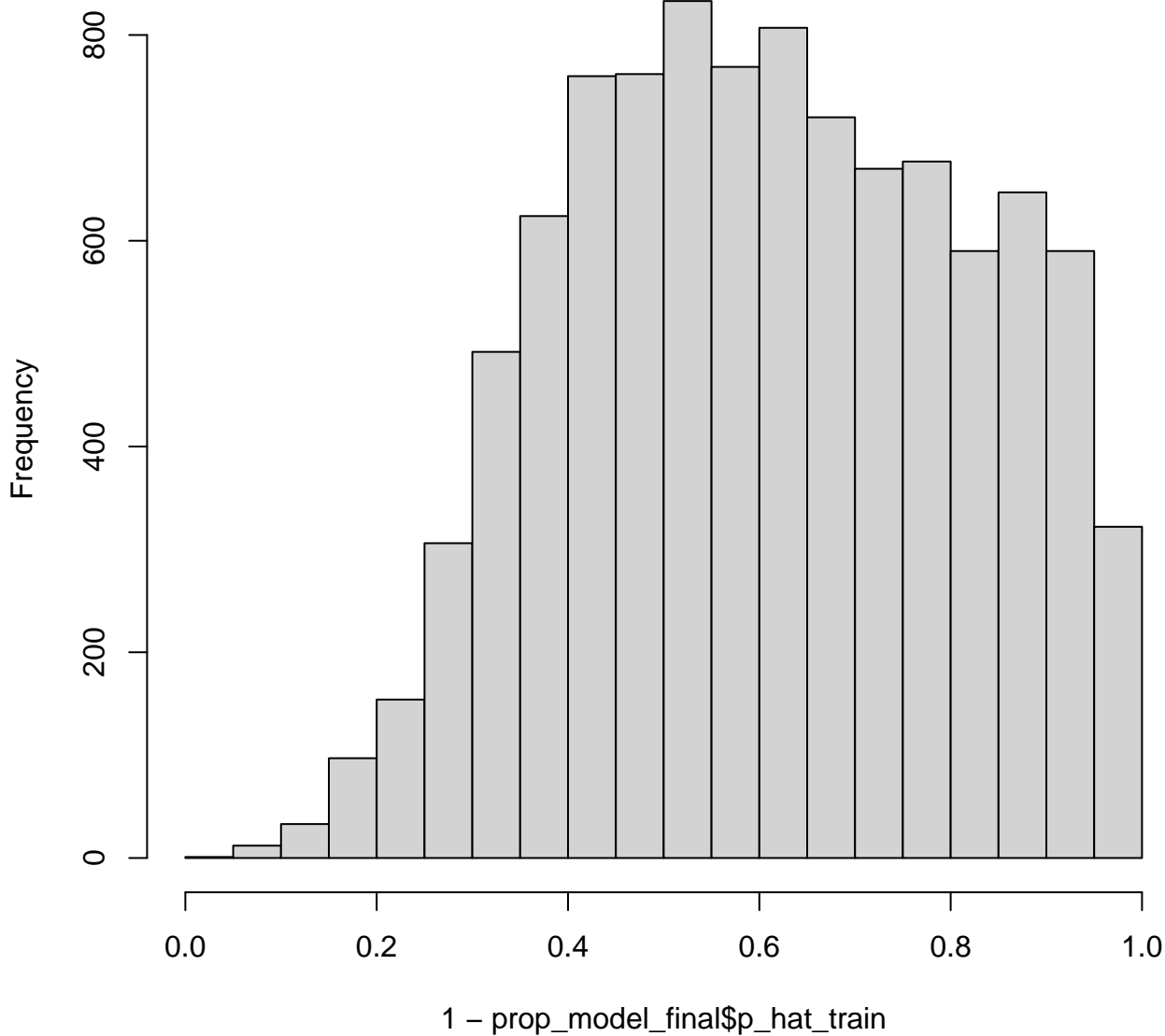
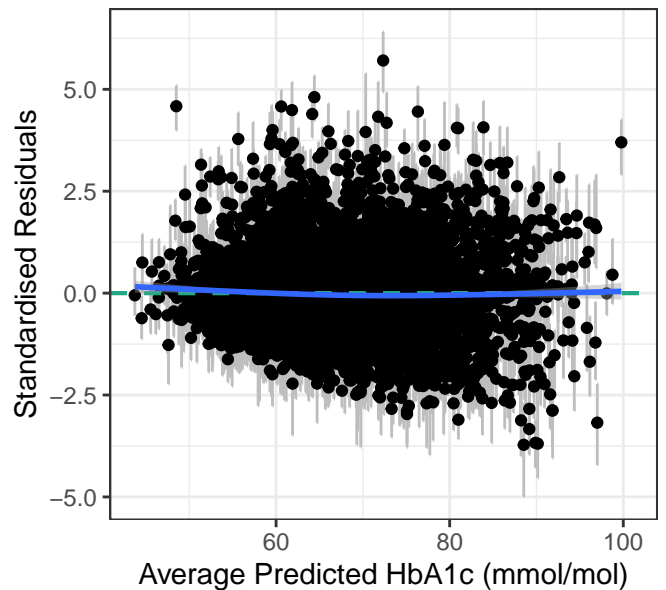
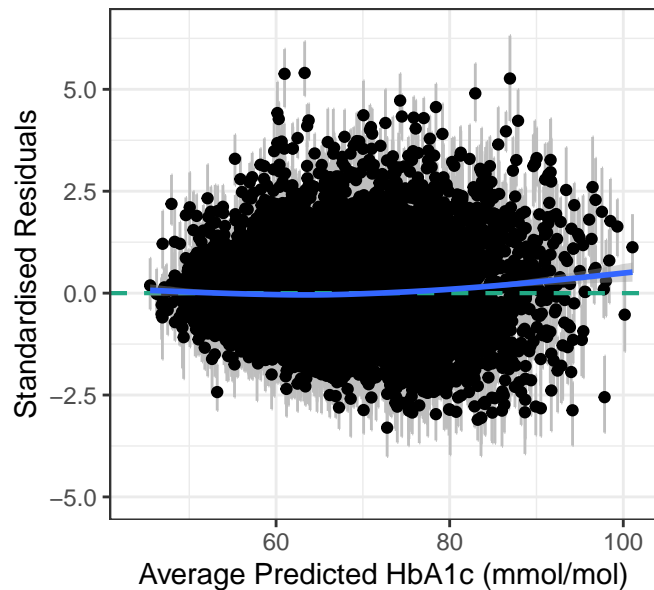
