FIG S1 eBURST of 1015 *Yersinia* isolates. A minimum number of identical loci of a group was set to zero, thus the entire population of three pathogenic *Yersinia* was included. The solid line represented one locus difference. *Y. enterocolitica , Y. pseudotuberculosis* and *Y.similis* subgroup were presented according to the algorithm. Group founders were marked in blue.