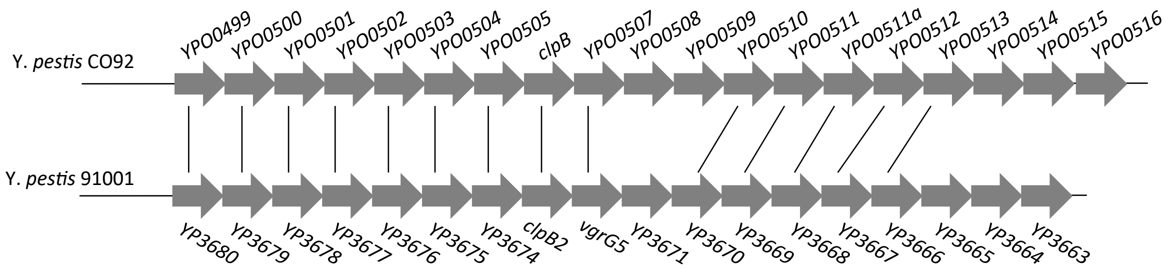
Supplementary Table S1. Oligonucleotide primers designed in this study

Target	Primers (forward/reverse, 5′–3′)
qPCR	
YP3680	*ATGACGCCTACCGGCTCAG/GATGCAGAGTGTTGATCAGGT*
16S rDNA	*CACACTGGAACTGAGACAC/TGCTTCTTCTGCGAGTAAC*
LacZ fusion	
YP3680	*GCAGGATCCGCATTCAAATCCAGCTCTGATT/GGTAAGCTTGAGAGGATGGACTCAATCT*
Primer extension	
YP3680	*/TCCTGATTTACATACCTGTTATA*
EMSA	
YP3680	*GCATTCAAATCCAGCTCTGATT/GAGAGGATGGACTCAATCT*



Supplementary Figure S1. Organization of the T6SS gene cluster in *Y. pestis*. Genes connected by the lines indicate alleles in the two *Yersinia* strains.

Supplementary Figure S2. Promoter structure of *YP3680*. Shown were translation/transcription starts, predicted core promoter −10 and −35 elements, predicted Shine-Dalgarno (SD) sequences for ribosomal binding, and RovA and RcsAB box-like sequences.