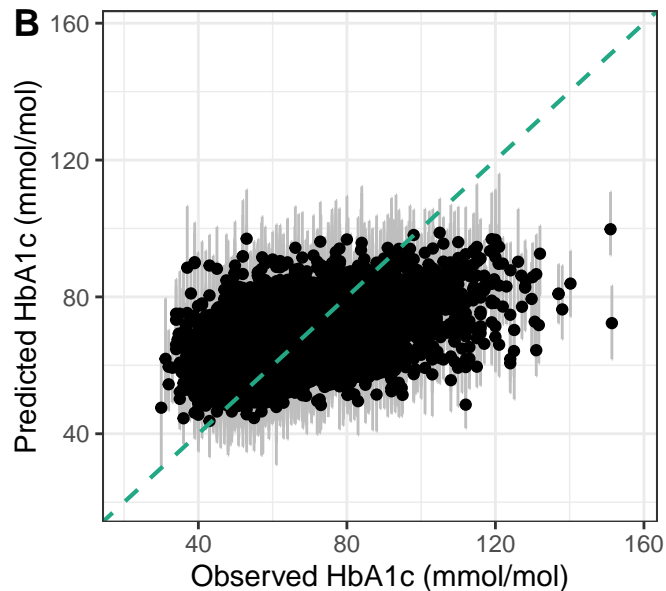
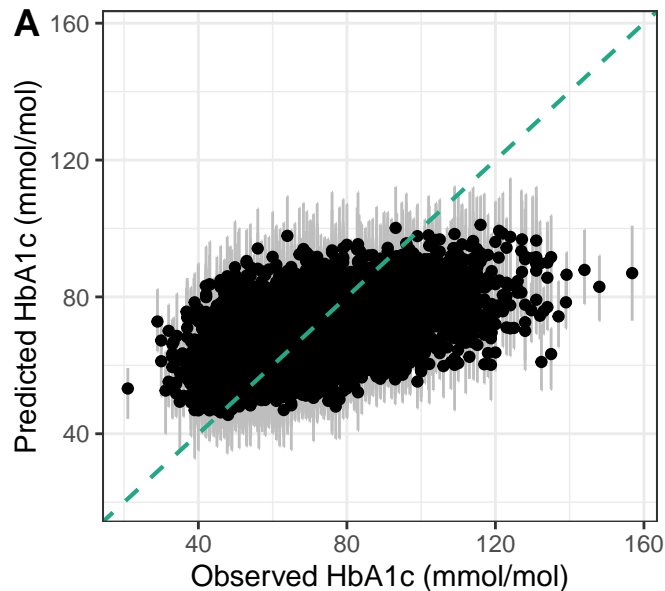
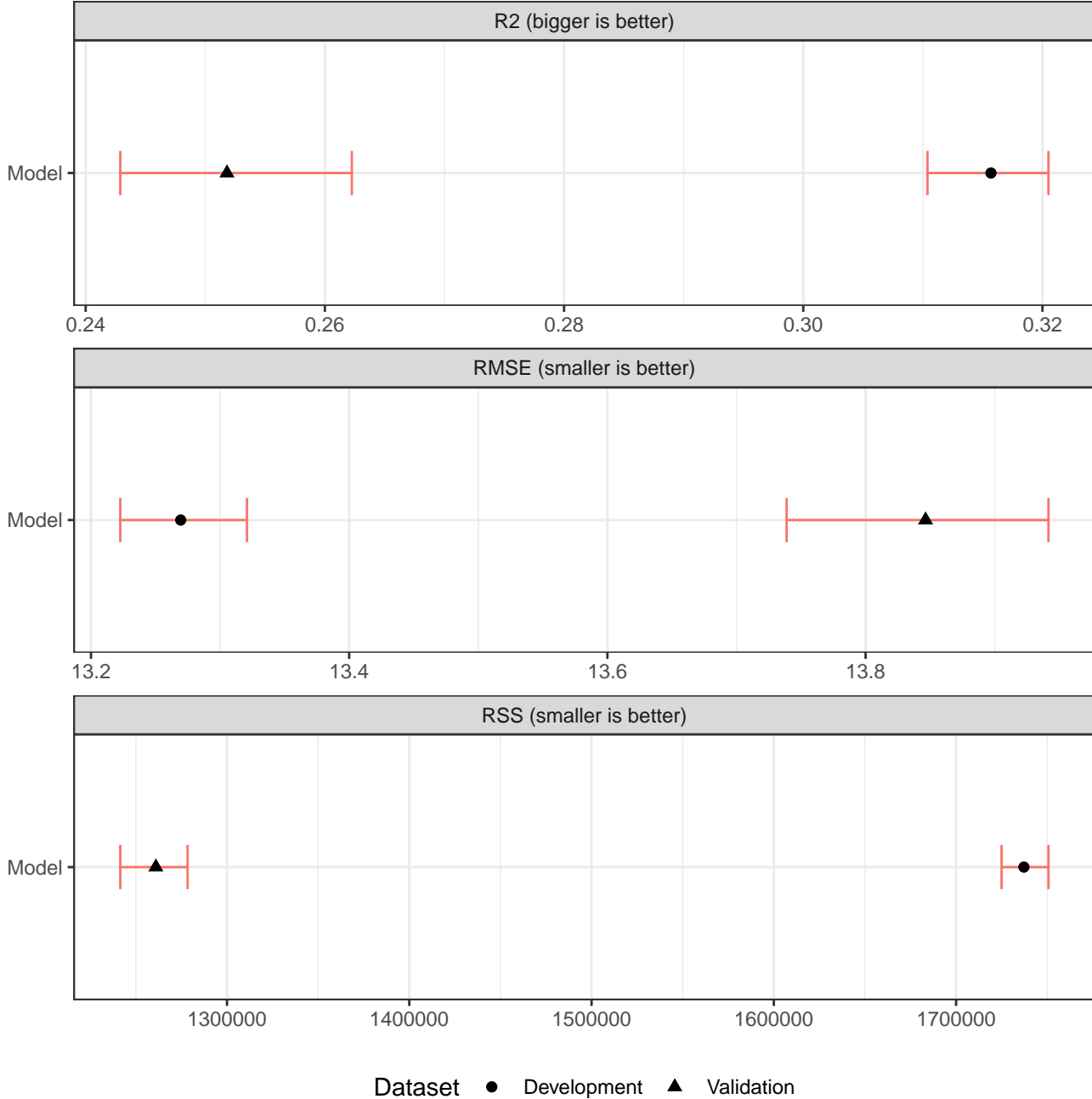


