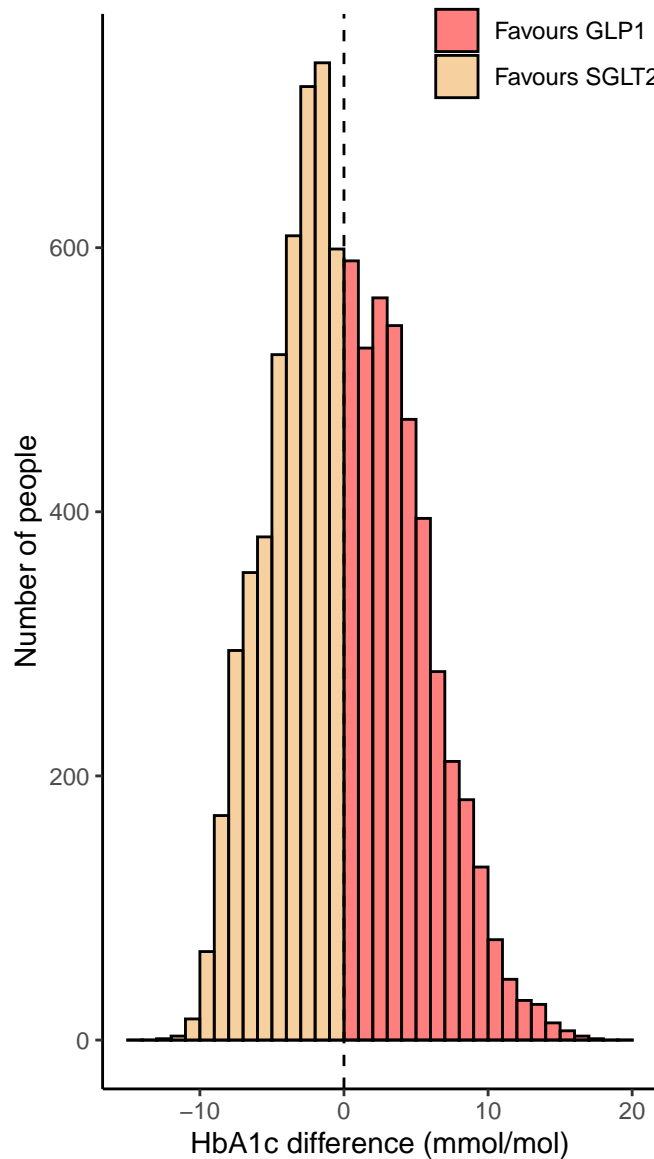
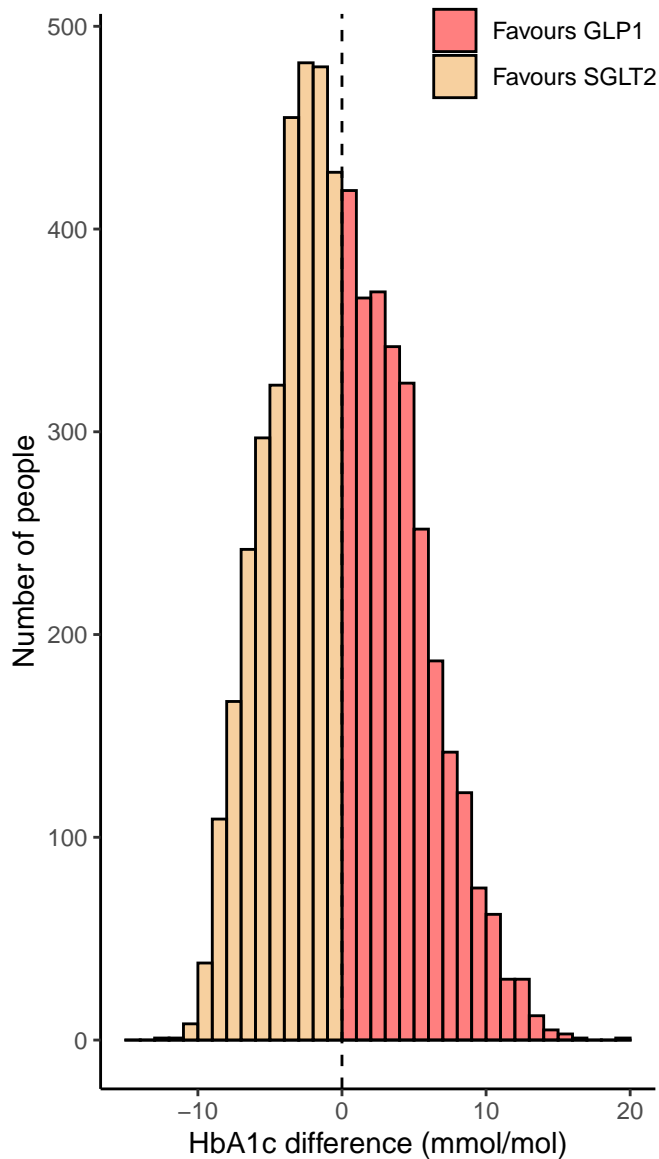
