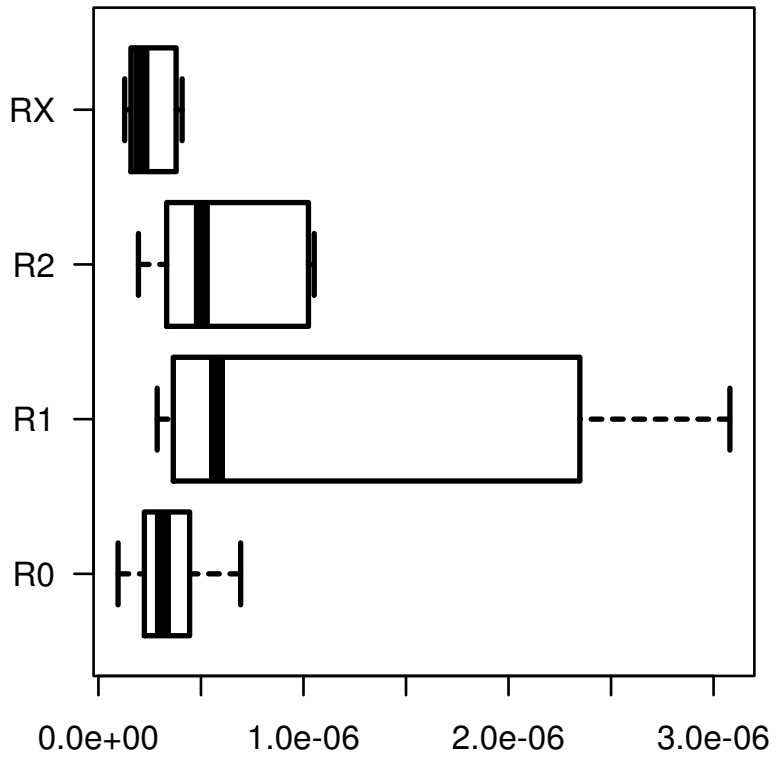


RESIDUAL_TUMOR

MUTATIONRATE_SILENT



MUTATION_RATE
Kruskal-Wallis P = 0.00556