Histogram of $1 - \text{prop_model_final}\p_hat_train

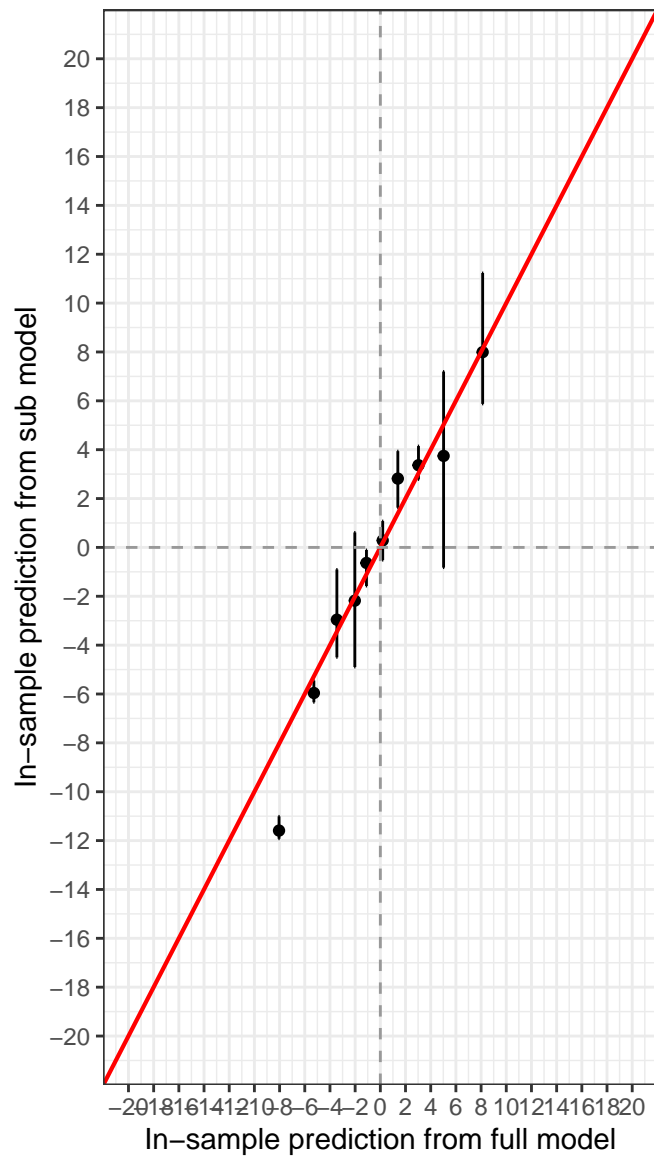
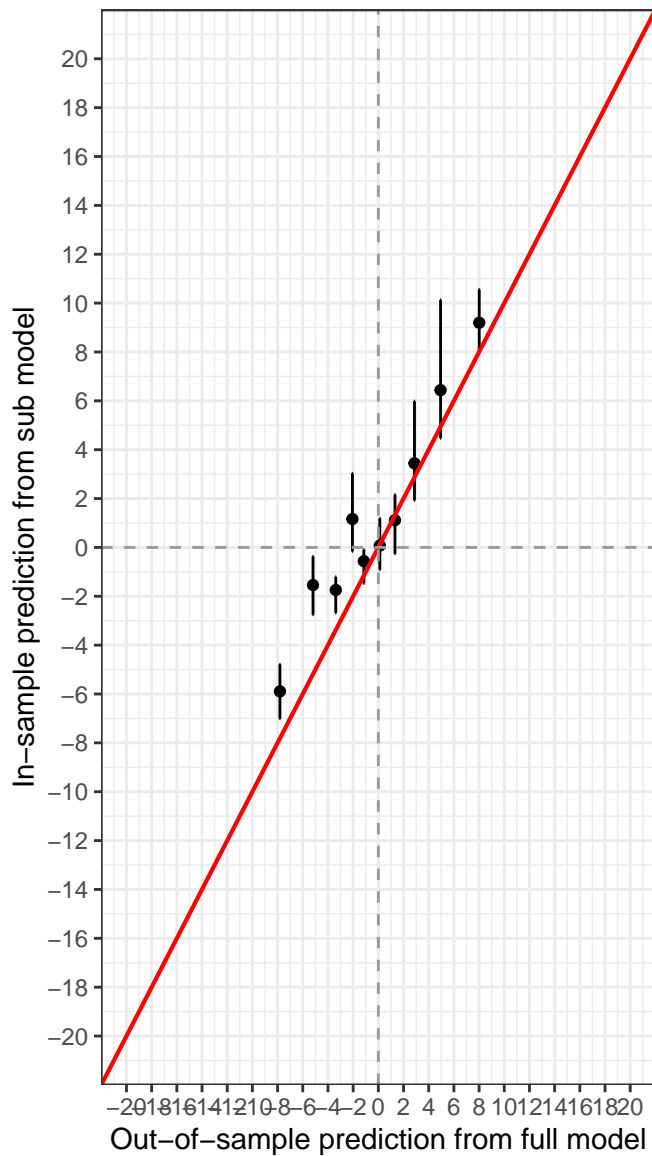


