The number of patients with severe fever with thrombocytopenia syndrome (SFTS) gradually increased in 2020, peaked in 2021, and then progressively declined in subsequent years. A summary of articles and cases related to various TBDs (Figure 2B) showed that 43% of the articles focused on SFTS, 18% on spotted fever group rickettsiosis (SFGR), and 14% on human granulocytic anaplasmosis (HGA). This finding indicates a shift in research attention toward TBDs in China, with SFTS attracting significant interest among researchers. Furthermore, SFTS accounted for the greatest number of cases, representing 66% (10,080/15,261) of the total TBD cases reported in the country. This underscores SFTS as the most prominent health concern among TBDs in China.

### **Distribution of TBDs in China**

We assessed the geographical distribution of TBD cases across Chinese mainland (Figure 2C). This

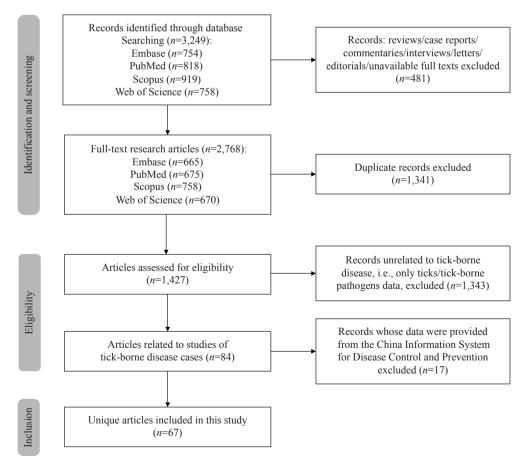


FIGURE 1. Flowchart illustrating the search and screening process for articles on tick-borne diseases in China.

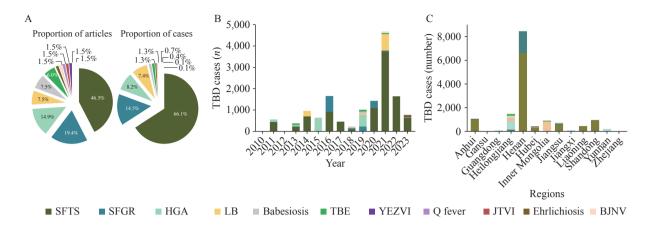


FIGURE 2. Distribution of articles and yearly cases of tick-borne diseases in China, 2003–2023. (A) The proportion of articles and cases related to tick-borne diseases; (B) Number of cases per tick-borne disease across the years; (C) Incidence of tick-borne diseases reported across Chinese mainland.

Abbreviation: SFTS=severe fever with thrombocytopenia syndrome; SFGR=spotted fever group rickettsiosis; HGA=human granulocytic anaplasmosis; LB=Lyme borreliosis; TBE=tick-borne encephalitis; YEZVI=Yezo virus infection; JTVI=Jingmen tick virus infection; BJNV=Beiji nairovirus.

analysis revealed a concentration of TBD cases primarily in provincial-level administrative divisions (PLADs) such as Henan, Heilongjiang, Anhui, Shandong, Inner Mongolia, and Jiangsu. These findings align with the known distribution of ticks and tick-borne pathogens (7). Additionally, SFTS alone accounted for 10,080 of the 15,261 total TBD cases reported in China, followed by SFGR and HGA, which accounted for 2,212 and 1,257 cases, respectively. Notably, 61.2% (6,611/10,800) of SFTS cases and 81.8% (1,810/2,212) of SFGR cases were reported in Henan Province, 37.9% (477/1,257) of HGA cases were reported in Heilongjiang Province, and 68.5% (771/1,126) of Lyme borreliosis (LB) cases and 74.9% (149/199) of tick-borne encephalitis (TBE) cases were reported in Inner Mongolia Autonomous

A comprehensive categorization of TBDs into viral, bacterial, and parasitic types is presented in Table 1. The clinical characteristics, diagnostic methods, and tick species for these diseases are summarized. In China, *Haemaphysalis longicornis*, *Ixodes persulcatus*, and *Haemaphysalis yeni* serve as the primary vectors for numerous tick-borne pathogens. This prevalence might be attributed to the favorable environmental conditions in China that are conducive to the proliferation of these particular tick species (7).

### **Major Tick-borne Diseases in China**

**SFTS** SFTS is a recently identified infectious disease that originated in China and is caused by a novel bunyavirus (65). In China, SFTSV has been detected

in Haemaphysalis longicornis ticks collected from livestock in areas where SFTS patients reside, establishing this species as the primary transmission vector (66). SFTS cases in China are primarily concentrated in Liaoning Province in northeastern China; Shandong Province along the eastern coast; Jiangsu and Zhejiang provinces in eastern China; and Henan, Hubei, and Anhui provinces in central China (7). According to this study, a total of 10,080 SFTS cases have been identified in China. The characteristic clinical manifestations include acute fever, fatigue, thrombocytopenia, leukopenia, and gastrointestinal symptoms such as nausea, vomiting, and diarrhea. In severe cases, multiple organ dysfunction may occur (10). Currently, no specific treatment or vaccine is available for this disease, making effective prevention measures crucial in endemic areas (66).

