

SUPPLEMENTARY MATERIAL

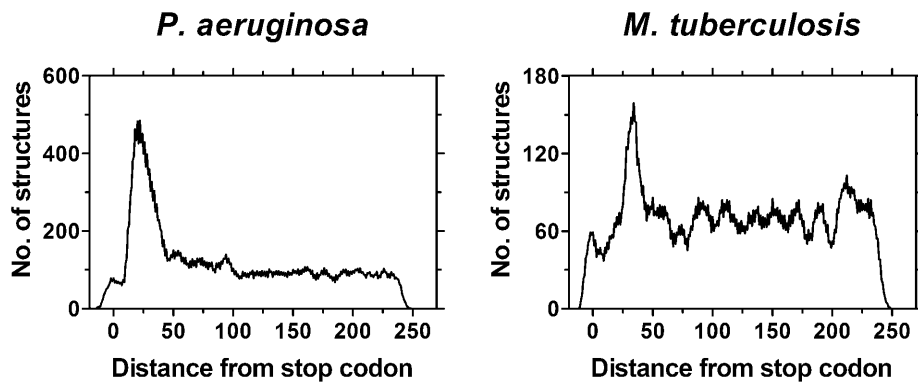


Figure S1. Distribution of Best Candidate structures in the region downstream (-20 to +270 nt from the stop codon) of coding regions. These regions are defined without restriction by the start position of downstream genes.