

**Table S6 EST libraries produced to facilitate gene calling**

Libraries	Pathogenicity	Conditions	Reads	Spans	Average Sequence Identity %
Pt-1C(BFP)	Yes-Race 1	In planta	10,083	3,088	
Pt-1C(BFP)	Yes-Race 1	Mixed culture	10,444	12,198	
SO3-P	Yes-Race 9	Mixed culture	10,185	12,115	98.08
SD20-NP	No-Race 4	Mixed culture	10,328	8,195	98.2