

Supplementary Information

596 Supplementary Figures

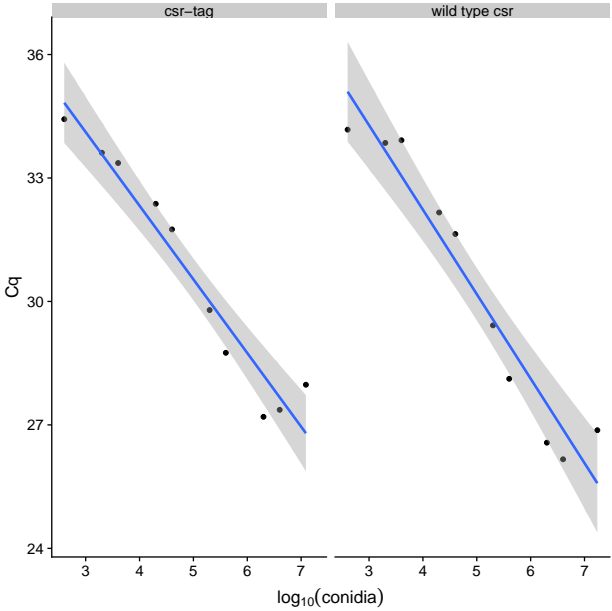


Figure S1: Correlation between number of conidia used in DNA extraction (log-scale) and number of PCR cycles required to detect the PCR-product. Differences between the alleles are not significant.

Supplementary Tables

Table S1: Primers used in the study.

Name	Sequence
csrL-f	TGC CAT GTT CTT CTT GAG CC
csrL-r	CCC ATG TTT GCG CGG ACC TGG AGA AGC GGC TGG ACT TAC ACT ATT ACA TGA ATT CGG ATG TTT GCG AAA AAG CTC TGG C
csrR-f	CAC TGC AAC TTT CTC CTG CGC CAG AGC TTT TTC GCA AAC ATC CGA ATT CAT GTA ATA GTG TAA GTC CAG CCG CTT CTC
csrR-r	GAC AAT GGT GGG CTT CTT GG
csr-hrm-f	CGT CAT CTC TCA AGC CCA CT
csr-hrm-r	GAG AAG CGG CTG GAC TTA CA