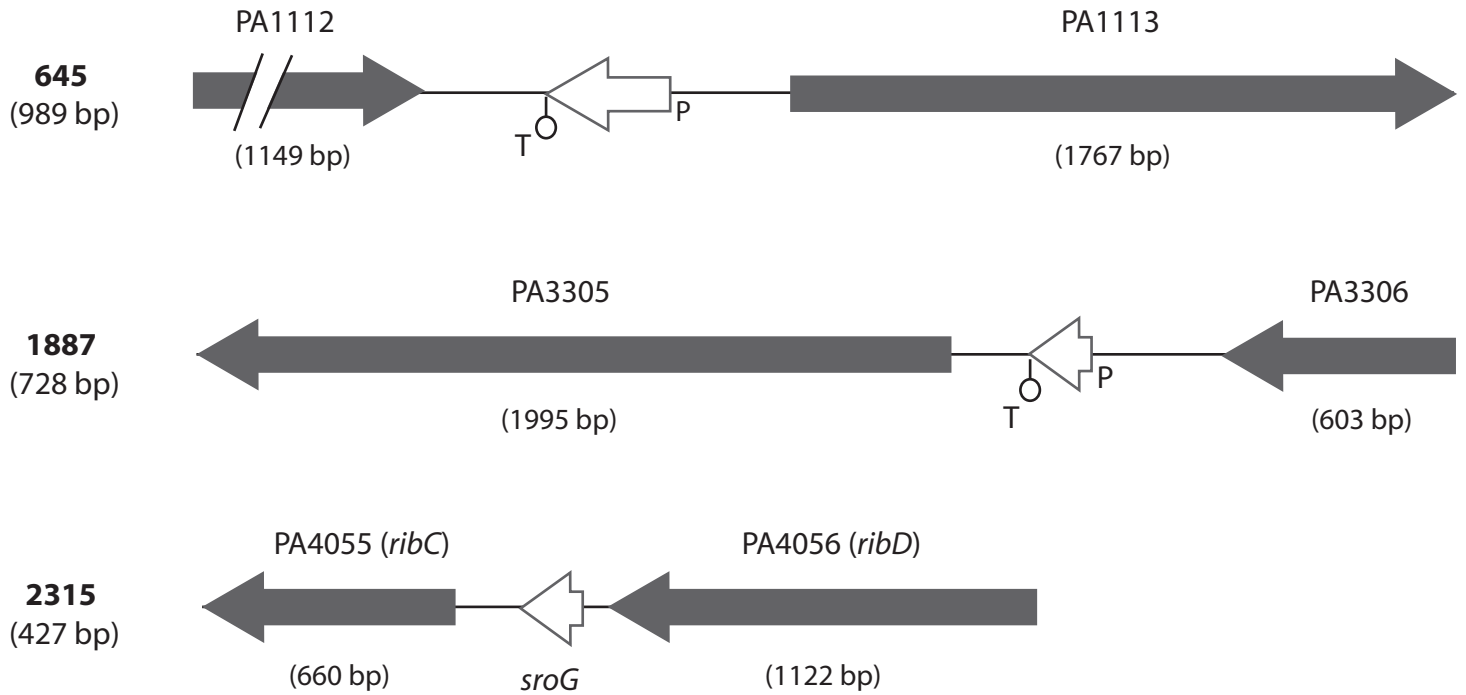


Intergenic
region:

0.5 kb 1 kb



Additional file 2: Predicted coordinates of three sRNA genes (645, 1887, 2315). These predictions are based on an analysis of putative RpoD-dependent promoters (P), putative ρ -independent terminators (T) or sequence homology (in case of *sroG*; Vogel *et al.*, 2003). The coordinates predicted are: 1205239 - 1205035 (IgR 645); 3705522 - 3705315 (IgR 1887); 4537006 - 4536845 (IgR 2315).