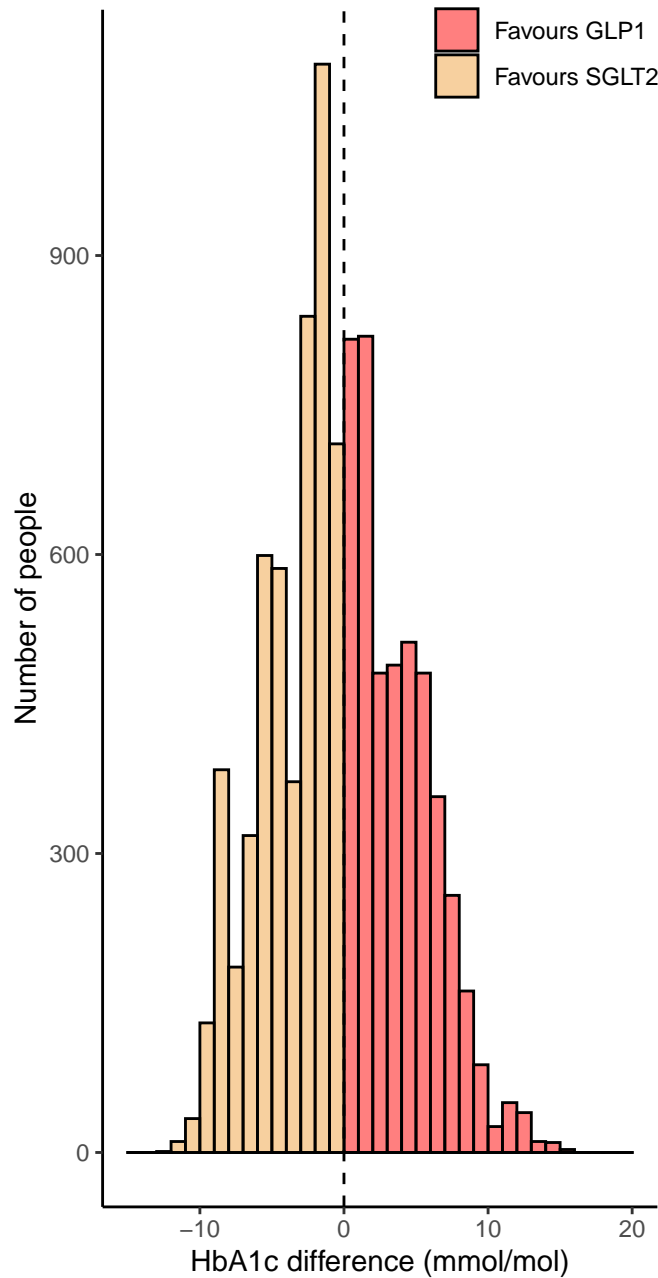
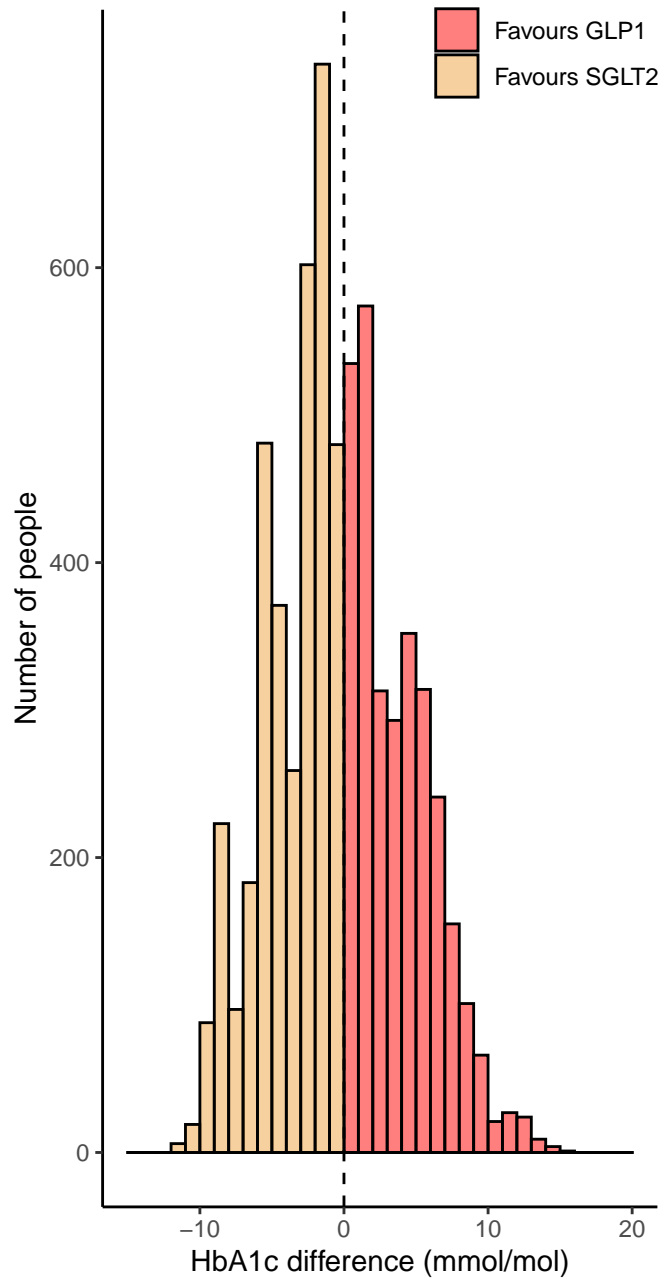
Residuals of Model 3

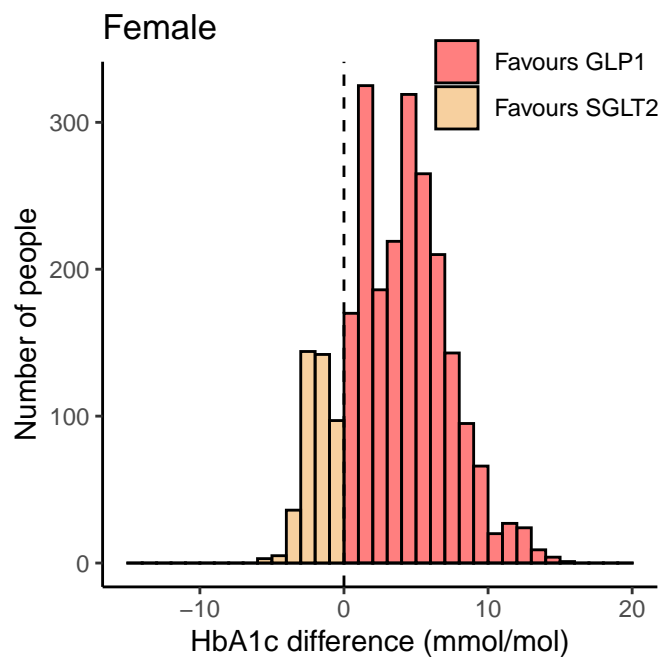
