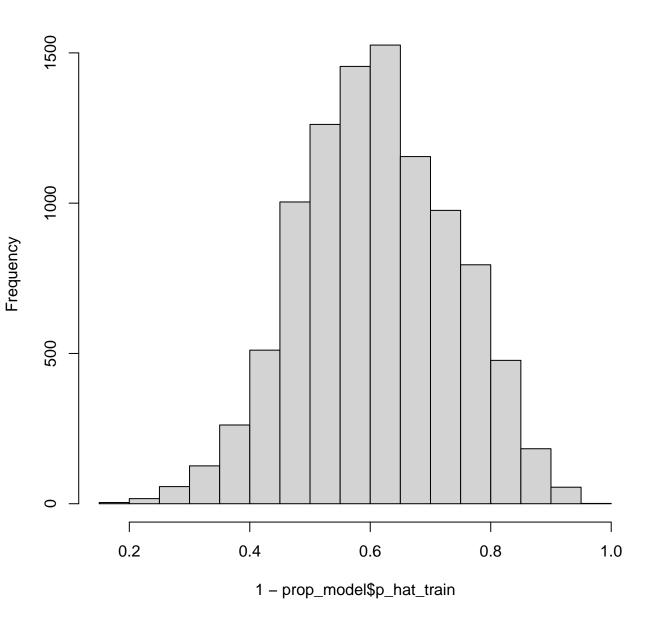
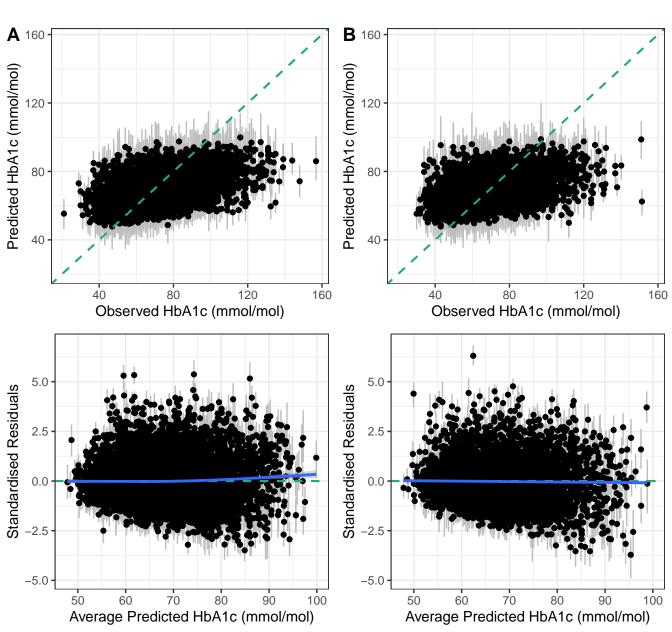
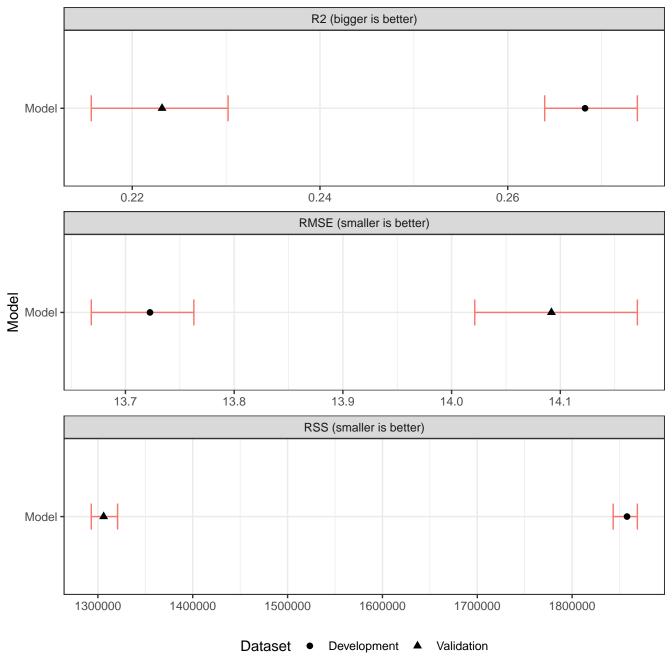
Histogram of 1 – prop_model\$p_hat_train

