



Epidemiology of Tularemia and *Francisella tularensis*

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Message from the Guest Editor

Dear Colleague,

Tularemia is an emerging disease affecting humans and animals in the northern hemisphere. The disease is caused by the bacterium *Francisella tularensis*, which has been isolated from many vertebrates, ticks, and biting insects.

Environmental changes and changing recreational activities can influence the reservoir, as well as transmission and spread of the disease. Whole-genome sequencing has helped to elucidate epidemiological patterns, but sequencing and the bioinformatics tools are not yet established in all affected countries.

This Special Issue wants to provide a comprehensive picture of the current knowledge on the epidemiological patterns and the tools to analyze them.

Manuscripts dealing with systemic review will only be acceptable in rare cases.

PD Dr. Herbert Tomaso
Guest Editor





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Message from the Editor-in-Chief

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