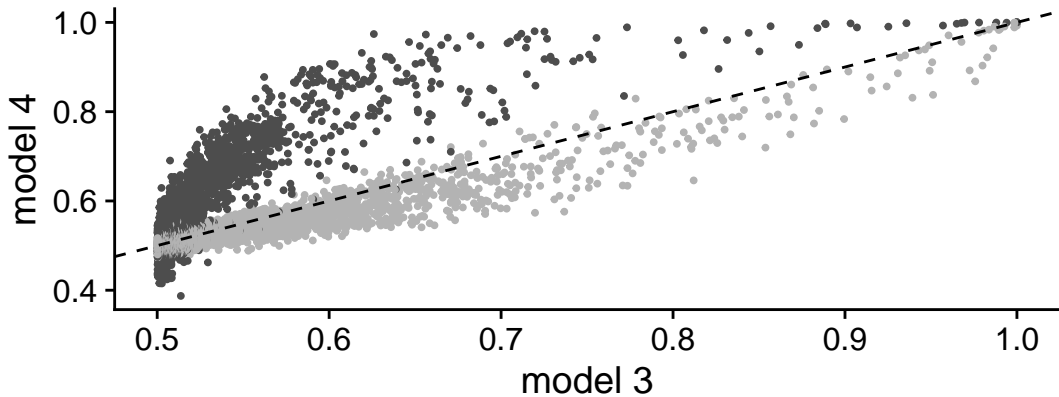


posterior probabilities for all evaluated replacements (HBVDüsseldorf – HBe)



within predicted epitope • no • yes