SFGR SFGR is a zoonosis caused by bacteria of the genus *Rickettsia* and is transmitted primarily by arthropod vectors, particularly ticks (67). This study identified 2,212 SFGR cases reported in China. The clinical presentation typically includes fever, rash, and other characteristic features such as eschars and lymphadenopathy (68). Early diagnosis and treatment are essential, with antibiotics commonly used to alleviate symptoms and prevent disease progression. Preventive measures include avoiding tick bites, practicing good personal hygiene, and maintaining clean living environments. Our analysis revealed that *Haemaphysalis yeni* and *H. longicornis* serve as the primary vectors for SFGR in China.

**HGA** HGA is an infectious bacterial disease caused

TABLE 1. Characteristics of TBDs reported in China, 2003–2023.

Category	TBD	Pathogen	Clinical symptom	Biochemical indicator	Identification method	Tick species	Citation
Viral diseases	SFTS	thrombocytopeni	nFever, anorexia, headache, muscular soreness, diarrhea, dizziness, ecchymosis, arthrodynia, central nervous system	Thrombocytopenia, leukocytopenia, elevated aminotransferase levels, elevated lactate dehydrogenase levels,	RT-PCR	H. longicornis	(8–41)
	TBE	Tick-borne encephalitis virus	symptoms Fever, headache, scentral nervous system symptoms	proteinuria, hematuresis Increased white blood cell count, neutrophilic leukocytosis, elevated aminotransferase levels	PCR	NA	
	Jingmen tick virus infection	Jingmen tick virus	Fever, headache, myodynia, pruritus, eschar, lymphadenectasis		RT-PCR	I. persulcatus	
	Yezo virus infection	Yezo virus	Fever, headache, dizziness, diminution of vision, chest distress, breathe hard arthrodynia	neutrophilic leukocytosis,	RT-PCR	I. persulcatus	
	Beiji nairovirus infection	Beiji nairovirus	Fever, headache, redness and swollen, pruritus, depression, coma, fatigue, myalgia, arthralgia, poor appetite, skin rashes, petechiae	Thrombocytosis, leukocytosis, elevated level of high sensitivity C-reactive protein, elevated aminotransferase levels, elevated lactate dehydrogenase levels	PCR I	NA	
Bacterial diseases	HGA	Anaplasma phagocytophilum	Fever, headache, dizziness, weakness, malaise, anorexia, nausea, chills, cough, abdominal pain, myalgia, ecchymosis, rash, lymphadenectasis	Leukocytopenia, thrombocytopenia, elevated aminotransferase levels, elevated lactate dehydrogenase levels	PCR, IFA	H. longicornis	(42–60)
	SFGR	Rickettsia rickettsii	Fever, headache, asitia, weakness, cough, dizziness, rash, eschar, lymphadenectasis	Thrombocytopenia, leukocytopenia, elevated aminotransferase levels	PCR, IFA	H. yeni,H. Iongicornis	
	LB	Borrelia burgdorferi	Frythema migrans, fever, headache, weak, naupathia, arthrodynia, asitia, lymphadenectasis, erythra, eschar	Elevated aminotransferase levels, thrombocytopenia, leukocytosis	PCR	NA	
	Ehrlichiosis	Ehrlichia spp.	Fever, rash, asthenia, anorexia, myalgia	Thrombocytopenia, leukocytopenia, elevated aminotransferase levels, increased C-reactive protein	PCR	H. longicornis	
	Q fever	Q fever Rickettsia	Fever, headache, muscular soreness, erythra, eschar		PCR	NA	
Parasitic diseases		Babesia spp.	Fever, headache, dizziness, naupathia, chills, arthralgia, lymphadenectasis, flu-like symptoms			NA	(61–64)

Abbreviation: TBD=tick-borne disease; SFTS=severe fever with thrombocytopenia syndrome; TBE=tick-borne encephalitis; BJNV=Beiji nairovirus; HGA=human granulocytic anaplasmosis; SFGR=spotted fever group rickettsiosis; LB=Lyme borreliosis; RT-PCR=reverse transcription polymerase chain reaction; PCR=polymerase chain reaction; IFA=immunofluorescence assay; H.=Haemaphysalis; I.=Ixodes; NA=no relevant results found.

by Anaplasma phagocytophilum (69). The primary transmission route is through tick bites, with blood transfusion representing another documented mode of

transmission (70–71). This study identified 1,257 HGA patients in China. Typical clinical manifestations include fever, headache, myalgia, fatigue, nausea,

vomiting, general malaise, and thrombocytopenia with associated bleeding abnormalities (42). HGA cases in China are primarily concentrated in the central and southeastern regions (72). The Infectious Diseases Society of America (IDSA) recommends empiric treatment with doxycycline for all patients with suspected HGA (73). In China, Haemaphysalis longicornis serves as the primary vector for HGA.

LB LB is a naturally occurring epidemic disease caused by *Borrelia burgdorferi* and is transmitted primarily by ticks, particularly those in the genus *Ixodes* (74). Although LB is the most commonly reported tick-borne illness in the United States, China has reported fewer cases, with a total of 1,126 documented in this study (7,75). The principal clinical manifestations include skin rash, erythema migrans, fever, fatigue, headache, and arthralgia. Most patients recover with effective treatment, and fatal outcomes are rare (76).