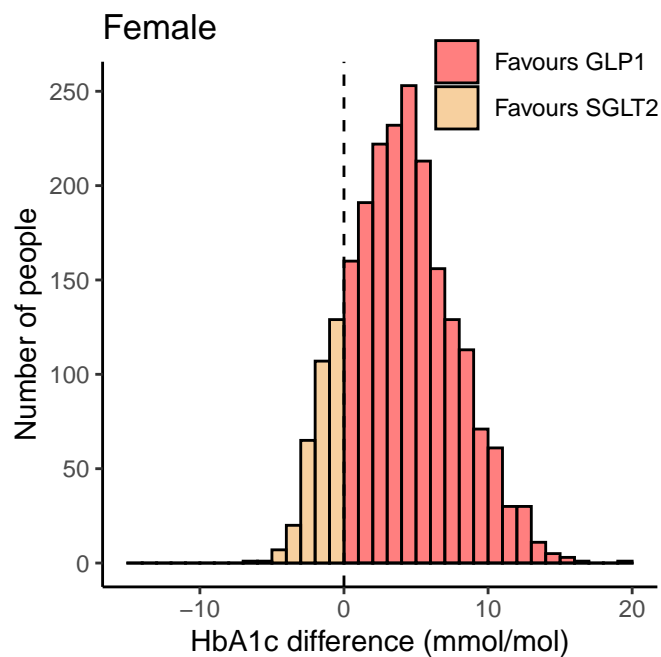
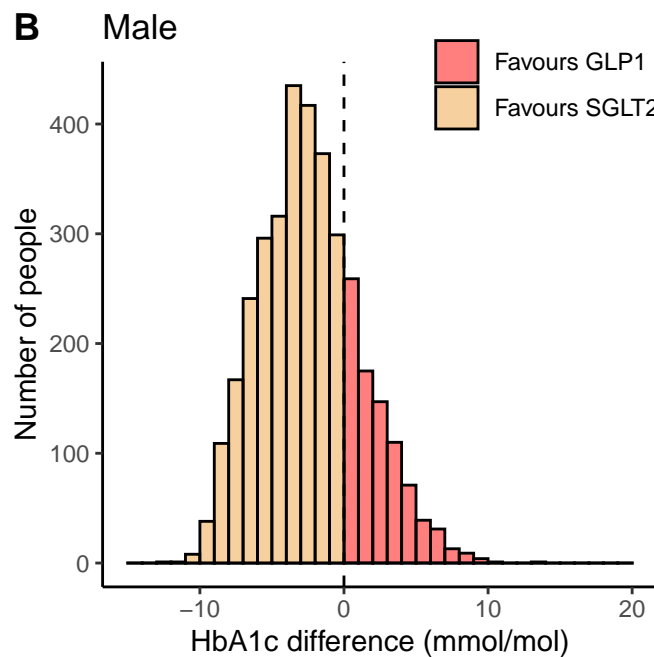
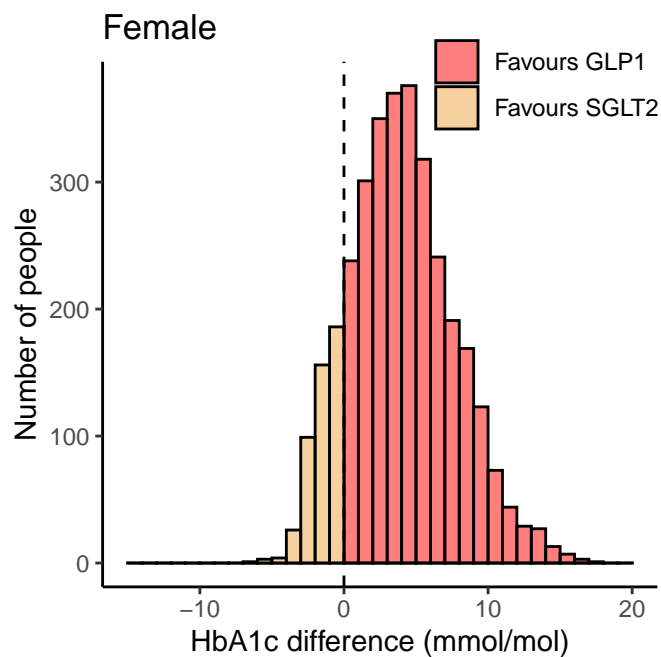
