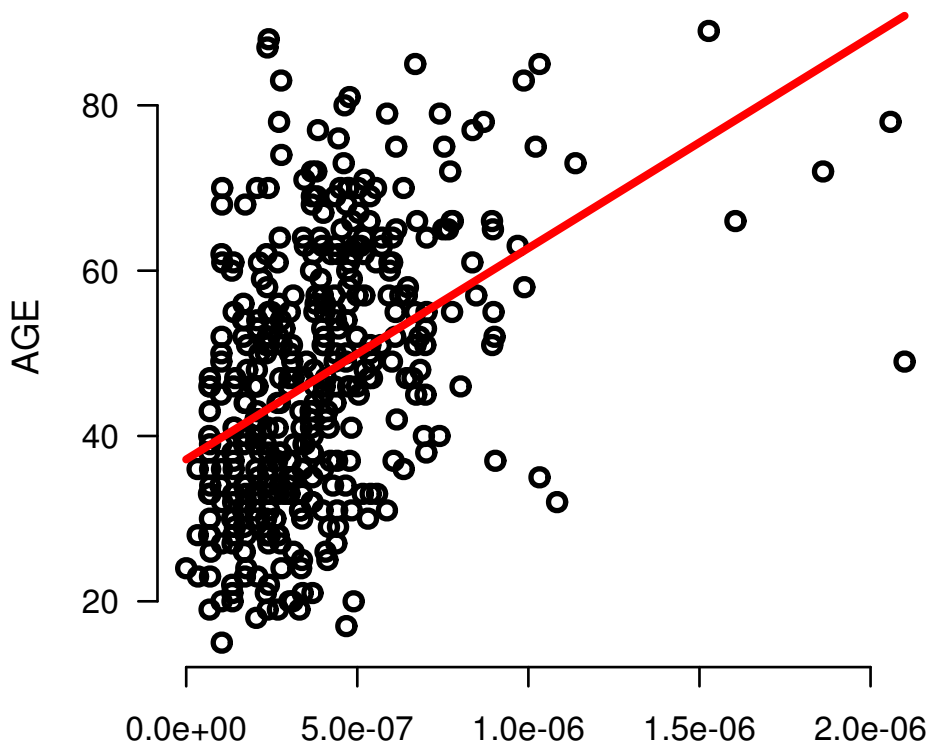


MUTATIONRATE_NONSYNONYMOUS



MUTATION_RATE
Spearman.Corr= 0.468, P=3.83e-23