**Babesiosis** Babesiosis is a zoonotic parasitic disease caused by Babesia, a blood protozoan that infects red blood cells and is transmitted to humans and animals through tick bites (77,78). Over the past 20 years, more than 100 patients in Zhejiang, Yunnan, Guangxi, and other Chinese PLADs have been infected with Babesia microti, suggesting that B. microti from field mice may be the primary pathogen causing human babesiosis in China (79). Clinical manifestations include fever, headache, myalgia and arthralgia, cutaneous pain or pruritus, lymphadenopathy, abdominal pain, diarrhea or constipation, dyspnea, anemia, and neurological symptoms such as dizziness, nausea, and blurred vision (80). While typically mild to moderate in severity, babesiosis can lead to severe outcomes or even death in elderly immunocompromised patients (81).

TBE, also known as forest encephalitis, is an acute infectious disease of the central nervous system caused by the tick-borne encephalitis virus of the genus Flavivirus. It is transmitted primarily through the bite of ticks in the genus Ixodes (82). After entering the body via a tick bite, the virus replicates and migrates to draining lymph nodes before spreading to the bloodstream and crossing the blood-brain barrier, resulting in central nervous system damage (83). Between 2007 and 2018, 3,364 cases were reported in Chinese mainland, predominantly in forested areas of northeastern China (84). The principal symptoms include fever, headache, myalgia, fatigue, nausea, vomiting, diarrhea, rash, and, in severe cases, neurological manifestations such altered as

consciousness, seizures, and paralysis (85).

### **DISCUSSION**

Due to China's vast geographical expanse and complex natural environment, combined with the diversity of tick vector species, pathogen types, and intermediate hosts, there likely exist even more types of diseases (TBDs) currently tick-borne than documented. This presents significant challenges for TBD prevention and control efforts in China. Research priorities regarding TBDs in China have evolved over time. Historically, diseases such as Crimean-Congo hemorrhagic fever, African swine fever, Q fever, and Lyme disease were the primary focus (86). However, SFTS has received considerable attention in recent years. SFTS is an emerging tickborne infectious disease first reported in China in 2009 and subsequently identified as a novel bunyavirus in 2010 (87). SFTS is particularly lethal, with the elderly patients and those with underlying conditions being susceptible to multiple organ failure following infection (36,88). Currently, there is no specific antiviral treatment or approved vaccine for SFTSV (89). The number of SFTS cases increased gradually in 2020, peaked in 2021, and subsequently decreased each year thereafter. This trend may be attributed to the focus on COVID-19 prevention and control efforts, which diverted attention from surveillance and management. Additionally, the significant fluctuation in TBD incidence in recent years indicates that monitoring and control measures should be strengthened. The clinical features of TBDs underscore the urgency of addressing these diseases. Furthermore, pathogens of many TBDs, including bacteria and viruses, are transmitted primarily by H. longicornis, a tick species widely distributed throughout China (90). Recent research has also shown that Rhipicephalus sanguineus may serve as a vector for SFTSV (91). TBD cases are concentrated in developed agricultural or forestry areas such as Henan, Heilongjiang, and Inner Mongolia. The prevalence of TBDs in these regions is primarily due to favorable environmental factors, including abundant vegetation and livestock resources. These conditions not only provide suitable habitats for the complete tick life cycle from eggs to adults but also promote extensive agropastoral activities within local communities, further exacerbating disease spread (7,92). These

findings suggest that urgent measures are needed to control ticks and tick-borne pathogens.

A limitation of this study is that the literature search was restricted to English-language articles, potentially overlooking studies published in other languages.

### **CONCLUSION**

In summary, due to the rich diversity and widespread distribution of tick species in China, TBDs have become a significant threat to public health. Research on these infections, particularly SFTS, continues to face notable challenges. We urge the use of an integrated One Health approach, at the human-animal-environment interface, to enhance prevention and control measures for TBDs, particularly in regions with developed agriculture and forestry, as well as agropastoral activities.

Conflicts of interest: No conflicts of interest.

doi: 10.46234/ccdcw2025.150

\* Corresponding authors: Kokouvi Kassegne, ephremk@outmail.fr; Xiaonong Zhou, xiao-nong.zhou@sjtu.edu.cn.

Copyright © 2025 by Chinese Center for Disease Control and Prevention. All content is distributed under a Creative Commons Attribution Non Commercial License 4.0 (CC BY-NC).

Submitted: March 25, 2025 Accepted: June 17, 2025 Issued: June 27, 2025

### **REFERENCES**

- Boulanger N, Boyer P, Talagrand-Reboul E, Hansmann Y. Ticks and tick-borne diseases. Med Mal Infect 2019;49(2):87 – 97. https://doi. org/10.1016/j.medmal.2019.01.007.
- Nowak M, Cieniuch S, Stańczak J, Siuda K. Detection of Anaplasma phagocytophilum in Amblyomma flavomaculatum ticks (Acari: Ixodidae) collected from lizard Varanus exanthematicus imported to Poland. Exp Appl Acarol 2010;51(4):363 – 71. https://doi.org/10.1007/s10493-009-9332-5
- Cutler SJ, Vayssier-Taussat M, Estrada-Peña A, Potkonjak A, Mihalca AD, Zeller H. Tick-borne diseases and co-infection: current considerations. Ticks Tick Borne Dis 2021;12(1):101607. https://doi. org/10.1016/j.ttbdis.2020.101607.
- de la Fuente J, Estrada-Peña A, Rafael M, Almazán C, Bermúdez S, Abdelbaset AE, et al. Perception of ticks and tick-borne diseases worldwide. Pathogens 2023;12(10):1258. https://doi.org/10.3390/ PATHOGENS12101258.
- Ogden NH, Lindsay LR. Effects of climate and climate change on vectors and vector-borne diseases: ticks are different. Trends Parasitol 2016;32(8):646 – 56. https://doi.org/10.1016/j.pt.2016.04.015.
- 6. Ostfeld RS, Brunner JL. Climate change and Ixodes tick-borne diseases

