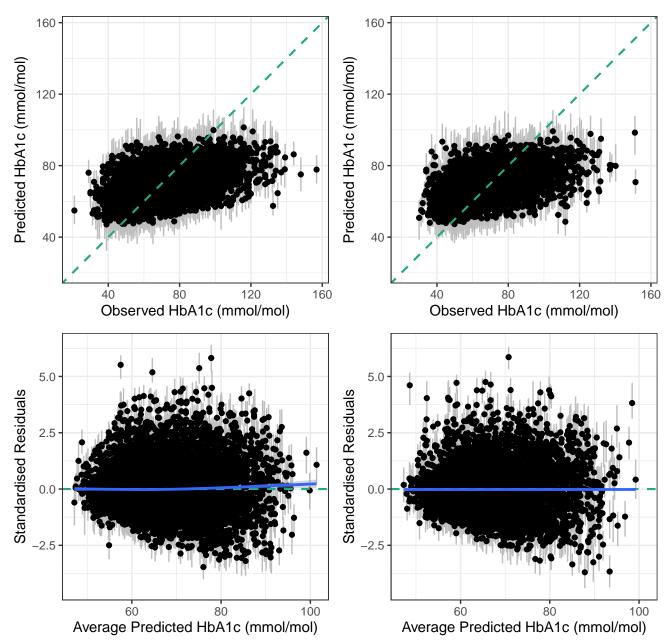
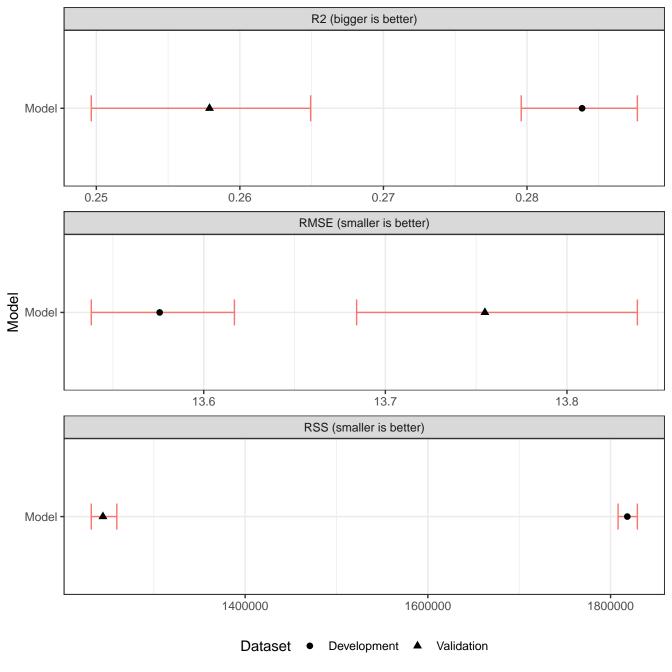
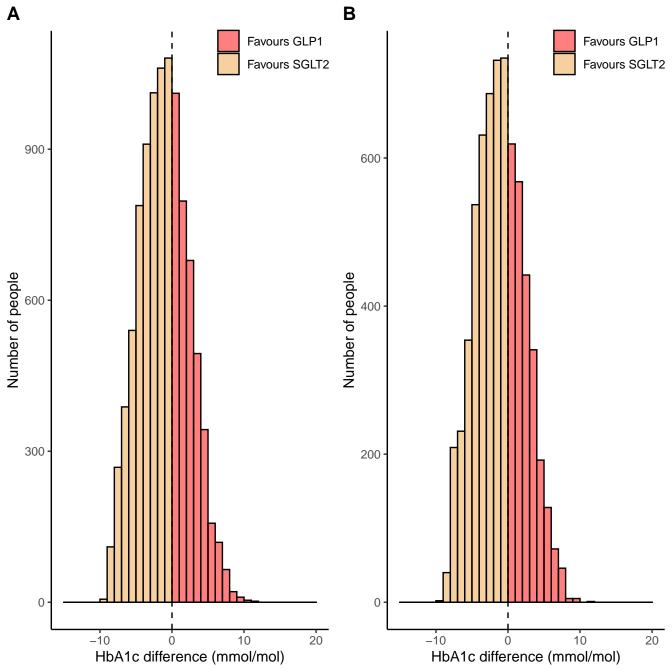
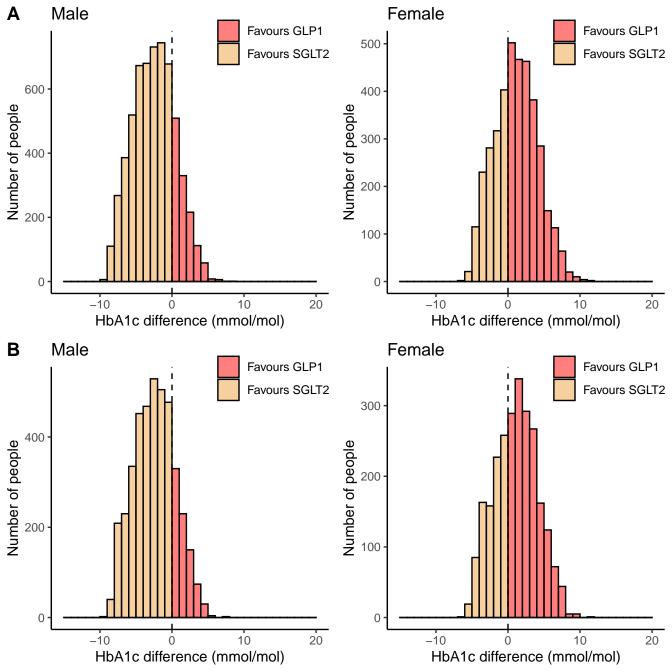
Residuals of Model 7