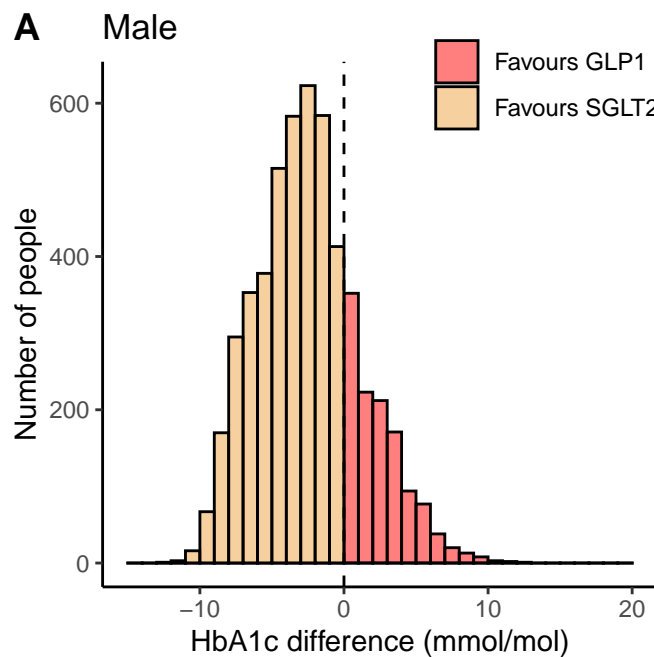
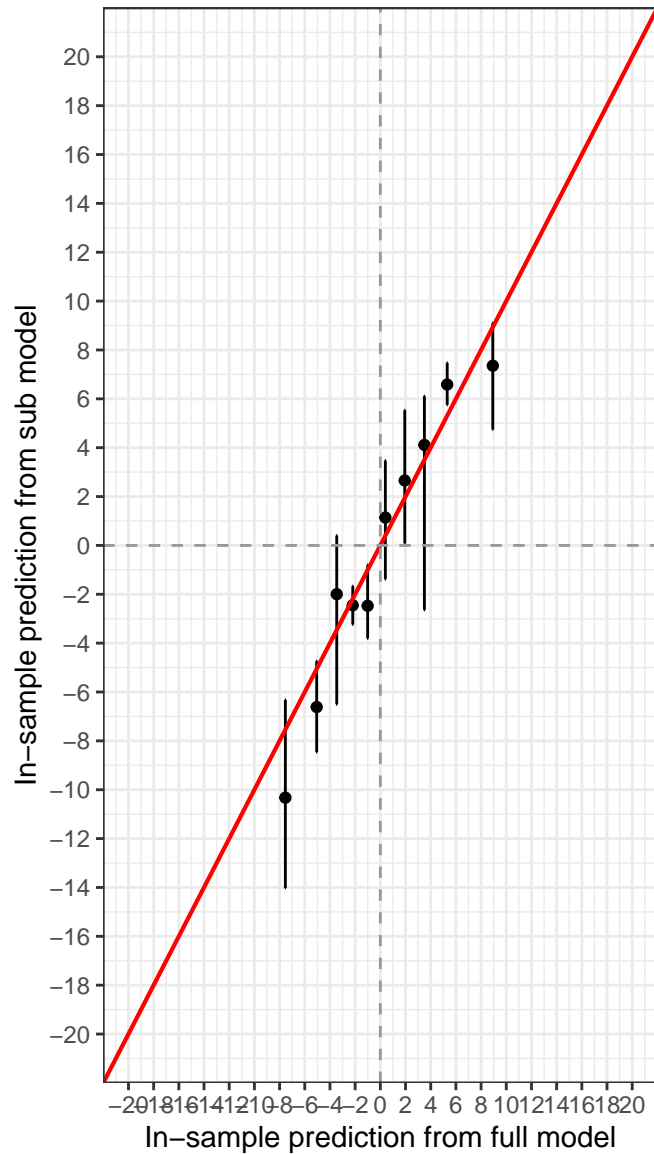
