

Supplementary Table S1. Information of mapping statistics for all the transcriptome reads to *Pst*-78 reference genome.

	Resistant bulk (count)	Susceptible bulk (count)	Resistant (%)	Susceptible (%)
Counted fragments	370,937	352,479	10.18	11.91
Unique fragments	342,405	329,772	9.40	11.15
Common fragments	28,532	22,707	0.78	0.77
Uncounted fragments	3,271,563	2,606,435	89.82	88.09
Total fragments	3,642,500	2,958,914	100	100

Note: Counted fragments represent reads that mapped to reference *Pst* genome, unique fragments mapped to single position on the genome, while non-specific (aligned to a few regions) had between 1-5 hits, uncounted fragments didn't have any hits in the reference *Pst* genome.