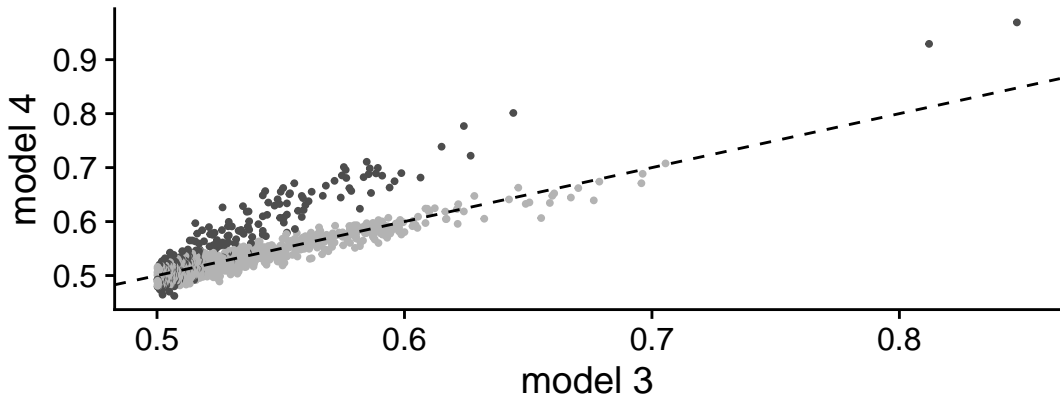


posterior probabilities for all evaluated replacements (HIVChina – Protease)



within predicted epitope • no • yes