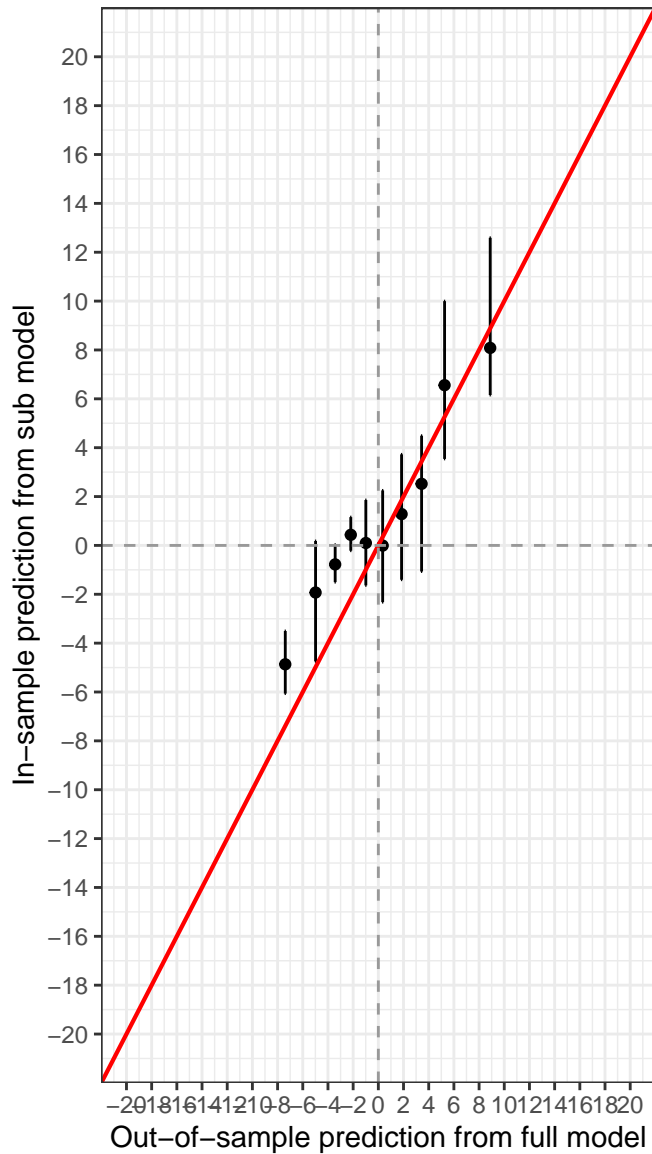


Model fitting: Complete data (Routine/No propensity score)

