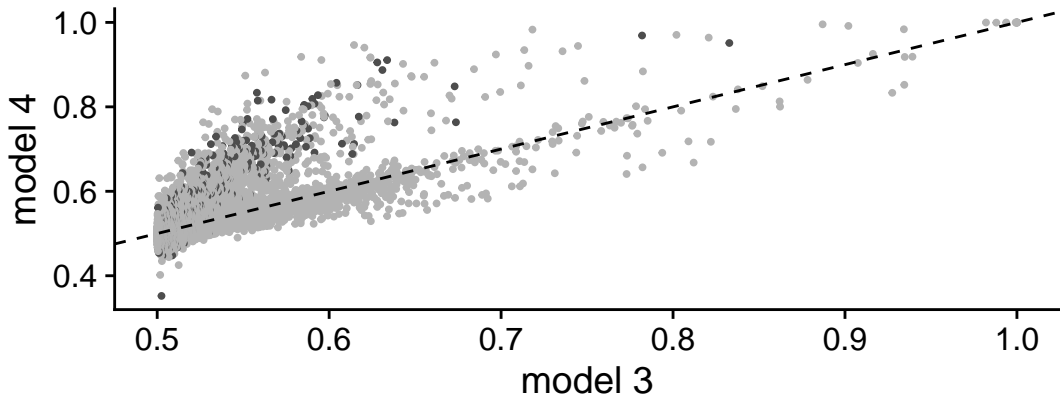


posterior probabilities for all evaluated replacements (HIVBrumme – vif)



within predicted epitope • no • yes