

Appendix Table 1. *Francisella tularensis* genomic sequences used for indel identification

<i>F. tularensis</i> subspecies	Strain	FSC* no.	ATCC† no.	Source of genomic sequence	Status	GenBank accession no.	Reference
<i>novicida</i>	U112	040‡	15482	University of Washington, Seattle, WA, USA	Finished	CP000439	Unpub.
<i>tularensis</i>	SCHU S4	237	–	Swedish/British/US consortium§	Finished	AJ749949	(1)
<i>mediasiatica</i>	GIEM 543	147	–	Los Alamos National Laboratory, Los Alamos, NM, USA	Unfinished	Pending	Unpub.
<i>holarctica</i>	OSU18	–	–	Baylor College of Medicine, Waco, TX, USA	Finished	CP000437	(2)
<i>holarctica</i>	LVS	458‡	29684	Lawrence Livermore National Laboratory, CA, USA	Finished	AM233362	Unpub.

**Francisella* Strain Collection.

†American Type Culture Collection.

‡Strains sequenced were not obtained from the *Francisella* Strain Collection.

§Member institutes of the sequencing consortium were: Swedish Defence Research Agency, Sweden; Uppsala and Umeå Universities, Sweden, Defence Science and Technology Laboratories, Salisbury, UK; Walter Reed Army Institute of Research, Bethesda, Maryland, USA.

References

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- Petrosino JF, Xiang Q, Karpathy SE, Jiang H, Yerrapragada S, Liu Y, et al. [Chromosome rearrangement and diversification of *Francisella tularensis* revealed by the type B \(OSU18\) genome sequence.](#) J Bacteriol. 2006;188:6977–85.