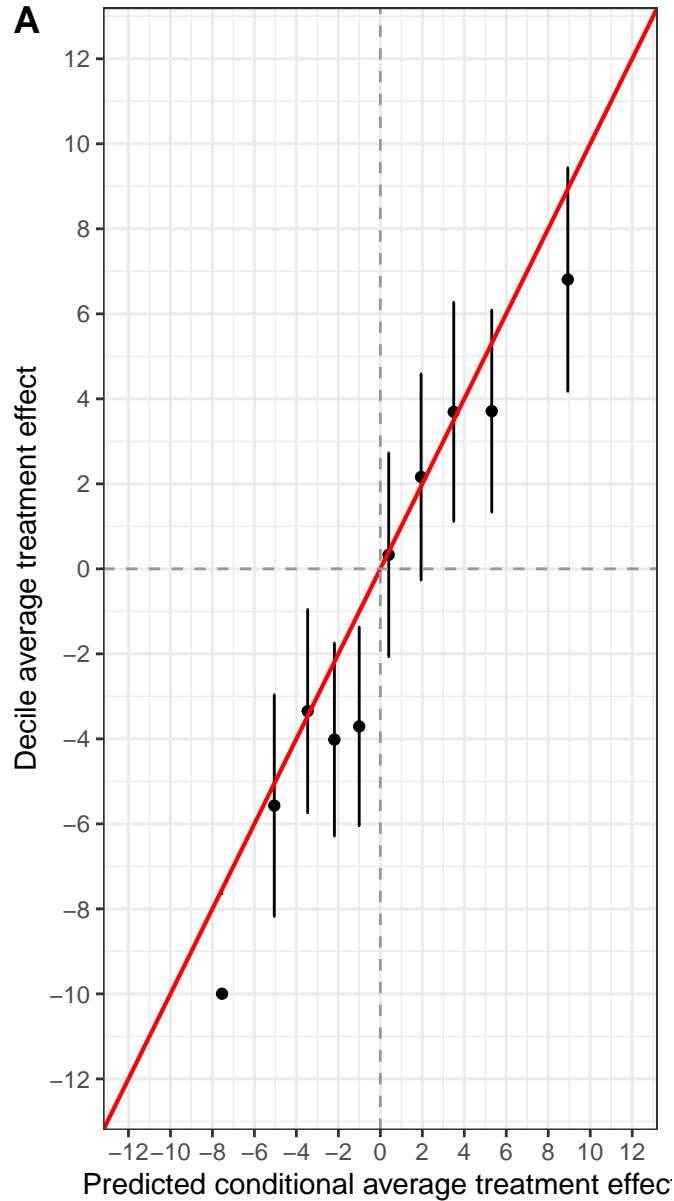
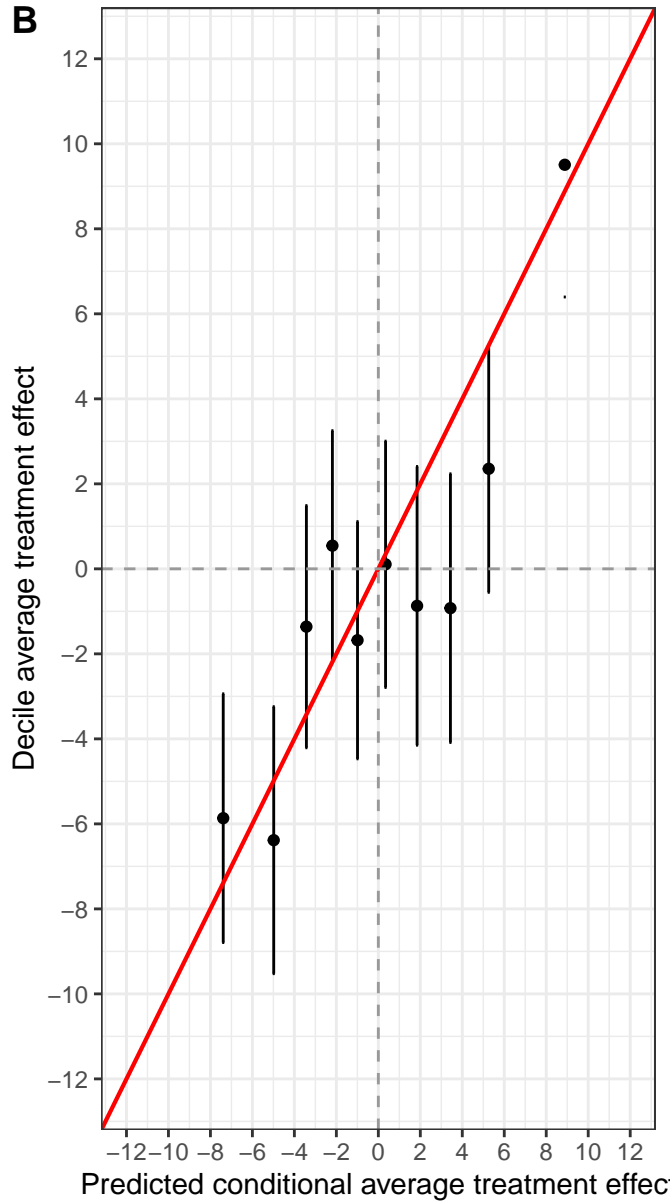
A**B**



