



Figure S1: Relationships between *Xcc* strains analysed by MLSA or based on diversity of the type III secretomes. (A) Neighbor-joining tree of the concatenated sequences of *efP* and *glnA*. (B) Dice similarity indices calculated from presence/absence of gene coding for type III secreted proteins were used to construct a neighbor-joining tree showing the genetic relationships between *Xcc* predicted type III secretomes. Bootstrap values higher than 40% are indicated. *Xcc* strains which genome sequence is available are boxed in black. On the right, brackets indicate AFLP-based subgroups of *Xcc* (Figure 1). *X. campestris* pv. *raphani* strain CFBP5828 was used to root the tree.

Figure S1
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