

ОБЗОРЫ ЛИТЕРАТУРЫ

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OVERVIEW OF TICK-ASSOCIATED RICKETTSIALES BACTERIA IN JAPAN

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Rickettsiales bacteria are obligatory intracellular parasites in vertebrates and/or invertebrates, and some cause severe febrile illness in humans as well as livestock. In Japan, "*Rickettsia japonica*" and "*Orientia tsutsugamushi*" are well known as causative agents for "Japanese spotted fever" transmitted by ticks and "*Tsutsugamushi* disease" transmitted by mites, respectively. In recent years, many researchers have reported the presence of several tick-associated *Rickettsiales* bacteria in Japan, mostly by molecular-based analyses. Some of these might be public health or veterinary significance. Such *Rickettsiales* bacteria include spotted fever group rickettsiae other than *R. japonica*, *Ehrlichia* spp., *Anaplasma* spp., and *Neoehrlichia* sp. We will summarize and introduce the representatives of these *Rickettsiales* bacteria present in Japan.

Key word: *Rickettsiales*, *Rickettsia*, *Ehrlichia*, *Anaplasma*, *Neoehrlichia*

ОБЗОР БАКТЕРИЙ ПОРЯДКА RICKETTSIALES, АССОЦИИРОВАННЫХ С ИКСОДОВЫМИ КЛЕЩАМИ В ЯПОНИИ

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Бактерии порядка *Rickettsiales* являются облигатными внутриклеточными паразитами позвоночных и беспозвоночных животных и некоторые из них вызывают тяжелые лихорадочные заболевания у человека и домашних животных. В Японии хорошо известны микроорганизмы *Rickettsia japonica* и *Orientia tsutsugamushi*, которые являются возбудителями Японской пятнистой лихорадки, переносимой иксодовыми клещами и лихорадки Цуцугамуши, переносимой соответственно краснотелковыми клещами. В последние годы появляется много сообщений о выявлении, преимущественно молекулярно-генетическими методами, в Японии нескольких новых бактерий порядка *Rickettsiales*, ассоциированных с клещами. Некоторые из этих бактерий могут иметь существенное ветеринарное или медицинское значение. Подобные бактерии включают риккетсий группы клещевых пятнистых лихорадок не относящихся к *R. japonica*, *Ehrlichia* spp., *Anaplasma* spp., и *Neoehrlichia* sp. В данном сообщении характеризуются основные представители этих групп, обитающие на территории Японии.

Ключевые слова: *Rickettsiales*, *Rickettsia*, *Ehrlichia*, *Anaplasma*, *Neoehrlichia*

The order *Rickettsiales* includes the genera *Rickettsia*, *Orientia*, *Ehrlichia*, *Anaplasma*, and *Neoehrlichia* etc. [1, 2]. In Japan, before the first report of "Japanese spotted fever (JSF)" caused by *Rickettsia japonica* in 1984 [3], "*Tsutsugamushi* disease" caused by *Orientia tsutsugamushi* had only been known as a rickettsiosis. *R. japonica* and *O. tsutsugamushi* are transmitted by ticks and mites, respectively. In recent years, many isolation and molecular analyses revealed the presence of tick-associated *Rickettsiales* bacteria such as spotted fever group rickettsiae other than *R. japonica*, *Ehrlichia* spp., *Anaplasma* spp., and *Neoehrlichia* sp. in ticks or wild mammals. The representatives of these *Rickettsiales* bacteria in Japan are summarized as below.

SPOTTED FEVER GROUP (SFG) RICKETTSIAE

So far, SFG rickettsiae which were identified in Japan are *R. japonica* [4], *R. helvetica* [5], *R. tamurae* [6], *R. asiatica* [7], *Candidatus R. tarasevichiae* [8],

and *R. heilonjiangensis* [9]. *Rickettsia japonica* was isolated from tick species of *Haemaphysalis histricis*, *H. cornigera*, *H. flava*, *H. longicornis*, *Dermacentor taiwanensis*, and *Ixodes ovatus* [10]. *R. helvetica* was also isolated from ticks of *Ixodes ovatus*, *I. persulcatus*, and *I. monospinosus* [5] and also detected from Sika deer (*Cervus nippon yesoensis*) in Hokkaido, a northern part of Japan [11]. *R. tamurae* was isolated from *Amblyomma testudinarium* ticks [6]. *R. asiatica* was isolated from *I. ovatus* ticks [7]. *Candidatus R. tarasevichiae* was detected by PCR from *I. persulcatus* ticks in Hokkaido [8]. *R. heilonjiangensis* was isolated from *H. concinna* ticks [9]. Of these, *R. japonica* cause severe febrile illness in humans frequently in Japan [12]. Human infection by *R. heilonjiangensis* has recently been confirmed [9]. Additionally, *R. tamurae* was found to cause rickettsiosis, but it seems to be rare. The immunocompromised hosts are probably more sensitive to *R. tamurae* infection [13].

EHRlichia spp.

Ehrlichia muris have firstly been isolated from a wild mouse of *Eothenomys kageus* (synonym of *E. smithi*) and reported in 1995 in Japan [14]. Later, several nonclassified genetic variants of the genus *Ehrlichia* have been found in Japan. Of these, the *Ehrlichia* HF strain has well studied. The HF strain was initially isolated and described from *I. ovatus* tick in 2000 [15]. The genetic variants of HF strain were detected by molecular analysis from wild mice of *Apodemus argenteus*, *A. speciosus*, and *E. smithi* [16] and dogs [17]. These bacteria cause fetal infection to immunocompetent laboratory mice. Recently, the *Ehrlichia* variant closely related to *E. ewingii* was identified from *H. longicornis* ticks [18]. *E. chaffeensis* which is well known as a human pathogen was detected from Sika deer (*Cervus nippon nippon*) and described in 2009 [19], although there is no report of human infection in Japan to date.

ANAPLASMA spp.

We have initially detected *Anaplasma phagocytophilum*, a human pathogen, from *I. persulcatus* and *I. ovatus* ticks inhabiting Japan in 2005 [20]. Recently, we confirmed that *H. formosensis*, *H. longicornis*, and *H. megaspinosus* were also potential arthropod vectors for transmission of *A. phagocytophilum* [21]. The genetic variants of *A. phagocytophilum* have also been identified from Sika deer and boars in Japan [22, 23]. Besides *A. phagocytophilum*, *A. centrale* and *A. bovis* were detected from deer and *Haemaphysalis* ticks [22]. *A. platys* was identified from dog in Japan [24].

NEOEhrlichia sp.

Candidatus Neoehrlichia mikurensis and its genetic variants have firstly discovered from wild rodents (*Rattus norvegicus*) and *I. ovatus* ticks in Japan in 2004 [2]. Later, we and the others were also detected from wild mice of *A. argenteus*, *A. speciosus*, and *E. smithi* [25, 26]. Recently, the human cases of *Candidatus N. mikurensis* infection have been reported in Europe [27-29]. In the most cases, the disease developed in immunocompromised individuals. The vector candidates are *Ixodes* ticks such as *I. ricinus*, *I. persulcatus*, and *I. ovatus* in Europe and Asia [2, 30, 31].

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