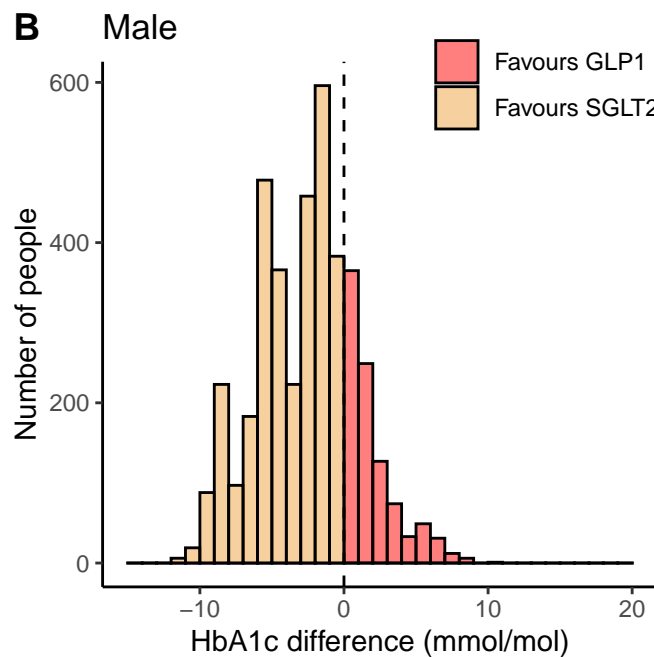
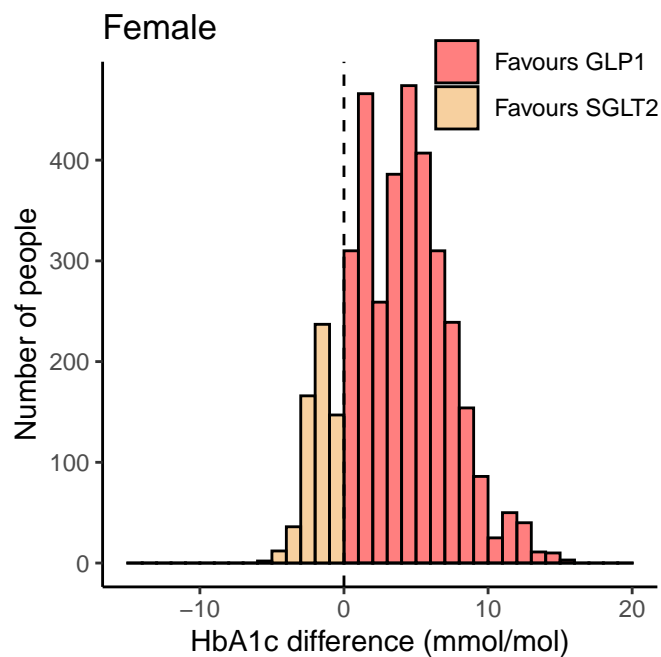
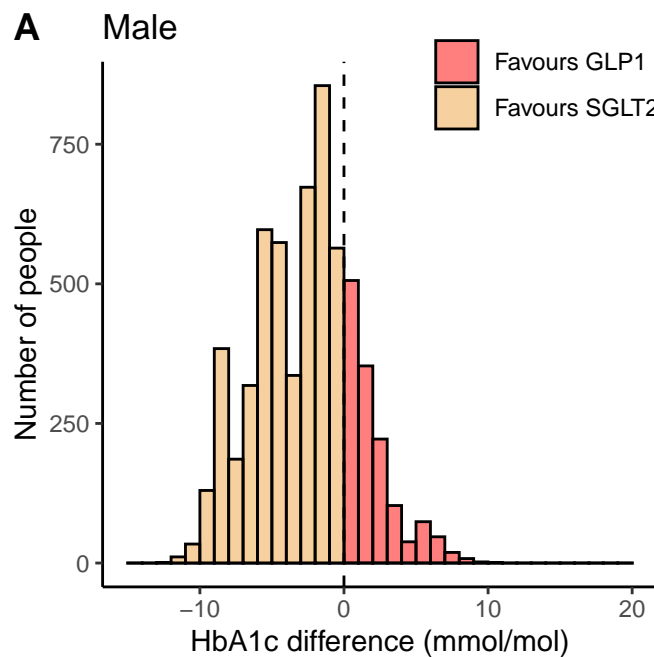




