

**Genomic DNA from *Francisella tularensis* subsp. *novicida*, Strain ΔIglD**

**Catalog No. NR-13362**

**Product Description:** Genomic DNA was isolated from a preparation of *Francisella tularensis* (*F. tularensis*) subsp. *novicida*, strain ΔIglD. *F. tularensis* subsp. *novicida*, strain ΔIglD is a transposon mutant of the wild-type strain U112, in which the *igID* gene region has been replaced with a mini-Tn5 insert, rendering it resistant to kanamycin.

**Lot<sup>1</sup>: 59882648**

**Manufacturing Date: 11APR2011**

TEST	SPECIFICATIONS	RESULTS
<b>Sequencing of 16S Ribosomal RNA Gene</b> (~ 1370 base pairs)	Identical to BEI Resources NR-9713 Consistent with <i>F. tularensis</i>	Identical to BEI Resources NR-9713 Consistent with <i>F. tularensis</i>
<b>Agarose Gel Electrophoresis</b>	High molecular weight chromosomal DNA	High molecular weight chromosomal DNA (Figure 1)
<b>Concentration by PicoGreen<sup>®</sup> Measurement</b>	0.7 to 1.5 µg in 25 to 100 µL per vial	0.9 µg in 32 µL per vial (28 µg/mL)
<b>Functional Activity by PCR Amplification</b> 16S ribosomal RNA gene	~ 1500 bp amplicon	~ 1500 bp amplicon
<b>Molecular Subtyping by PCR Amplification of Subspecies-Specific Sequence from Extracted DNA<sup>2,3</sup></b>	~ 1500 bp amplicon (subsp. <i>tularensis</i> ) ~ 900 bp amplicon (subsp. <i>holarctica</i> ) ~ 3300 bp amplicon (subsp. <i>novicida</i> )	~ 3300 bp amplicon (subsp. <i>novicida</i> )
<b>OD<sub>260</sub>/OD<sub>280</sub> Ratio</b>	1.7 to 2.0	2.0
<b>Bacterial Inactivation</b> 10% of total yield plated on Chocolate Agar <sup>4,5</sup>	No viable bacteria detected	No viable bacteria detected

<sup>1</sup>NR-13362 was produced by inoculation of NR-9713 (Lot 58893070) into Brain Heart Infusion Broth and grown 24 hours at 37°C. Broth inoculum was added to Chocolate agar Kolles which were grown 24 hours at 37°C and genomic DNA was extracted using proprietary technology.

<sup>2</sup>Broekhuijsen, M., et al. "Genome-Wide DNA Microarray Analysis of *Francisella tularensis* Strains Demonstrates Extensive Genetic Conservation within the Species but Identifies Regions that are Unique to the Highly Virulent *F. tularensis* subsp. *tularensis*." *J. Clin. Microbiol.* 41 (2003): 2924-2931. PubMed: 12843022.

<sup>3</sup>The Molecular Subtyping by PCR Amplification of Subspecies-Specific Sequence from Extracted DNA was performed on NR-13362 lot 58893072 which was extracted from the same material that was used to produce NR-9713 lot 58893070. NR-13362 lot 59882648 was extracted from NR-9713 lot 58893070.

<sup>4</sup>Incubated for 7 days at 37°C and aerobic atmosphere

<sup>5</sup>An extraction procedure was used that has been shown to consistently inactivate 100% of Gram-positive and Gram-negative bacteria.

**Date:** 11 NOV 2011

**Signature:**



**Title:** Technical Manager, BEI Authentication or designee

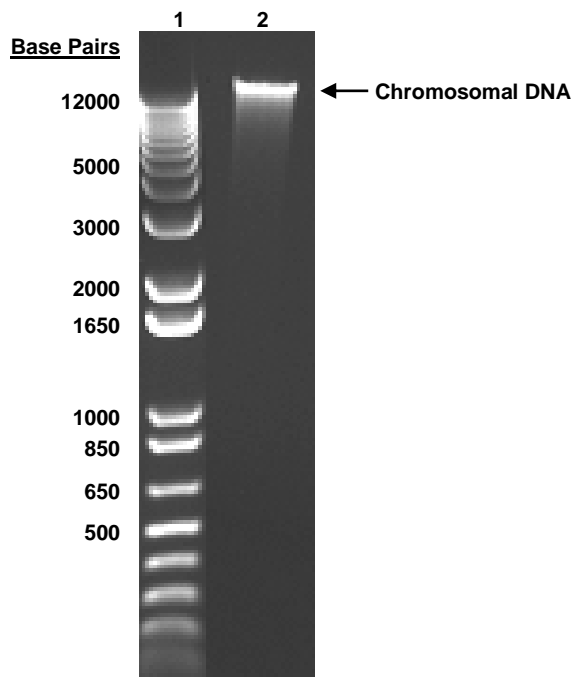
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Figure 1



Lane 1: Invitrogen™ TrackIt 1 Kb Plus DNA Ladder™  
Lane 2: 200 ng of HM-13362