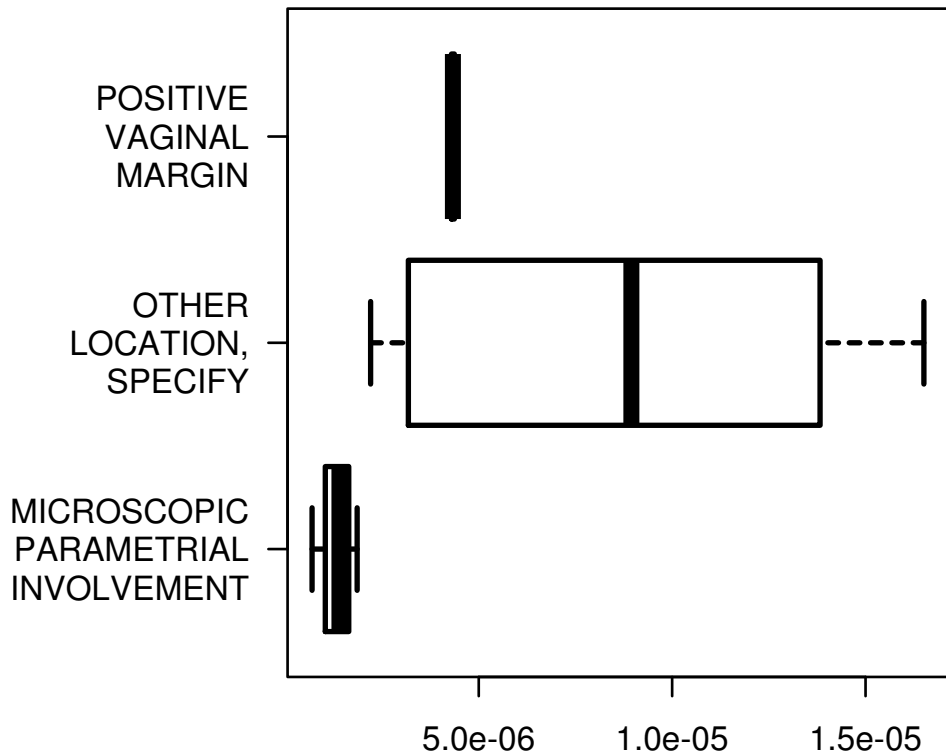


MUTATIONRATE_NONSYNONYMOUS

POSITIVE_MARGINS



MUTATION_RATE
Kruskal-Wallis P = 0.0207