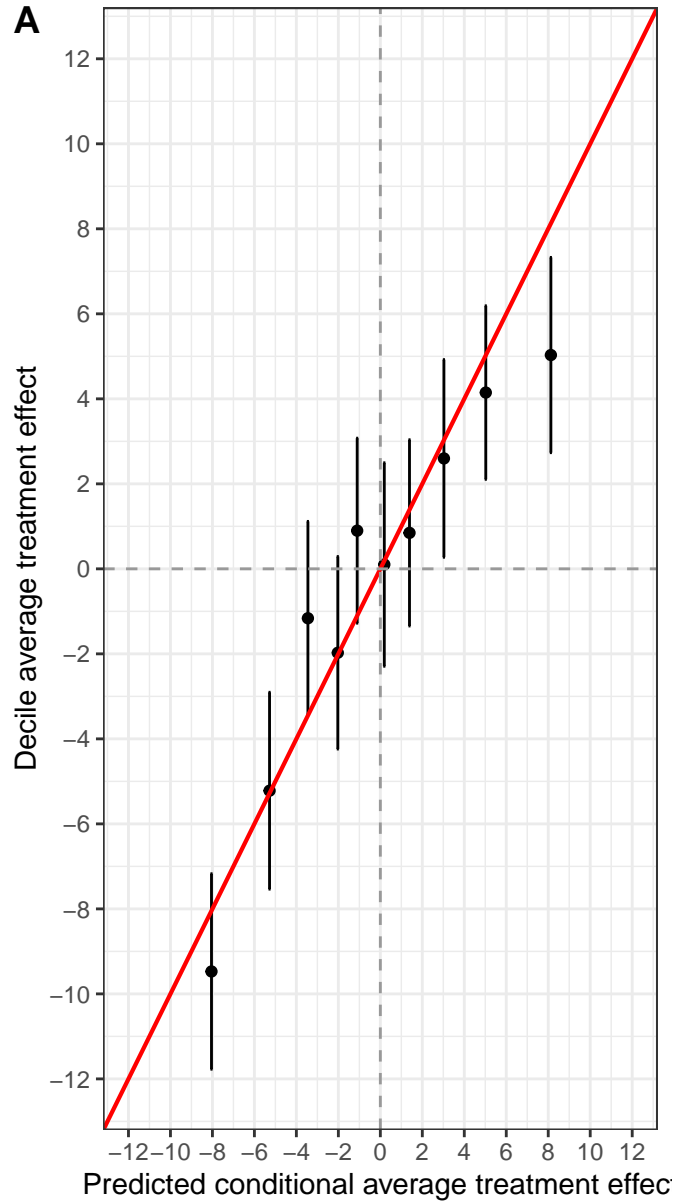
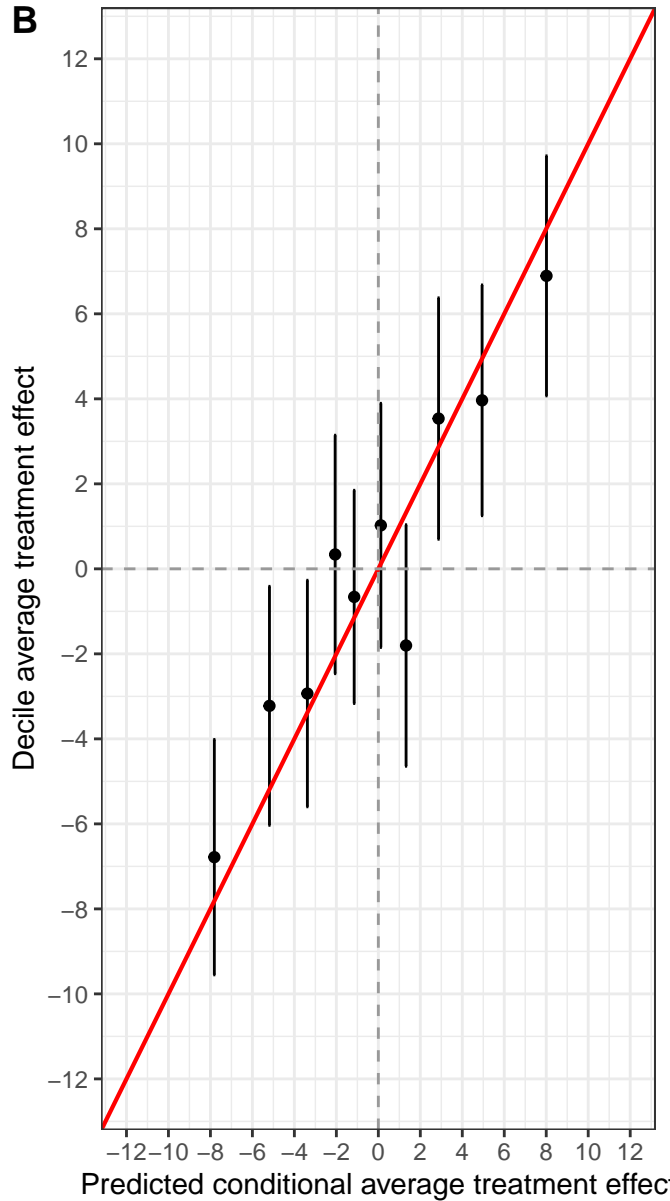
Effect submodels

A**B**

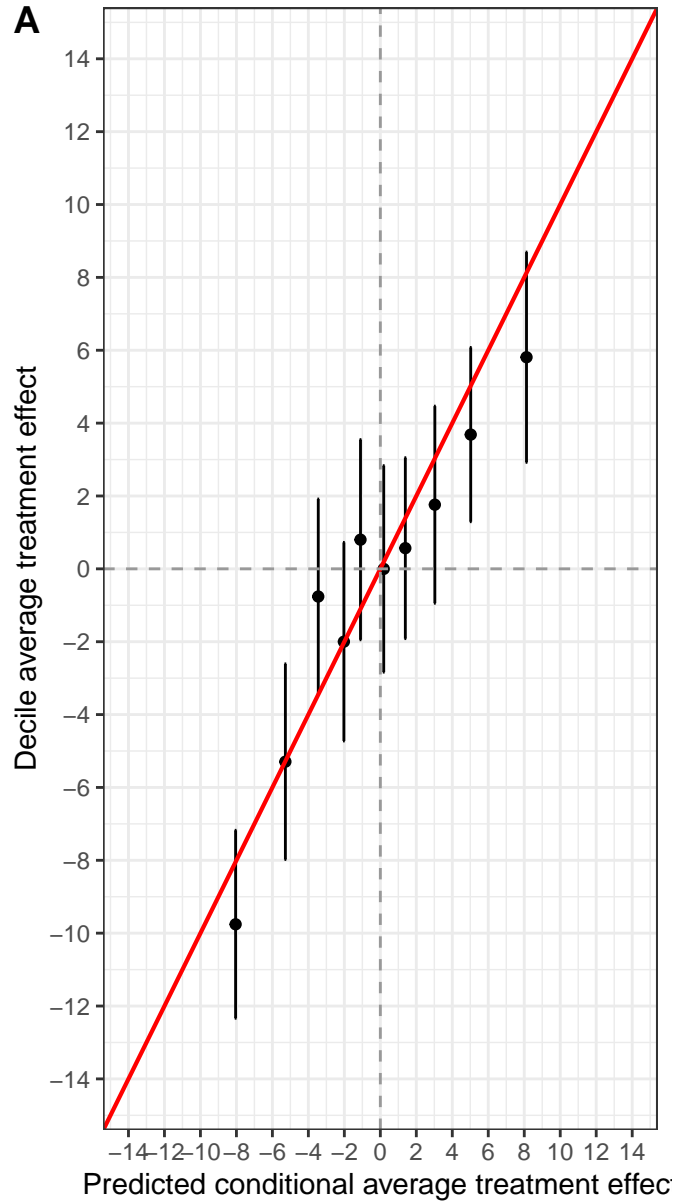