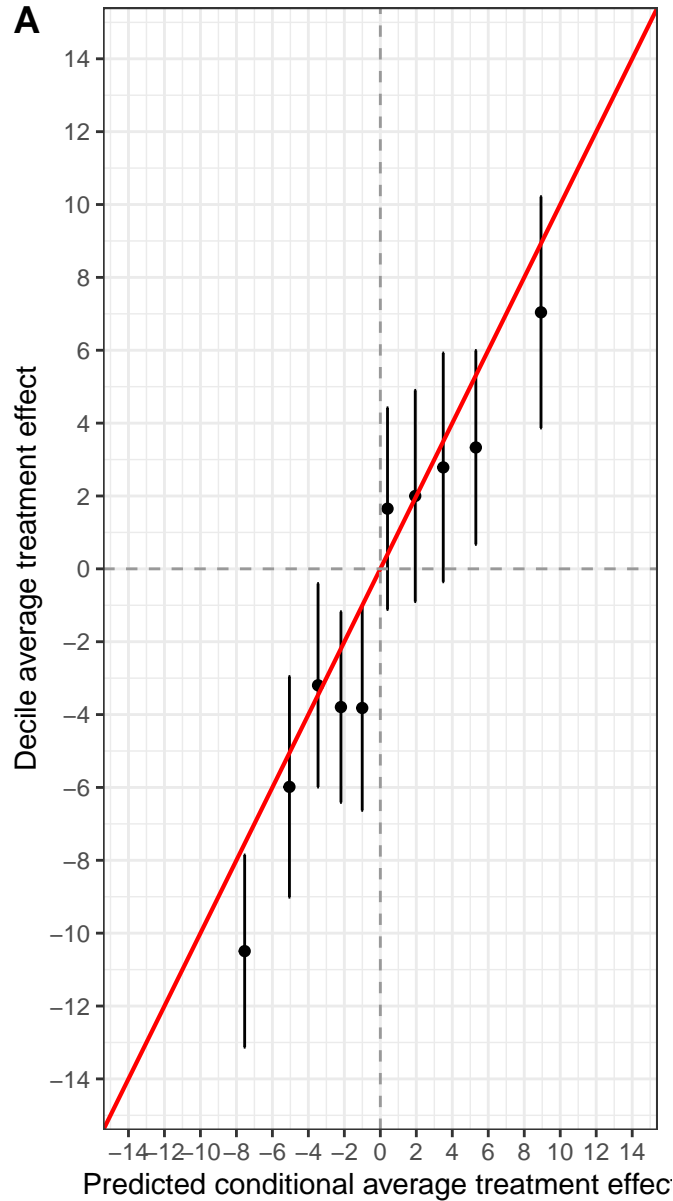
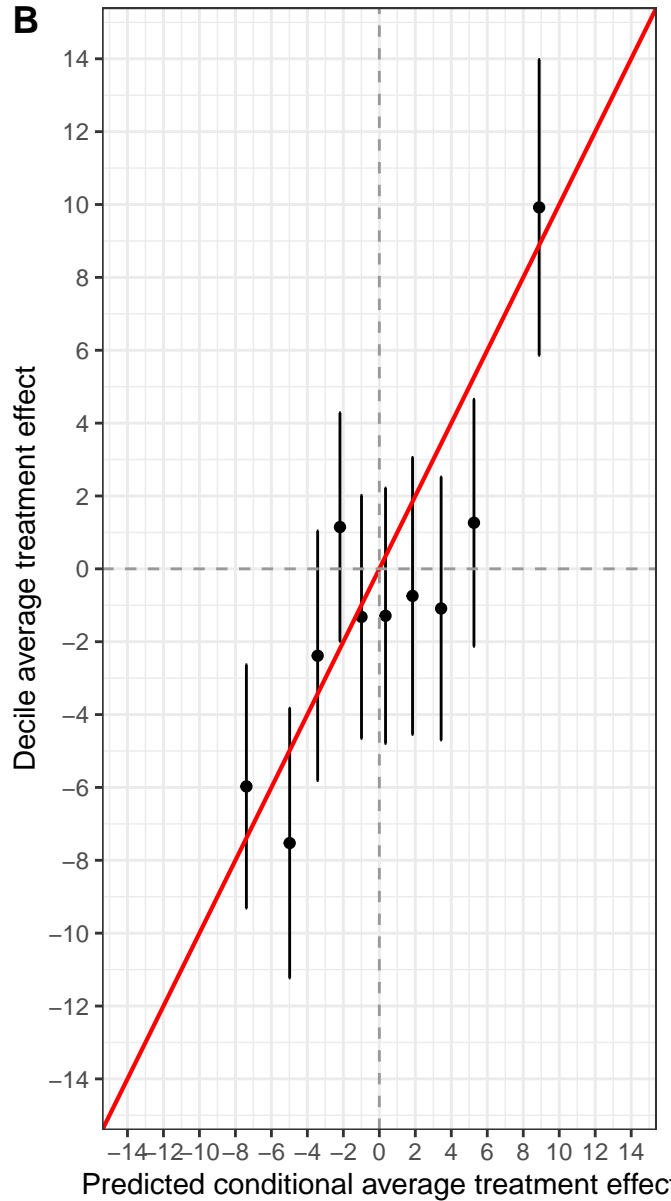
Effect submodels

