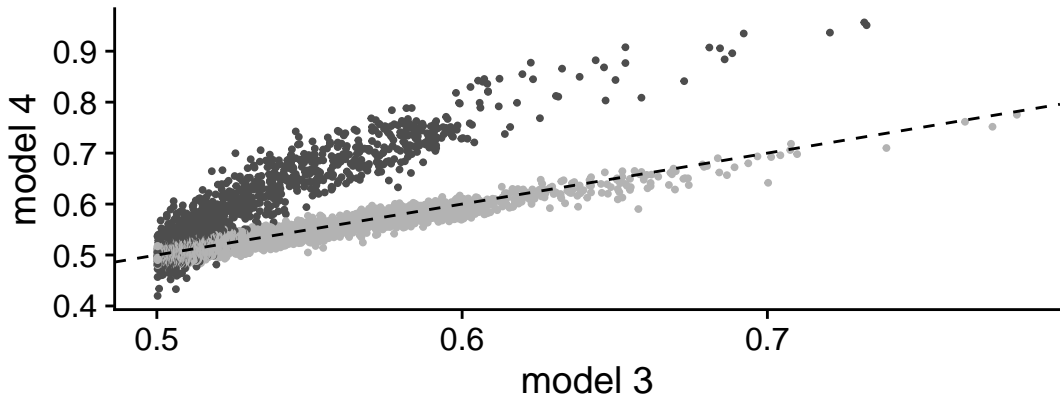


# posterior probabilities for all evaluated replacements (HIVChina – gag)



within predicted epitope   •   no   •   yes