- of humans. Philos Trans Roy Soc B: Biol Sci 2015;370(1665): 20140051. https://doi.org/10.1098/rstb.2014.0051.
- Zhao GP, Wang YX, Fan ZW, Ji Y, Liu MJ, Zhang WH, et al. Mapping ticks and tick-borne pathogens in China. Nat Commun 2021;12(1):1075. https://doi.org/10.1038/s41467-021-21375-1.
- Si BY, Jiang T, Zhang Y, Deng YQ, Huo QB, Zheng YC, et al. Complete genome sequence analysis of tick-borne encephalitis viruses isolated in northeastern China. Arch Virol 2011;156(8):1485 – 8. https://doi.org/10.1007/s00705-011-1031-y.
- Xu BL, Liu LC, Huang XY, Ma H, Zhang Y, Du YH, et al. Metagenomic analysis of fever, thrombocytopenia and leukopenia syndrome (FTLS) in Henan Province, China: discovery of a new bunyavirus. PLoS Pathog 2011;7(11):e1002369. https://doi.org/10. 1371/journal.ppat.1002369.
- Yu XJ, Liang MF, Zhang SY, Liu Y, Li JD, Sun YL, et al. Fever with thrombocytopenia associated with a novel bunyavirus in China. N Engl J Med 2011;364(16):1523 – 32. https://doi.org/10.1056/ NEJMoa1010095.
- 11. Zhang YZ, Zhou DJ, Xiong YW, Chen XP, He YW, Sun QZ, et al. Hemorrhagic fever caused by a novel tick-borne *Bunyavirus* in Huaiyangshan, China. Chin J Epidemiol 2011;32(3):209 20. https://doi.org/10.3760/cma.j.issn.0254-6450.2011.03.001.
- Kang XP, Li YC, Wei JJ, Zhang Y, Bian C, Wang K, et al. Elevation of matrix metalloproteinase-9 level in cerebrospinal fluid of tick-borne encephalitis patients is associated with IgG extravassation and disease severity. PLoS One 2013;8(11):e77427. https://doi.org/10.1371/ journal.pone.0077427.
- Lam TTY, Liu W, Bowden TA, Cui N, Zhuang L, Liu K, et al. Evolutionary and molecular analysis of the emergent severe fever with thrombocytopenia syndrome virus. Epidemics 2013;5(1):1 – 10. https://doi.org/10.1016/j.epidem.2012.09.002.
- Ding F, Guan XH, Kang K, Ding SJ, Huang LY, Xing XS, et al. Risk factors for bunyavirus-associated severe fever with thrombocytopenia syndrome, China. PLoS Negl Trop Dis 2014;8(10):e3267. https://doi. org/10.1371/journal.pntd.0003267.
- Ding SJ, Niu GY, Xu XH, Li JP, Zhang XM, Yin HY, et al. Age is a critical risk factor for severe fever with thrombocytopenia syndrome. PLoS One 2014;9(11):e111736. https://doi.org/10.1371/journal.pone. 0111736.
- 16. Liu K, Cui N, Fang LQ, Wang BJ, Lu QB, Peng W, et al. Epidemiologic features and environmental risk factors of severe fever with thrombocytopenia syndrome, Xinyang, China. PLoS Negl Trop Dis 2014;8(5):e2820. https://doi.org/10.1371/journal.pntd.0002820.
- 17. Xing XS, Guan XH, Liu L, Zhan JB, Jiang HB, Liu L, et al. Natural transmission model for severe fever with thrombocytopenia syndrome bunyavirus in villages of Hubei Province, China. Medicine 2016;95(4): e2533. https://doi.org/10.1097/MD.0000000000002533.
- Huang DY, Jiang YP, Liu XP, Wang B, Shi JM, Su Z, et al. A cluster of symptomatic and asymptomatic infections of severe fever with thrombocytopenia syndrome caused by person-to-person transmission. Am J Trop Med Hyg 2017;97(2):396 – 402. https://doi.org/10.4269/ ajtmh.17-0059.
- Li ZF, Hu JL, Cui LB, Hong Y, Liu JW, Li PF, et al. Increased prevalence of severe fever with thrombocytopenia syndrome in Eastern China clustered with multiple genotypes and reasserted virus during 2010-2015. Sci Rep 2017;7(1):6503. https://doi.org/10.1038/s41598-017-06853-1.
- Wu YY, Fu GM, Guo S, Ye L, Hou J, Wang JN, et al. The effects of health education and promotion with regard to severe fever with thrombocytopenia syndrome (SFTS) in rural residents: a pilot study in China. Biosci Trends 2018;11(6):697 – 701. https://doi.org/10.5582/ bst.2017.01252.
- Xing XS, Guan XH, Liu L, Xu JQ, Li GM, Zhan JB, et al. A casecontrol study of risk sources for severe fever with thrombocytopenia syndrome in Hubei Province, China. Int J Infect Dis 2017;55:86 – 91. https://doi.org/10.1016/j.ijid.2017.01.003.
- 22. Jia B, Wu WH, Huang R, Wang GY, Song PX, Li Y, et al. Characterization of clinical features and outcome for human-to-human

<sup>&</sup>lt;sup>1</sup> School of Global Health, Chinese Centre for Tropical Diseases Research, Shanghai Jiao Tong University School of Medicine, Shanghai, China; <sup>2</sup> School of Public Health, Shanghai Jiao Tong University School of Medicine, Shanghai, China.

