Table S1. Primers used in this study.

Primer	Sequence and Description
1483promF	5’-GAGAATTCGTCCCACGTGTCGTTCT-3’
	binds 379 bp upstream of *secY* (*msmeg_1483*), contains engineered EcoRI site, used to amplify *secY* promoter region
1483promR	5’-GAGAATTCCCGTTCGCAGCGACGAGATG-3’
	binds at bp 34 of *secY* (*msmeg_1483*) coding sequence, contains engineered EcoRI site, used to amplify *secY* promoter
secYFNotI	5’-GAGCGGCCGCGTCCCACGTGTCGTTCTC-3’
	binds 379 bp upstream of *secY* (*msmeg_1483*), contains an engineered NotI site, used to amplify *secY* with its native promoter
secYREcoRV	5’-GAGATATCAGGCGGTCCGAGCAGAAC-3’
	binds 23 bp downstream of *secY* (*msmeg_1483*), contains an engineered EcoRV site, used to amplify *secY* with its native promoter
secY5’RACE-R1	5’-GACCTGCGCGATGCATTGCTGAAC-3’
	binds at bp 153 of *secY* (*msmeg_1483*) coding sequence, used to reverse transcribe *secY* transcript 5’ ends to cDNA for 5’ RLM-RACE
secY5’RACE-R2	5’-GGATTACCAGGCCCAAGGTGAACAG-3’
	binds at bp 79 of *secY* (*msmeg_1483*) coding sequence, used to amplify *secY* transcript 5’ ends for 5’ RLM-RACE
secY5’RACE-R3	5’-GTGGGTCCAGCGTCAATGGTCAAG-3’
	binds 51 bp upstream of *secY* (*msmeg_1483*), used to amplify *secY* transcript 5’ ends for 5’ RLM-RACE
secYF1	5’-GTGCTTTCGGCTTTCATCTC-3’
	binds at bp 1 of *secY* (*msmeg_1483*) coding sequence, used for quantitative RT-PCR
secYR1	5’-CGCGATGCATTGCTGAAC-3’
	binds at bp 147 of *secY* (*msmeg_1483*) coding sequence, used for quantitative RT-PCR
rpoBF	5’-GTCTCTAGCCAGAGCAAGTC-3’
	binds at bp 25 of *rpoB* (*msmeg1367*) coding sequence, used for quantitative RT-PCR
rpoBR	5’-TCGAAGGAATCCGTCTGAAC-3’
	binds at bp 158 of *rpoB* (*msmeg1367*) coding sequence, used for quantitative RT-PCR