



Figure S1. Analysis of *MDR1* expression by quantitative real-time RT-PCR

(A) Relative fold change in *MDR1* expression in the fluconazole-resistant isolates F5 and G5 and two independently derived *mrr1*Δ mutants of each parental strain as compared with the matched fluconazole-susceptible isolates F1 and G1, respectively. Shown are the means and standard errors from three experiments.

(B) Relative fold change in *MDR1* expression in strains expressing mutated *MRR1* alleles (*MRR1*^{F5} and *MRR1*^{G5}) as compared with strains expressing the corresponding non-mutated alleles (*MRR1*^{F2-1} and *MRR1*^{G2-2}, respectively). Shown are the means and standard errors from three experiments performed with two independently constructed strains in each case.