- transmitted severe fever with thrombocytopenia syndrome. Infect Dis 2018;50(8):601 8. https://doi.org/10.1080/23744235.2018.1449962.
- Xu HH, Wei YY, Ma HQ, Liu YY, Zhang YL, Hu LF, et al. Alterations of gut microbiome in the patients with severe fever with thrombocytopenia syndrome. Front Microbiol 2018;9:2315. https:// doi.org/10.3389/fmicb.2018.02315.
- 24. Hu JL, Li ZF, Cai JP, Liu DL, Zhang XF, Jiang RJ, et al. A cluster of bunyavirus-associated severe fever with thrombocytopenia syndrome cases in a coastal plain area in China, 2015: identification of a previously unidentified endemic region for severe fever with thrombocytopenia bunyavirus. Open Forum Infect Dis 2019;6(6): ofz209. https://doi.org/10.1093/ofid/ofz209.
- Jia N, Liu HB, Ni XB, Bell-Sakyi L, Zheng YC, Song JL, et al. Emergence of human infection with Jingmen tick virus in China: a retrospective study. eBioMedicine 2019;43:317 – 24. https://doi.org/ 10.1016/j.ebiom.2019.04.004.
- Qi R, Qin XR, Wang L, Han HJ, Cui F, Yu H, et al. Severe fever with thrombocytopenia syndrome can masquerade as hemorrhagic fever with renal syndrome. PLoS Negl Trop Dis 2019;13(3):e0007308. https:// doi.org/10.1371/journal.pntd.0007308.
- Tian B, Qu D, Sasaki A, Chen JY, Deng BC. Acute pancreatitis in patients with severe fever with thrombocytopenia syndrome virus infection. Pancreatology 2020;20(8):1631 – 6. https://doi.org/10.1016/ j.pan.2020.09.024.
- Zhang YY, Liu DX, Fan WB, Yu JH, Liu L, Du WJ, et al. Hazardous alcohol consumption and aging synergistically increase the risk of death in patients with severe fever with thrombocytopenia syndrome. Ticks Tick Borne Dis 2020;11(6):101505. https://doi.org/10.1016/j.ttbdis. 2020.101505.
- 29. Du YD, Mi ZH, Xie YP, Lu DS, Zheng HJ, Sun H, et al. Insights into the molecular basis of tick-borne encephalitis from multiplatform metabolomics. PLoS Negl Trop Dis 2021;15(3):e0009172. https://doi.org/10.1371/journal.pntd.0009172.
- Hu LF, Kong QX, Liu YY, Li JJ, Bian TT, Ma XJ, et al. Time course of severe fever with thrombocytopenia syndrome virus and antibodies in patients by long-term follow-up study, China. Front Microbiol 2021;12:744037. https://doi.org/10.3389/fmicb.2021.744037.
- Wang YC, Wei ZK, Lv XL, Han SZ, Wang ZD, Fan CF, et al. A new nairo-like virus associated with human febrile illness in China. Emerg Microbes Infect 2021;10(1):1200 – 8. https://doi.org/10.1080/ 22221751.2021.1936197.
- 32. You EQ, Wang L, Zhang L, Wu JJ, Zhao KF, Huang F. Epidemiological characteristics of severe fever with thrombocytopenia syndrome in Hefei of Anhui Province: a population-based surveillance study from 2011 to 2018. Eur J Clin Microbiol Infect Dis 2021;40(5): 929 39. https://doi.org/10.1007/s10096-020-04098-x.
- Zhao J, Lu QB, Li H, Yuan Y, Cui N, Yuan C, et al. Sex differences in case fatality rate of patients with severe fever with thrombocytopenia syndrome. Front Microbiol 2021;12:738808. https://doi.org/10.3389/ fmicb.2021.738808.
- 34. Dai ZN, Peng XF, Li JC, Zhao J, Wu YX, Yang X, et al. Effect of genomic variations in severe fever with thrombocytopenia syndrome virus on the disease lethality. Emerg Microbes Infect 2022;11(1):1672 82. https://doi.org/10.1080/22221751.2022.2081617.
- 35. Han XH, Ma Y, Liu HY, Li D, Wang Y, Jiang FH, et al. Identification of severe fever with thrombocytopenia syndrome virus genotypes in patients and ticks in Liaoning Province, China. Parasit Vectors 2022;15 (1):120. https://doi.org/10.1186/s13071-022-05237-3.
- Zhao J, Ge HH, Wang G, Lin L, Yuan Y, Xu YL, et al. Fatal patients with severe fever with thrombocytopenia syndrome in China. Int J Infect Dis 2022;125:10 – 6. https://doi.org/10.1016/j.ijid.2022.10.008.
- Liu T, Zhang NN, Li HW, Hou ST, Liu XW. Analysis of severe fever with thrombocytopenia syndrome cluster in East China. Virol J 2023;20(1):199. https://doi.org/10.1186/s12985-023-02155-3.
- Ren YT, Tian HP, Xu JL, Liu MQ, Cai K, Chen SL, et al. Extensive genetic diversity of severe fever with thrombocytopenia syndrome virus circulating in Hubei Province, China, 2018-2022. PLoS Negl Trop Dis 2023;17(9):e0011654. https://doi.org/10.1371/journal.pntd.0011654.

