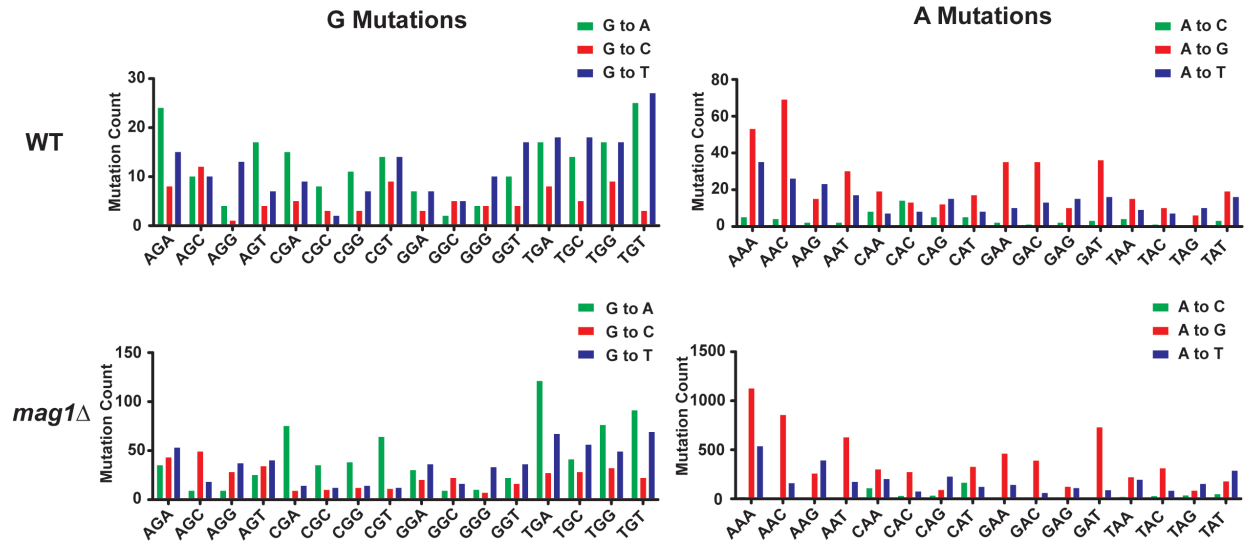


Supplemental Fig. S8



Supplemental Figure S8. The base substitution pattern of MMS-induced mutations in WT and *mag1*Δ yeast was determined for each of 32 possible trinucleotide contexts.