

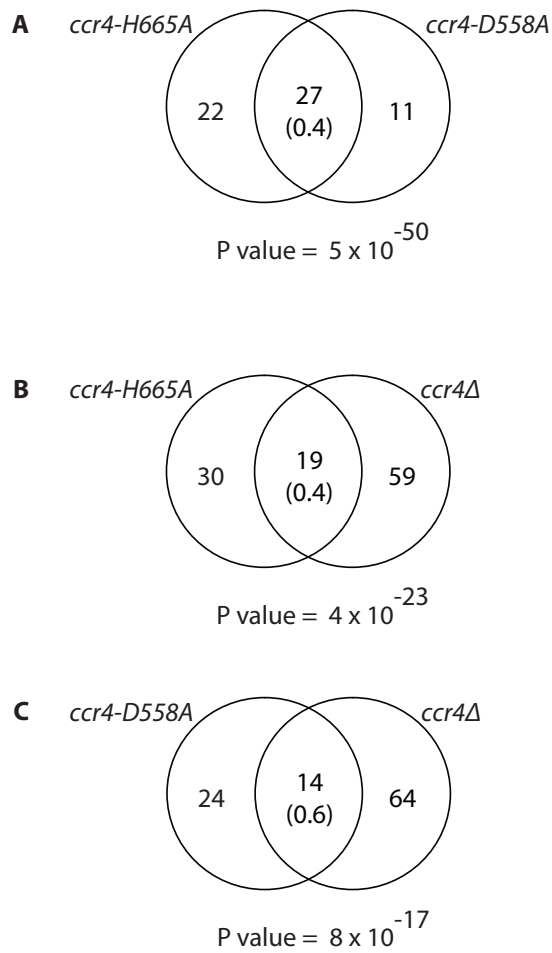


Figure S3. Microarray profiles of *ccr4* catalytic site mutants.

Venn diagrams comparing the genes overexpressed in *ccr4Δ*, *ccr4-H665A* and *ccr4-D558A* mutants. The numbers in parentheses indicate the expected overlap if randomly generated lists of the corresponding sizes were used.