- Zheng HX, Geng Y, Gu CG, Li M, Mao MX, Wan YW, et al. A reservoir computing with boosted topology model to predict encephalitis and mortality for patients with severe fever with thrombocytopenia syndrome: a retrospective multicenter study. Infect Dis Ther 2023;12(5):1379 91. https://doi.org/10.1007/s40121-023-00808-v.
- Fei X, Feng BB, Fang K, Ren WH. Risk factors for mortality in severe fever with thrombocytopenia syndrome patients with central nervous system complications. Med Sci Monit 2023;29:e938427. https://doi. org/10.12659/MSM.938427.
- Lv XL, Liu ZY, Li L, Xu WB, Yuan YX, Liang XJ, et al. Yezo virus infection in tick-bitten patient and ticks, northeastern China. Emerg Infect Dis 2023;29(4):797 – 800. https://doi.org/10.3201/eid2904. 220885.
- 42. Li HY, Zhou Y, Wang WJ, Guo DM, Huang S, Jie SH. The clinical characteristics and outcomes of patients with human granulocytic anaplasmosis in China. Int J Infect Dis 2011;15(12):e859 66. https://doi.org/10.1016/j.ijid.2011.09.008.
- Zhang LJ, Cui F, Wang LL, Zhang LL, Zhang JS, Wang SW, et al. Investigation of anaplasmosis in Yiyuan County, Shandong Province, China. Asian Pac J Trop Med 2011;4(7):568 – 72. https://doi.org/10. 1016/S1995-7645(11)60148-X.
- 44. Wang Y, Chen CF, Zhang LJ. Molecular characterization of msp2/p44 of Anaplasma phagocytophilum isolated from infected patients and Haemaphysalis longicornis in Laizhou Bay, Shandong Province, China. PLoS One 2013;8(10):e78189. https://doi.org/10.1371/journal.pone. 0078189.
- Zhang LJ, Wang GQ, Liu QH, Chen CF, Li J, Long B, et al. Molecular analysis of *Anaplasma phagocytophilum* isolated from patients with febrile diseases of unknown etiology in China. PLoS One 2013;8(2): e57155. https://doi.org/10.1371/journal.pone.0057155.
- 46. Ni XB, Jia N, Jiang BG, Sun T, Zheng YC, Huo QB, et al. Lyme borreliosis caused by diverse genospecies of *Borrelia burgdorferi* sensu lato in northeastern China. Clin Microbiol Infect 2014;20(8):808 – 14. https://doi.org/10.1111/1469-0691.12532.
- 47. Jiao XY, Fan ZC, Li YZ, Tang YT, Ke CW. Clinical and laboratory features parameters of human granulocytic anaplasmosis (HGA) in patients admitted to hospital in Guangdong Province, China. Trop Doct 2015;45(4):209 13. https://doi.org/10.1177/00494755155 79772
- 48. Li H, Zheng YC, Ma L, Jia N, Jiang BG, Jiang RR, et al. Human infection with a novel tick-borne *Anaplasma* species in China: a surveillance study. Lancet Infect Dis 2015;15(6):663 70. https://doi.org/10.1016/S1473-3099(15)70051-4.
- Li H, Cui XM, Cui N, Yang ZD, Hu JG, Fan YD, et al. Human infection with novel spotted fever group rickettsia genotype, China, 2015. Emerg Infect Dis 2016;22(12):2153 – 6. https://doi.org/10. 3201/eid2212.160962.
- Liu W, Li H, Lu QB, Cui N, Yang ZD, Hu JG, et al. *Candidatus* rickettsia tarasevichiae infection in eastern central China: a case series. Ann Intern Med 2016;164(10):641 – 8. https://doi.org/10.7326/M15-2572.
- Jiang BG, Jia N, Jiang JF, Zheng YC, Chu YL, Jiang RR, et al. *Borrelia miyamotoi* infections in humans and ticks, northeastern China. Emerg Infect Dis 2018;24(2):236 41. https://doi.org/10.3201/eid2402. 160378.
- 52. Li H, Zhang PH, Huang Y, Du J, Cui N, Yang ZD, et al. Isolation and identification of *Rickettsia raoultii* in human cases: a surveillance study in 3 medical centers in China. Clin Infect Dis 2018;66(7):1109 15. https://doi.org/10.1093/cid/cix917.
- Lu QY, Yu JP, Yu LQ, Zhang YJ, Chen YT, Lin MA, et al. Rickettsia japonica infections in humans, Zhejiang Province, China, 2015. Emerg Infect Dis 2018;24(11):2077 – 9. https://doi.org/10.3201/eid2411. 170044
- 54. Gaowa, Wulantuya, Yin XH, Guo SC, Ding CL, Cao MZ, et al. Spotted fever group rickettsiae in Inner Mongolia, China, 2015-2016. Emerg Infect Dis 2018;24(11):2105 7. https://doi.org/10.3201/eid2411.162094.

