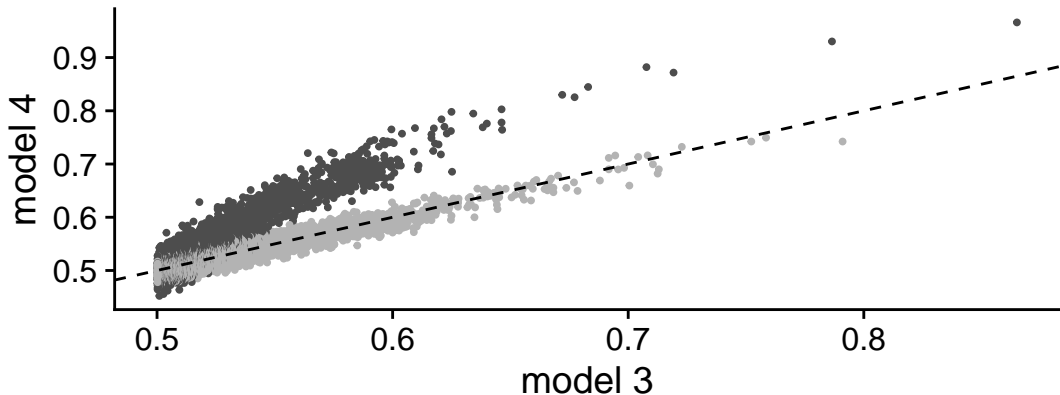


posterior probabilities for all evaluated replacements (HIVChina – pol)



within predicted epitope • no • yes