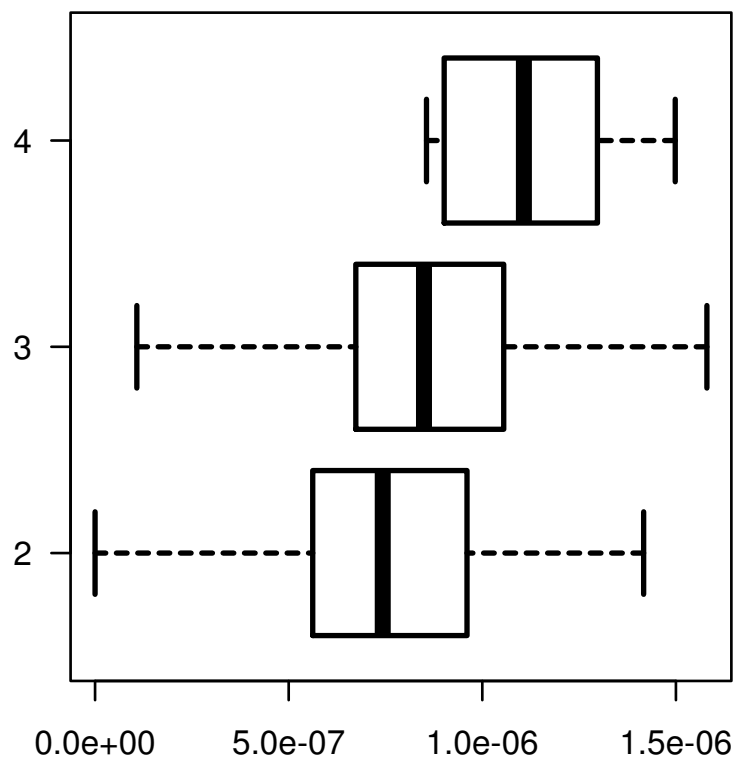


MUTATIONRATE_NONSYNONYMOUS
PATHOLOGY_T_STAGE



MUTATION_RATE
Spearman.Corr= 0.2, P=7.67e-06