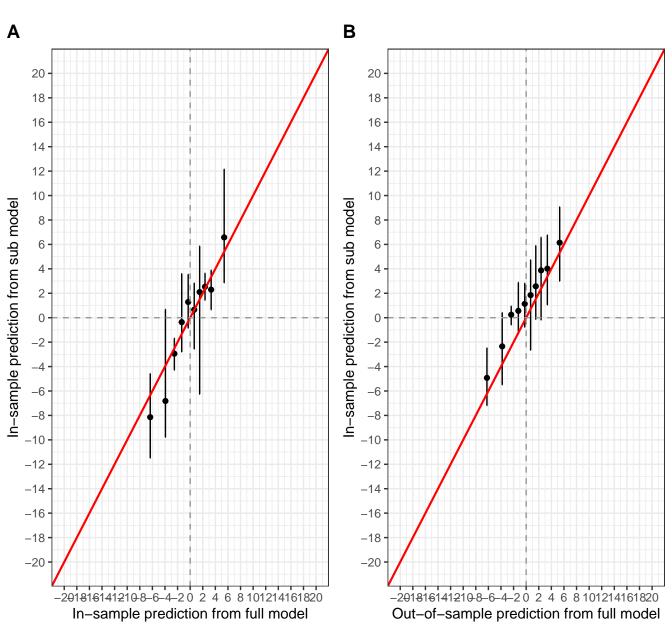


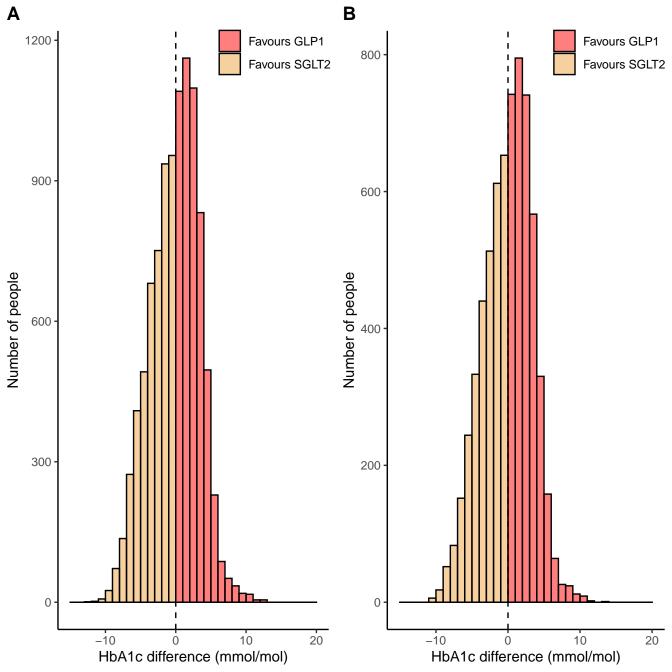
Residuals of Model 2

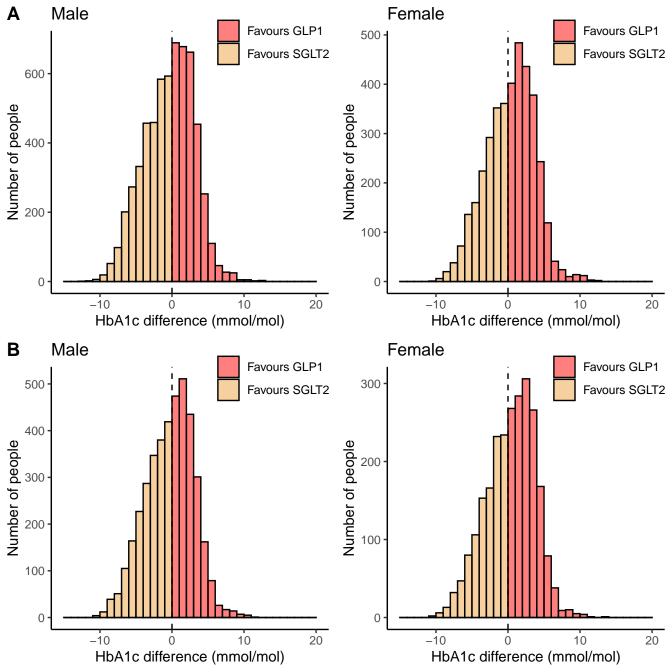




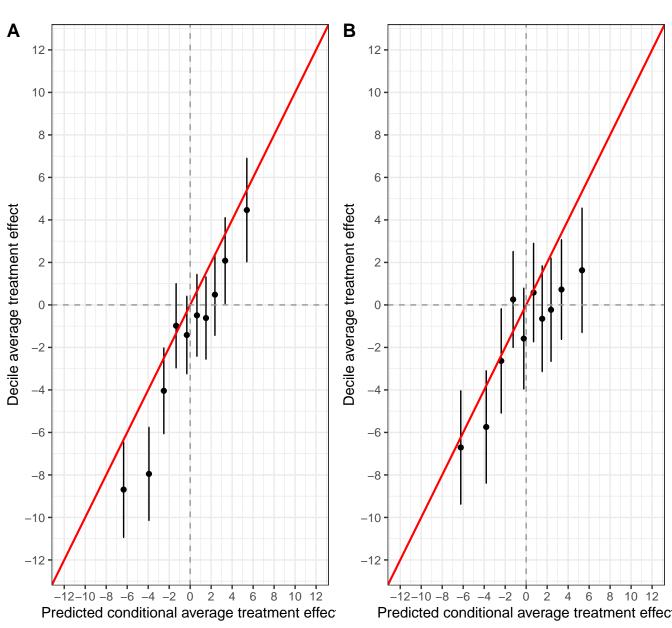
Effect submodels



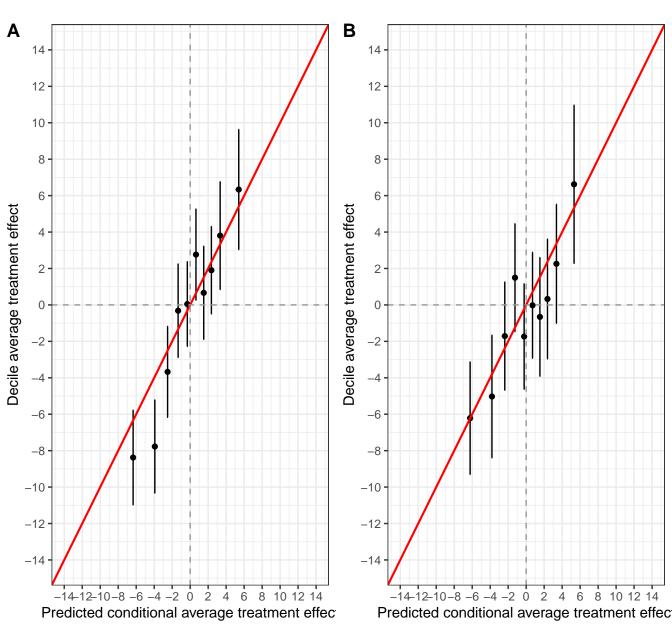




Effects validation lm(hba1c~drugclass+prop_score)



Effects validation prop score matching



Effects validation prop score inverse weighting

