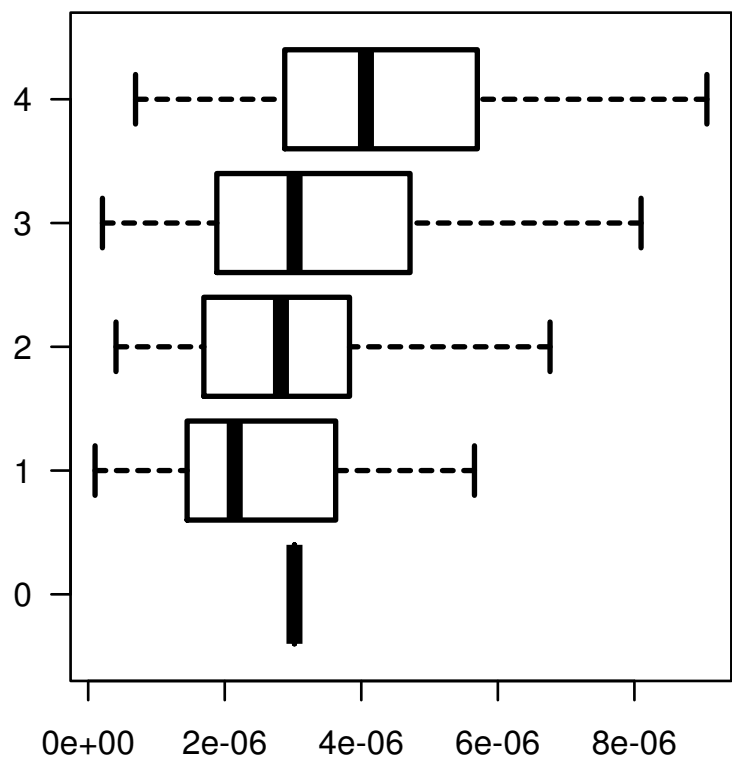


MUTATIONRATE_NONSYNONYMOUS
T_HOLOGY_T_STAGE



MUTATION_RATE
Spearman.Corr= 0.314, P=1.31e-11