- Li H, Zhang PH, Du J, Yang ZD, Cui N, Xing B, et al. *Rickettsia japonica* infections in humans, Xinyang, China, 2014-2017. Emerg Infect Dis 2019;25(9):1719 22. https://doi.org/10.3201/eid2509. 171421.
- 56. Wang F, Ma MB, Luo SY, Yan M, Tao LY, Liu AH, et al. Seroprevalence of tick-borne *Anaplasma phagocytophilum* infection in healthy adult population and patients with acute undifferentiated fever from the Yunnan Province of China. Vector Borne Zoonotic Dis 2019;19(8):576 81. https://doi.org/10.1089/vbz.2018.2389.
- Jia N, Liu HB, Zheng YC, Shi WQ, Wei R, Chu YL, et al. Cutaneous immunoprofiles of three spotted fever group rickettsia cases. Infect Immun 2020;88(4):e00686 – 19. https://doi.org/10.1128/IAI.00686-19
- Li H, Li XM, Du J, Zhang XA, Cui N, Yang ZD, et al. *Candidatus* rickettsia xinyangensis as cause of spotted fever group rickettsiosis, Xinyang, China, 2015. Emerg Infect Dis 2020;26(5):985 8. https://doi.org/10.3201/eid2605.170294.
- Gao Y, Lv XL, Han SZ, Wang W, Liu Q, Song MX. First detection of Borrelia miyamotoi infections in ticks and humans from the northeast of Inner Mongolia, China. Acta Trop 2021;217:105857. https://doi.org/ 10.1016/j.actatropica.2021.105857.
- Lyu Y, Shen Y, Hu CY, Xu PP, Qin W, Gong L, et al. The first reported outbreak of an undetermined species of human infection with spotted fever group Rickettsia in Lu'an, China. Acta Trop 2021;223: 106072. https://doi.org/10.1016/j.actatropica.2021.106072.
- 61. Jiang JF, Zheng YC, Jiang RR, Li H, Huo QB, Jiang BG, et al. Epidemiological, clinical, and laboratory characteristics of 48 cases of "*Babesia venatorum*" infection in China: a descriptive study. Lancet Infect Dis 2015;15(2):196 203. https://doi.org/10.1016/S1473-3099 (14)71046-1.
- 62. Jia N, Zheng YC, Jiang JF, Jiang RR, Jiang BG, Wei R, et al. Human babesiosis caused by a *Babesia* Crassa-like pathogen: a case series. Clin Infect Dis 2018;67(7):1110 9. https://doi.org/10.1093/cid/ciy212.
- 63. Wang JM, Zhang SD, Yang JQ, Liu JL, Zhang DK, Li YQ, et al. Babesia divergens in human in Gansu province, China. Emerg Microbes Infect 2019;8(1):959 – 61. https://doi.org/10.1080/22221751.2019. 1635431.
- Zhao L, Jiang RR, Jia N, Ning NZ, Zheng YC, Huo QB, et al. Human case infected with *Babesia venatorum*: a 5-year follow-up study. Open Forum Infect Dis 2020;7(3):ofaa062. https://doi.org/10.1093/ofid/ ofaa062.
- Li J, Li S, Yang L, Cao PF, Lu JH. Severe fever with thrombocytopenia syndrome virus: a highly lethal bunyavirus. Crit Rev Microbiol 2021;47 (1):112 – 25. https://doi.org/10.1080/1040841X.2020.1847037.
- 66. Liu Q, He B, Huang SY, Wei F, Zhu XQ. Severe fever with thrombocytopenia syndrome, an emerging tick-borne zoonosis. Lancet Infect Dis 2014;14(8):763 – 72. https://doi.org/10.1016/S1473-3099 (14)70718-2.
- Robinson MT, Satjanadumrong J, Hughes T, Stenos J, Blacksell SD. Diagnosis of spotted fever group *Rickettsia* infections: the Asian perspective. Epidemiol Infect 2019;147:e286. https://doi.org/10.1017/ S0950268819001390.
- Parola P, Paddock CD, Socolovschi C, Labruna MB, Mediannikov O, Kernif T, et al. Update on tick-borne rickettsioses around the world: a geographic approach. Clin Microbiol Rev 2013;26(4):657 – 702. https://doi.org/10.1128/CMR.00032-13.
- MacQueen D, Centellas F. Human granulocytic anaplasmosis. Infect Dis Clin North Am 2022;36(3):639 – 54. https://doi.org/10.1016/j. idc.2022.02.008.
- Dumic I, Jevtic D, Veselinovic M, Nordstrom CW, Jovanovic M, Mogulla V, et al. Human granulocytic anaplasmosis-a systematic review of published cases. Microorganisms 2022;10(7):1433. https://doi.org/ 10.3390/MICROORGANISMS10071433.
- 71. Jin HT, Wei F, Liu Q, Qian J. Epidemiology and control of human granulocytic anaplasmosis: a systematic review. Vector Borne Zoonotic Dis 2012;12(4):269 74. https://doi.org/10.1089/vbz.2011.0753.
- 72. Zhang SY, Hai R, Li WY, Li GH, Lin GY, He JR, et al. Seroprevalence