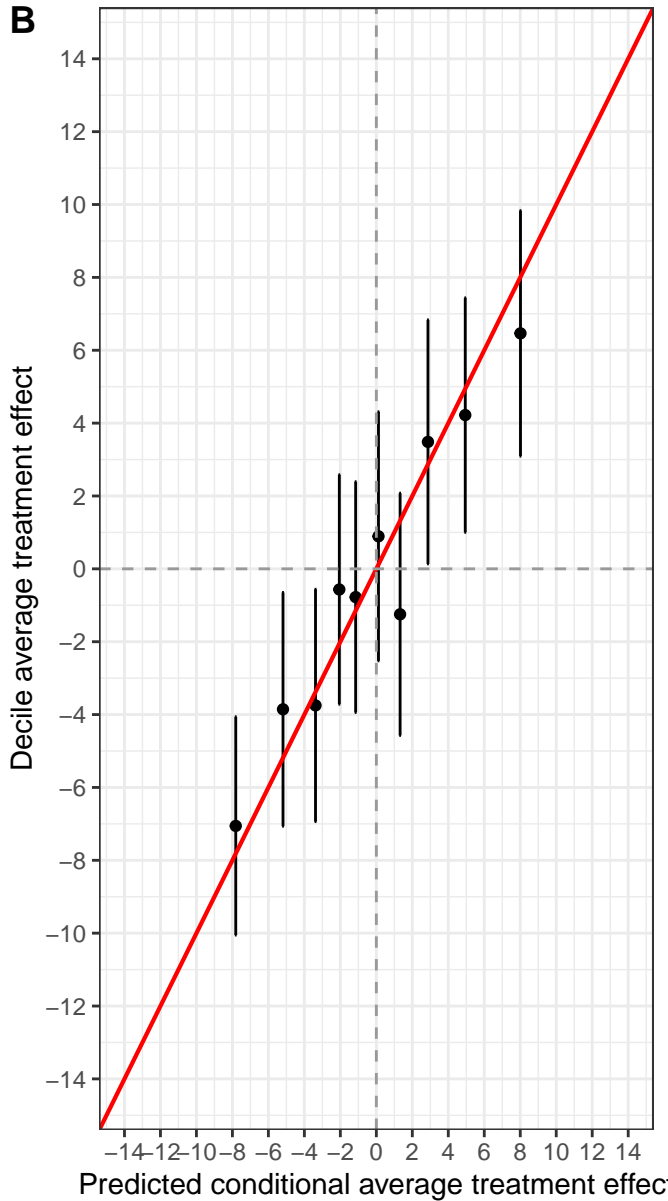
A**B**



Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

