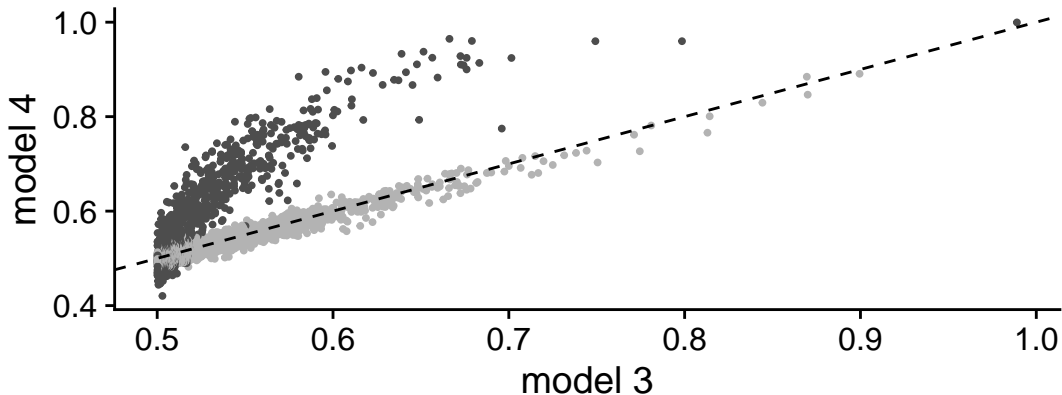


# posterior probabilities for all evaluated replacements (HDVRoggendorf – delta)



within predicted epitope • no • yes