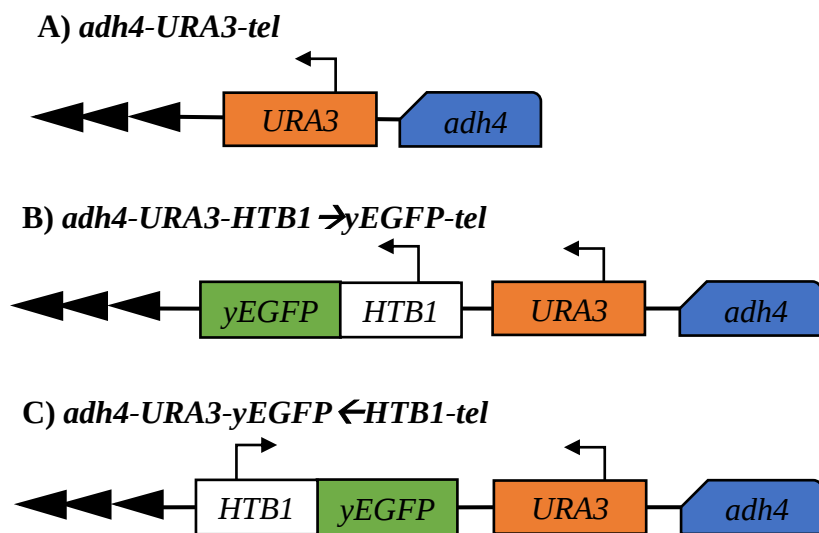
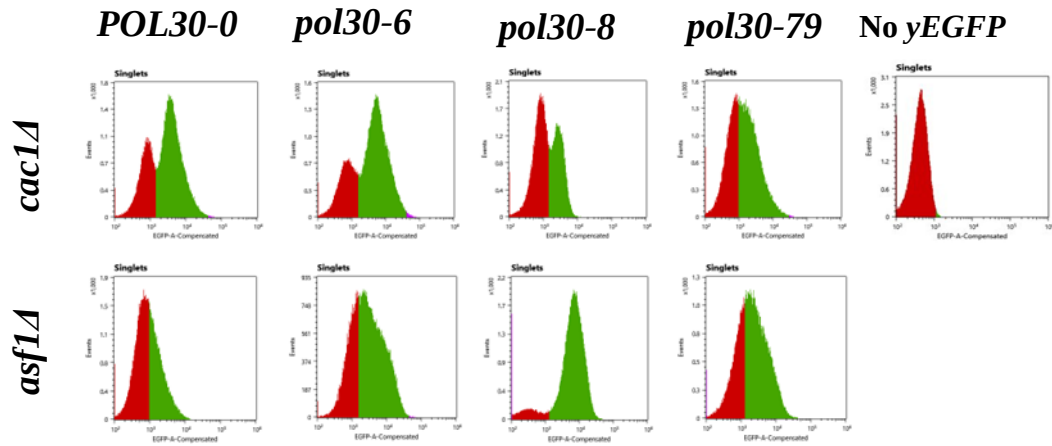


Supplemental Figure 1. Schematic of the reporter fragments used in this study.

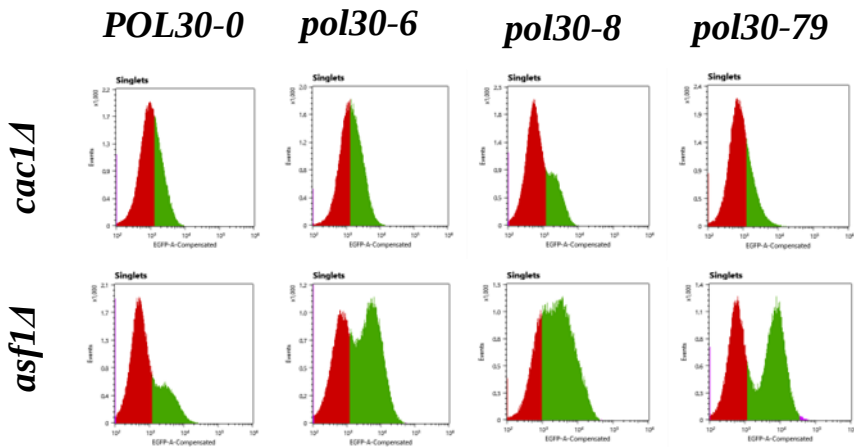


Supplemental Figure 2. yEGFP signal distribution by flow cytometry using A) *adh4-URA3-HTB1* →yEGFP-tel B) *adh4-URA3-yEGFP* ←HTB1-tel fragments. Wild-type to mutant %yEGFP-negative ratio using both fragments are shown in log scale in C) *cac1Δ* and D) *asf1Δ* backgrounds.

