posterior probabilities for all evaluated replacements (Arevir – RT)
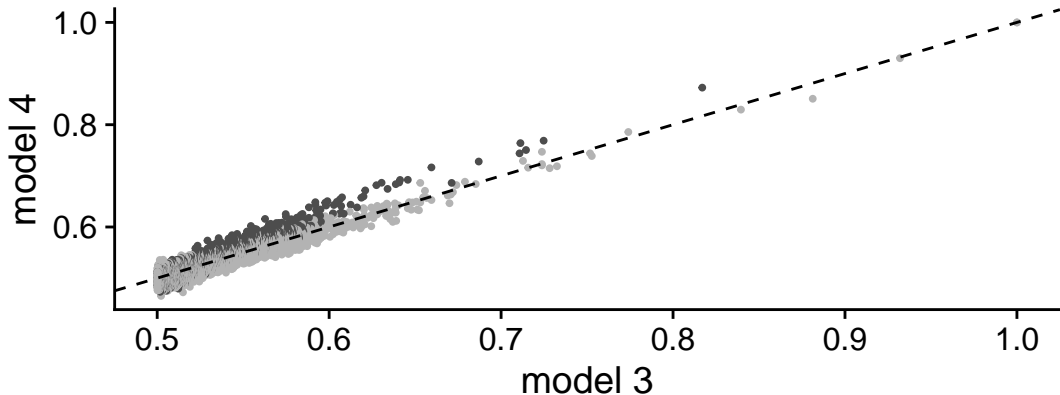
within predicted epitope • no • yes