- of human granulocytotropic anaplasmosis in central and southeastern China. Am J Trop Med Hyg 2009;81(2):293 5. https://doi.org/10.4269/ajtmh.2009.81.293.
- Hamburg BJ, Storch GA, Micek ST, Kollef MH. The importance of early treatment with doxycycline in human ehrlichiosis. Medicine 2008;87(2):53 – 60. https://doi.org/10.1097/MD.0b013e318168da1d.
- Wu XB, Na RH, Wei SS, Zhu JS, Peng HJ. Distribution of tick-borne diseases in China. Parasit Vectors 2013;6:119. https://doi.org/10.1186/ 1756-3305-6-119.
- 75. Mead PS. Epidemiology of Lyme disease. Infect Dis Clin North Am 2015;29(2):187 210. https://doi.org/10.1016/j.idc.2015.02.010.
- Stanek G, Strle F. Lyme borreliosis-from tick bite to diagnosis and treatment. FEMS Microbiol Rev 2018;42(3):233 – 58. https://doi.org/ 10.1093/femsre/fux047.
- 77. Vannier E, Krause PJ. Human babesiosis. N Engl J Med 2012;366(25): 2397 407. https://doi.org/10.1056/NEJMra1202018.
- Gray JS, Estrada-Peña A, Zintl A. Vectors of babesiosis. Annu Rev Entomol 2019;64:149 – 65. https://doi.org/10.1146/annurev-ento-011118-111932.
- Chen ZT, Li HQ, Gao XG, Bian AN, Yan HR, Kong DL, et al. Human Babesiosis in China: a systematic review. Parasitol Res 2019;118(4):1103 – 12. https://doi.org/10.1007/s00436-019-06250-9.
- Voisin O, Monpierre L, Le Lorc'h E, Pilmis B, Le Monnier A, Mourad JJ, et al. A typical babesiosis in an immunocompetent patient. Ann Biol Clin (Paris) 2021;79(5):456 9. https://doi.org/10.1684/abc.2021. 1675.
- 81. Stein E, Elbadawi LI, Kazmierczak J, Davis JP. Babesiosis surveillance Wisconsin, 2001-2015. MMWR Morb Mortal Wkly Rep 2017;66(26): 687 91. https://doi.org/10.15585/mmwr.mm6626a2.
- 82. Phipps LP, Johnson N. Tick-borne encephalitis virus. J Med Microbiol 2022;71(5):001492. https://doi.org/10.1099/jmm.0.001492.
- Johnson N, Migné CV, Gonzalez G. Tick-borne encephalitis. Curr Opin Infect Dis 2023;36(3):198 – 202. https://doi.org/10.1097/QCO. 0000000000000924.
- 84. Chen XJ, Li F, Yin QK, Liu WJ, Fu SH, He Y, et al. Epidemiology of tick-borne encephalitis in China, 2007-2018. PLoS One 2019;14(12): e0226712. https://doi.org/10.1371/journal.pone.0226712.
- Chiffi G, Grandgirard D, Leib SL, Chrdle A, Růžek D. Tick-borne encephalitis: a comprehensive review of the epidemiology, virology, and clinical picture. Rev Med Virol 2023;33(5):e2470. https://doi.org/10. 1002/rmv.2470.
- Luan YX, Gou JM, Zhong DJ, Ma L, Yin CS, Shu MF, et al. The tick-borne pathogens: an overview of China's situation. Acta Parasitol 2023;68(1):1 20. https://doi.org/10.1007/s11686-023-00658-1.
- 87. Madison-Antenucci S, Kramer LD, Gebhardt LL, Kauffman E. Emerging tick-borne diseases. Clin Microbiol Rev 2020;33(2):e00083 18. https://doi.org/10.1128/CMR.00083-18.
- Seo JW, Kim D, Yun N, Kim DM. Clinical update of severe fever with thrombocytopenia syndrome. Viruses 2021;13(7):1213. https://doi.org/ 10.3390/v13071213.
- 89. Kim D, Lai CJ, Cha I, Jung JU. Current progress of severe fever with thrombocytopenia syndrome virus (SFTSV) vaccine development. Viruses 2024;16(1):128. https://doi.org/10.3390/V16010128.
- Cun DJ, Wang Q, Yao XY, Ma B, Zhang Y, Li LH. Potential suitable habitats of *Haemaphysalis longicornis* in China under different climatic patterns. Chin J Schistosomiasis Control 2021;33(4):359 – 64. https:// doi.org/10.16250/j.32.1374.2021023.
- Yuan CF, Lu YJ, Li JQ, Chen C, Wang YH, Zheng AH, et al. Infection and transovarial transmission of severe fever with thrombocytopenia syndrome virus in *Rhipicephalus sanguineus* in Hainan Island, China. Integr Zool 2023;18(6):1009 – 13. https://doi.org/10.1111/1749-4877.12716.
- Fei SW, Zhao HQ, Yin JX, Wang L, Sun ZS, Zhang WG, et al. Molecular identification and genetic characterization of public health threatening ticks - Chongming Island, China, 2021-2022. China CDC Wkly 2023;5(37):815 - 21. https://doi.org/10.46234/ccdcw2023.156.