A**B**

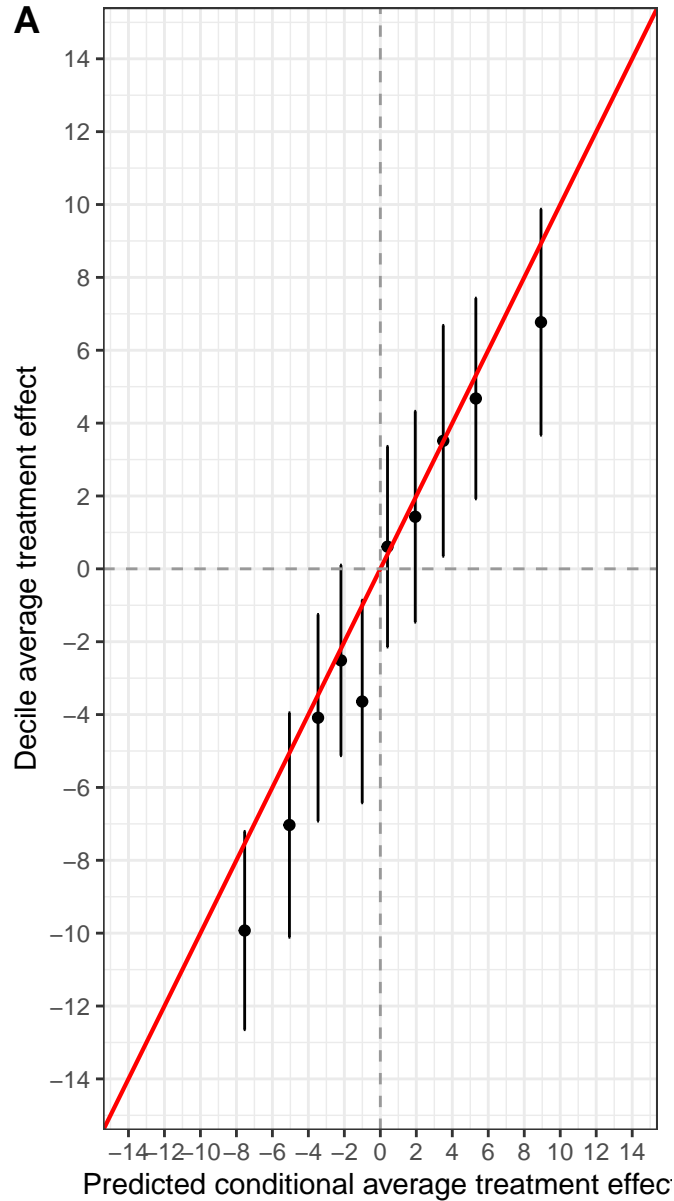
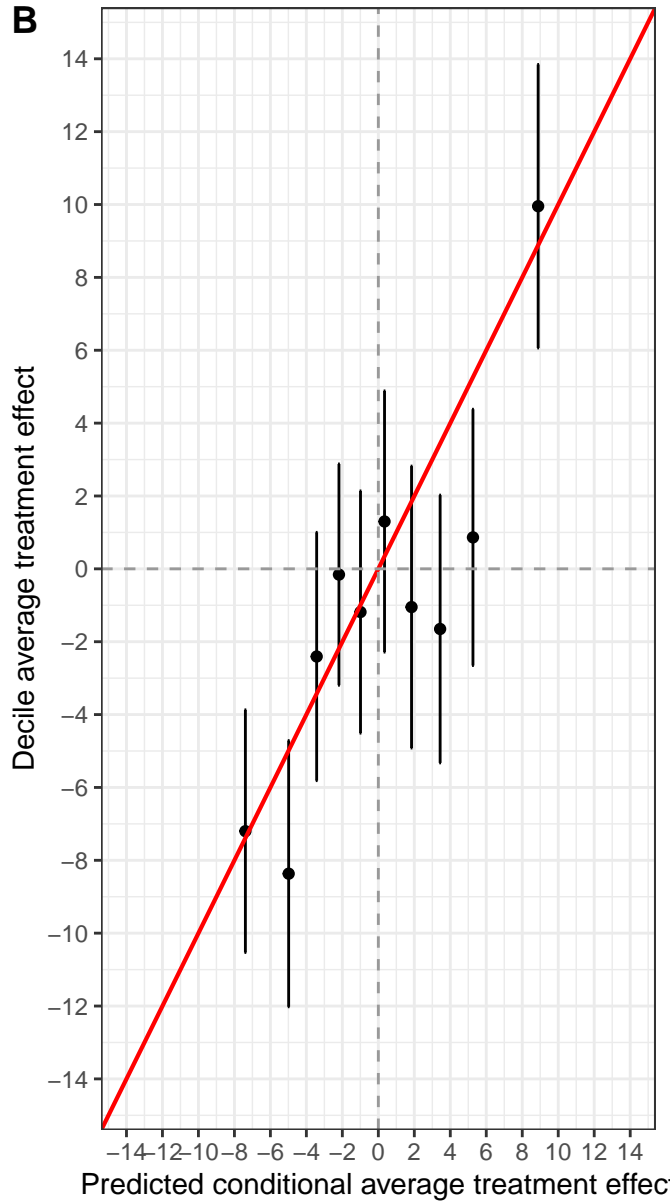