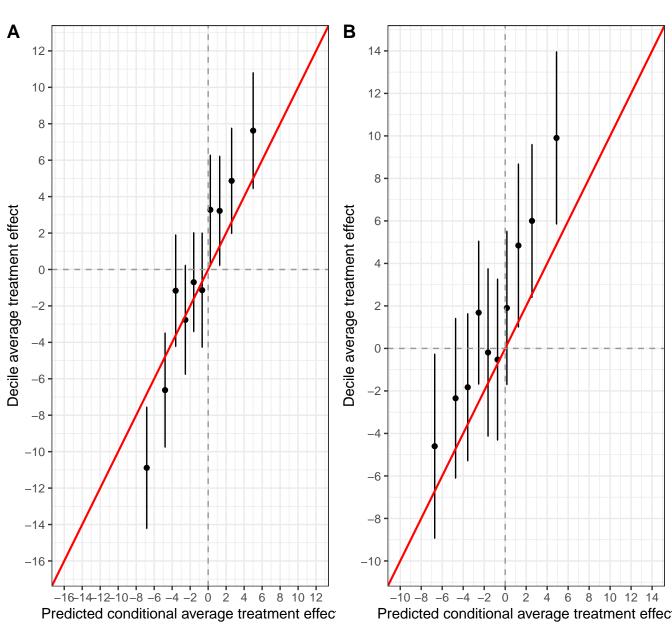




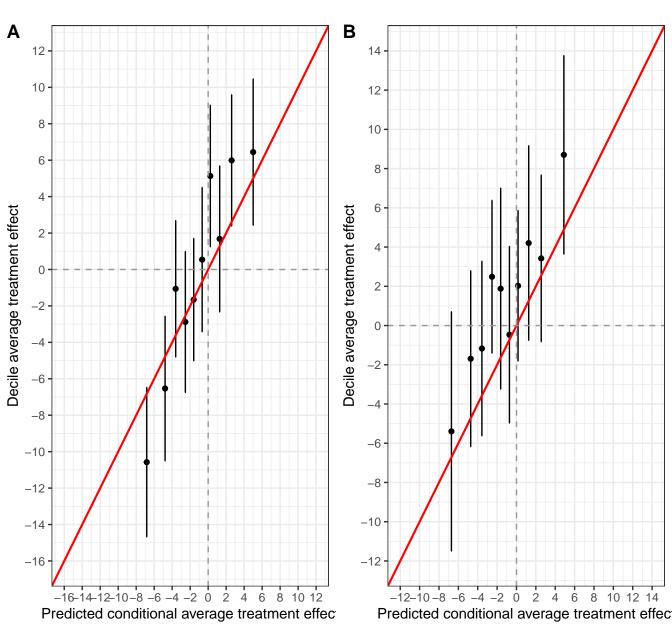




Effects validation lm(hba1c~drugclass+prop_score)



Effects validation prop score matching



Effects validation prop score inverse weighting