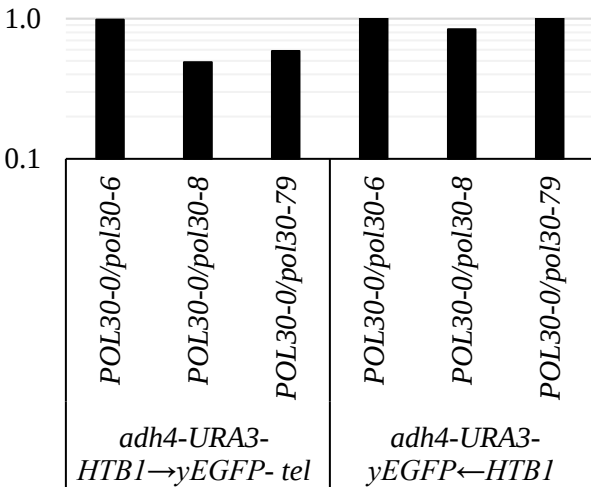
A) *adh4-URA3-HTB1* →yEGFP-tel



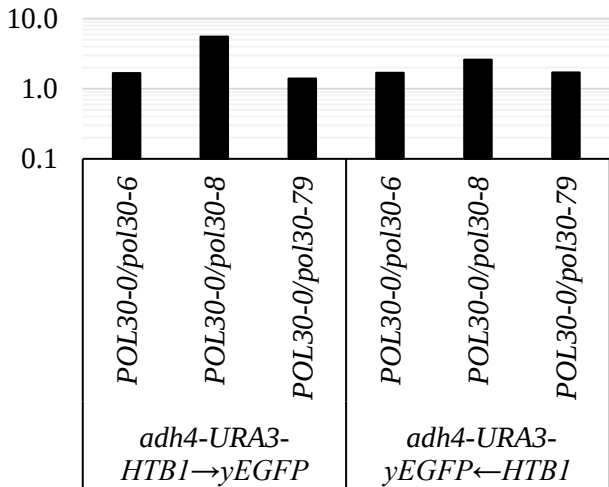
B) *adh4-URA3-yEGFP* ←HTB1-tel



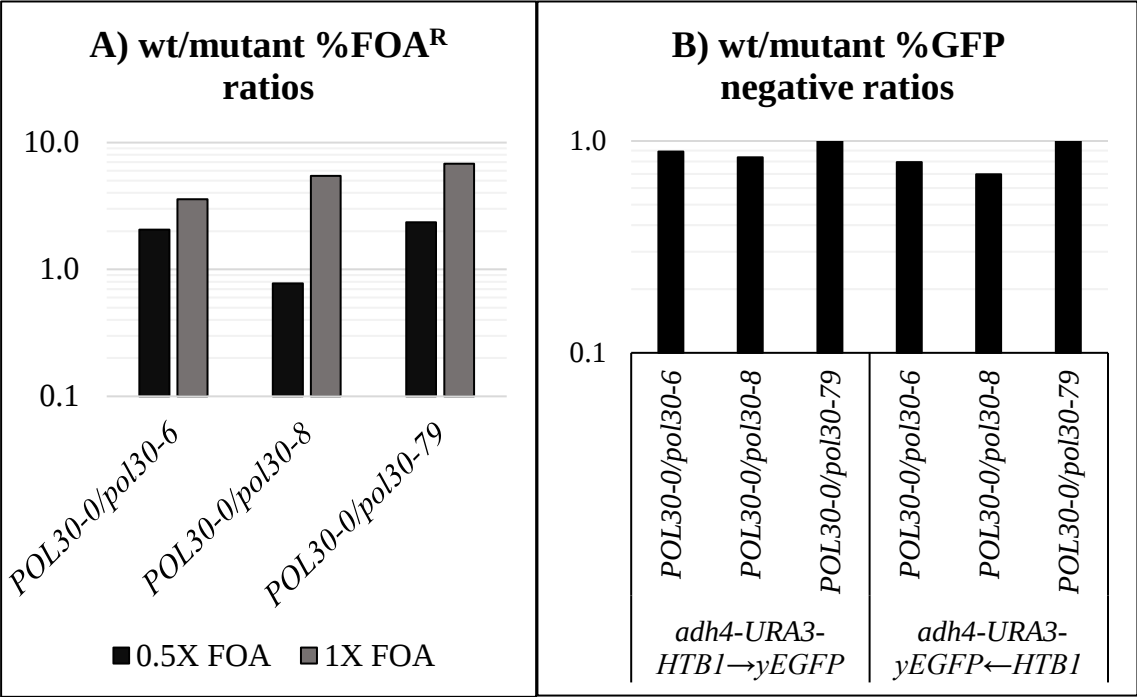
C) wt/mutant ratios in *cac1Δ*



D) wt/mutant ratios in *asf1Δ*



Supplemental Figure 3. Genetic interaction of *RRM3* with *pol30* mutants. Wt/mutant ratio of A) %FOA^R and B) %GFP negative cells are shown in *rrm3Δ* background on a log scale. C) yEGFP flow cytometry distribution image with both fragments in *rrm3Δ* strains



C) Flow cytometry distribution

