



Fig. S3. Genomic map and distribution of predicted TA systems of *P. syringae*. TA systems in the type strain DC3000 chromosome and plasmid are presented. TA systems not predicted in DC3000 but in other strains are presented as other TAs with a representative strain name attached. Except for the *abiEii* and *hok-sok* system, all TA systems belong to the type II class.