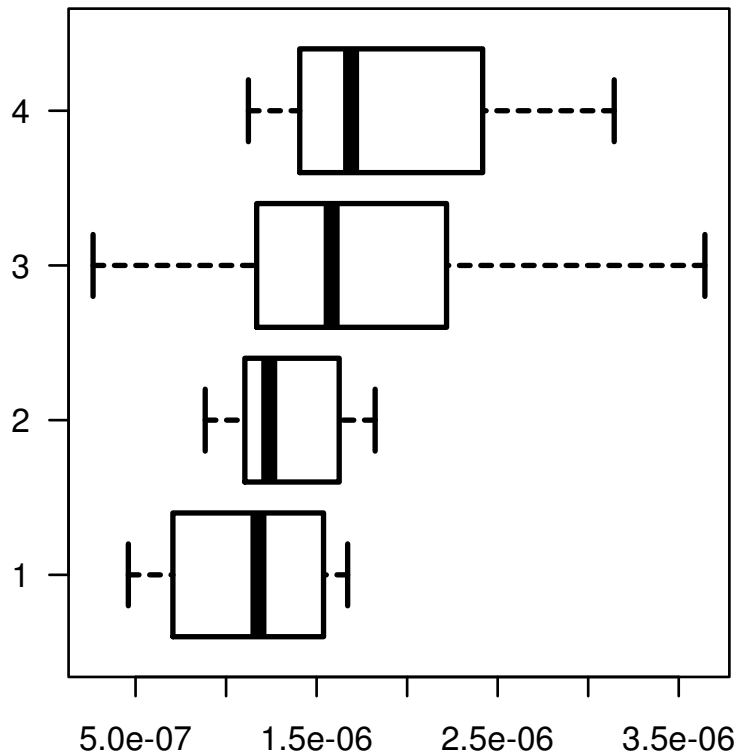


MUTATIONRATE_NONSYNONYMOUS
PATHOLOGY_T_STAGE



MUTATION_RATE
Spearman.Corr= 0.185, P=0.0256