

Figure S1. Bioinformatics analysis of total RNA from *Helicobacter pylori*. Total RNA was sequenced by using Illumina HiSeq2000. According to their genomic locations, tags in the IGR and AM were retained. Candidate sRNAs were defined as those with >20 reads. sRNA, small non-coding RNA; IGR, intergenic region; AM, antisense mRNA.

