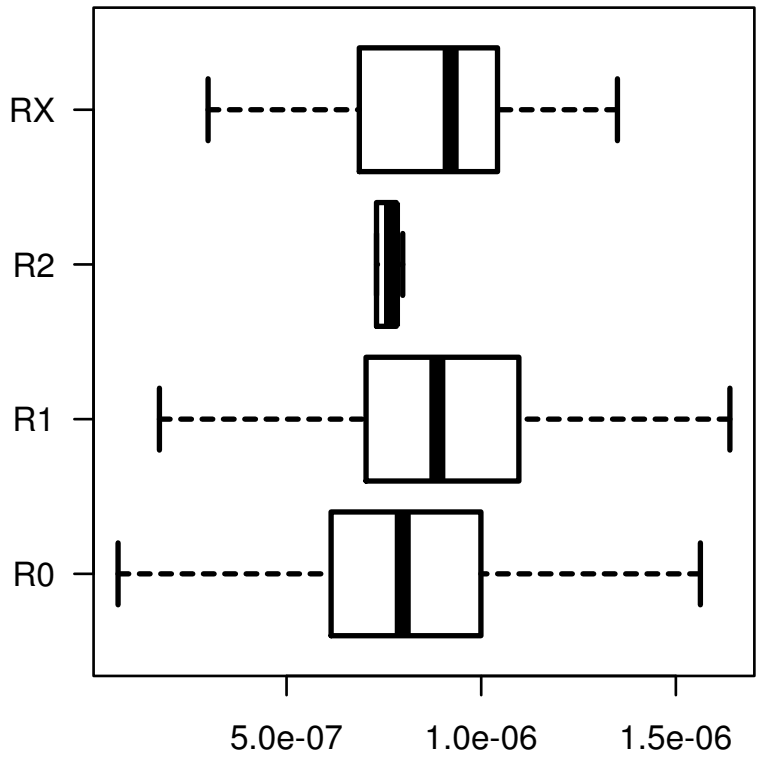


MUTATIONRATE_NONSYNONYMOUS

RESIDUAL_TUMOR



MUTATION_RATE
Kruskal-Wallis P = 0.0338