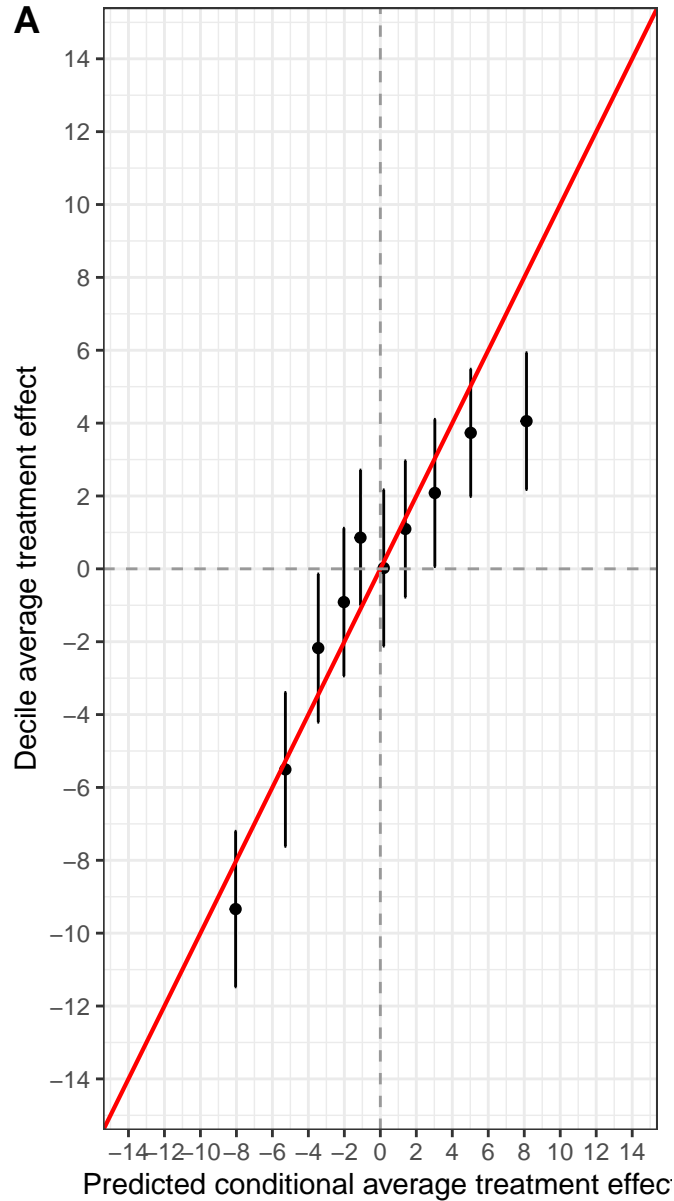
A**B**

Effects validation prop score matching

A**B**

Effects validation prop score inverse weighting

A



B

