

Position	974*	1017	1128	1200	1395	1482	1542	1560	1770	1816	1824	1893	1899
Ppat5Ahnd1	-	-	T	C	A	-	-	C	-	T	-	T	T
Ppat5Hind2	-	-	-	-	-	-	-	-	T	-	T	-	-
Ppat5Aswa1	C	-	-	-	-	-	-	-	-	-	-	-	-
Ppat5Emco5	-	-	-	-	-	-	-	-	-	-	-	-	-
Ppat5Hiks1	-	A	-	-	-	-	-	-	-	-	T	-	-
Ppat5Goco1	-	A	-	-	-	-	-	-	-	-	T	-	-
Ppat5Waco5	-	A	-	-	-	-	-	-	-	-	T	-	-
Ppat5Hind4	-	-	T	-	A	-	-	C	-	T	-	T	T
Ppat5Cala2	-	-	T	-	A	-	-	C	-	T	-	T	T
Ppat5Emoy2	-	A	-	-	-	-	-	-	-	-	T	-	-
Ppat5Noks1	-	-	T	-	A	-	A	C	-	T	-	T	T
Ppat5Wela3	-	-	-	-	-	-	-	-	-	-	-	-	-
Ppat5Bico1	-	-	-	-	A	-	-	C	-	T	-	T	-
Ppat5Emwa1-A	-	A	T	-	-	T	-	C	-	T	-	-	T
Ppat5Emwa1-B	-	-	T	-	A	-	-	-	-	T	T	T	-
Ppat5Maks9	-	-	-	-	-	-	-	-	-	-	T	-	-
Consensus	G	G	C	T	G	C	G	T	C	C	C	C	C

Supplemental Table 1. *Ppat5* polymorphisms in 15 isolates of *H. parastitica* (*Ppat5Cand5* was not analysed). - symbolises no nucleotide change, the allele has the nucleotide of the consensus sequence in this position. \* indicates a replacement change, all other polymorphisms are silent.