 Genome Sequence of *Bacillus anthracis* STI, a Sterne-Like Georgian/Soviet Vaccine Strain

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The *Bacillus anthracis* strain STI is a Soviet vaccine strain that lacks the pXO2 plasmid. Previous data indicate that this isolate forms a new branch within the *B. anthracis* sub-group originally identified as A. Br.008/009.

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This study was financially supported by the U.S. Department of Homeland Security (HSHQDC-10-C-00139). Portions of this work were performed using the Ames ancestor reference genome (6) as a template. The conserved nature of the *B. anthracis* genome is clearly visible in the identification of unique regions in new assemblies. A *de novo* assembly was also performed using the paired-end sequence data with CLC Bio Genomics WorkBench and the resulting analysis yielded 126 contigs. Both the template and *de novo* assembled contigs were submitted to the RAST annotation server for subsystem classification and functional annotation (http://metagenomics.anl.gov/) (8).

Nucleotide sequence accession numbers. The NCBI BioProject ID is PRJNA243052 and the GenBank accession numbers for this STI genome are BaA0485_chr (CP007660) and BaA0485_pXO1 (CP007661).

REFERENCES
