

Table S4. A complete list of known and novel sRNA regions predicted by sRNAscanner.

Bacterial strain (GenBank Acc. No)	<i>Known</i> (CSS > 14)	<i>Novel</i> (CSS > 14)	<i>Reference</i>
<i>Bacillus anthracis</i> Ames (AE016879)	49	535	File S1 Sheet 1-2
<i>Clostridium tetani</i> E88 (AE015927)	27	285	File S1 Sheet 3-4
<i>Chlamydia trachomatis</i> D/UW-3/Cxa (AE001273)	1	27	File S1 Sheet 5-6
<i>Helicobacter pylori</i> 26695 (AE000511)	2	107	File S1 Sheet 7-8
<i>Mycobacterium tuberculosis</i> CDC1551 (AE000516)	0	1	File S1 Sheet 9-10
<i>Pseudomonas aeruginosa</i> PAO1 (AE004091)	3	17	File S1 Sheet 11-12
<i>Salmonella enterica</i> Typhi CT18 (AL513382)	15	175	File S1 Sheet 13-14
<i>Staphylococcus aureus</i> N315 (BA000018)	17	253	File S1 Sheet 15-16
<i>Streptococcus pneumoniae</i> TIGR4 (AE005672)	9	190	File S1 Sheet 17-18
<i>Streptococcus pyogenes</i> M1 GAS (AE004092)	4	162	File S1 Sheet 19-20
<i>Yersinia pestis</i> KIMa (AE009952)	7	287	File S1 Sheet 21-22
<i>Salmonella enterica</i> Typhimurium LT2 (NC_003197)	24	135	File S1 Sheet 23-24
<i>Escherichia coli</i> K12-MG1655 (NC_000913)	22	170	File S1 Sheet 25-26