

Flea-Borne Rickettsiae in Almaty Oblast, Kazakhstan

Talgat Nurmakhanov¹, Yerlan Sansyzbayev¹, Heidi St. John², Christina Farris² and Allen Richards²

¹Kazakh Scientific Center for Quarantine and Zoonotic Disease, Almaty, Kazakhstan; ²Naval Medical Research Center, Silver Spring, MD, USA

Introduction

Flea-borne diseases in Kazakhstan have been a significant health risk to inhabitants and visitors for ages, particularly plague. Flea-borne rickettsial disease threats are unknown in Kazakhstan, we therefore initiated a study to detect and identify flea-borne rickettsiae among fleas collected in the Almaty Oblast, in southeastern Kazakhstan.

Methods

Fleas (n=248) were collected by members of the Taldykorgan anti-plague station from live captured rodents (i.e. the Great Gerbil-*Rhombomys opimus*) and from the rodent burrows collected at five Rayons (districts) within Almaty Oblast (province) during 2015. Fleas were identified morphologically by entomologic keys and then pooled together (1-50 fleas/pool) by species and host/rodent burrow. DNA was extracted from triturated fleas (PrepMan Ultra kit) and tested by genus- (*Rickettsia*), group- (*R. felis* genogroup), and species- (*Rickettsia typhi*, *Rickettsia felis* and *Candidatus Rickettsia asemboensis*) specific quantitative real-time PCR (qPCR) assays, Rick17b, RfelG, Rtyph, and Rasemb, respectively. With GPS coordinates and GIS (ArcGIS) a distribution map was developed.

Results

Of 248 fleas (*Coptopsylla lamellifer* 45, *Echidnophaga oschanini* 1, *Nosopsyllus laeviceps* 10, *Nosopsyllus tarsus* 1, *Nosopsyllus turkmenikus* 1, *Paradoxopsyllus teretifrons* 2, *Xenopsylla conformis* 1, *Xenopsylla gerbilli* 87, *Xenopsylla hirtipes* 26, and *Xenopsylla skrjabini* 74) assessed by qPCR 56 were identified as having: *Rickettsia* spp. only n=20, *R. felis* genogroup n=8, *R. felis* n=1, and *Ca. R. asemboensis* n=27. *X. gerbilli* was the flea most frequently found to be infected with a rickettsiae (44 of 87;50.6%) and 25 of the 44 rickettsia-infected fleas (56.8%) were infected by *Ca. R. asemboensis*. *X. hirtipes* was the next most commonly infected flea (4 of 26; 15.4%). One flea was infected with *R. felis*, and none were infected with *R. typhi*. *R. felis* and *R. typhi* cause flea-borne spotted fever and murine typhus, respectively.

Conclusions

Fleas captured from *R. opimus* or at their burrows were infected with rickettsiae. Most commonly found rickettsia-infected flea species was *X. gerbilli* and the most commonly found rickettsia was *Ca. R. asemboensis*. Future studies may include testing these and other fleas samples for the presence of other disease agents including *Bartonella* spp. and *Yersinia pestis*.

Keywords

flea-borne disease; rickettsiae; Kazakhstan

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