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

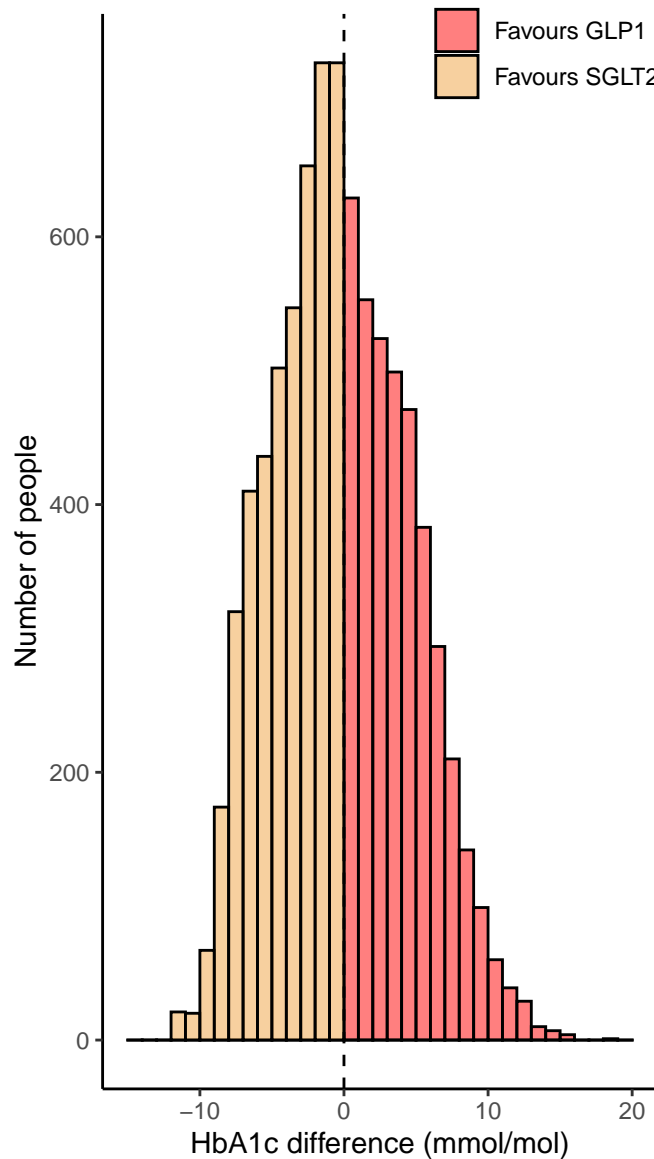
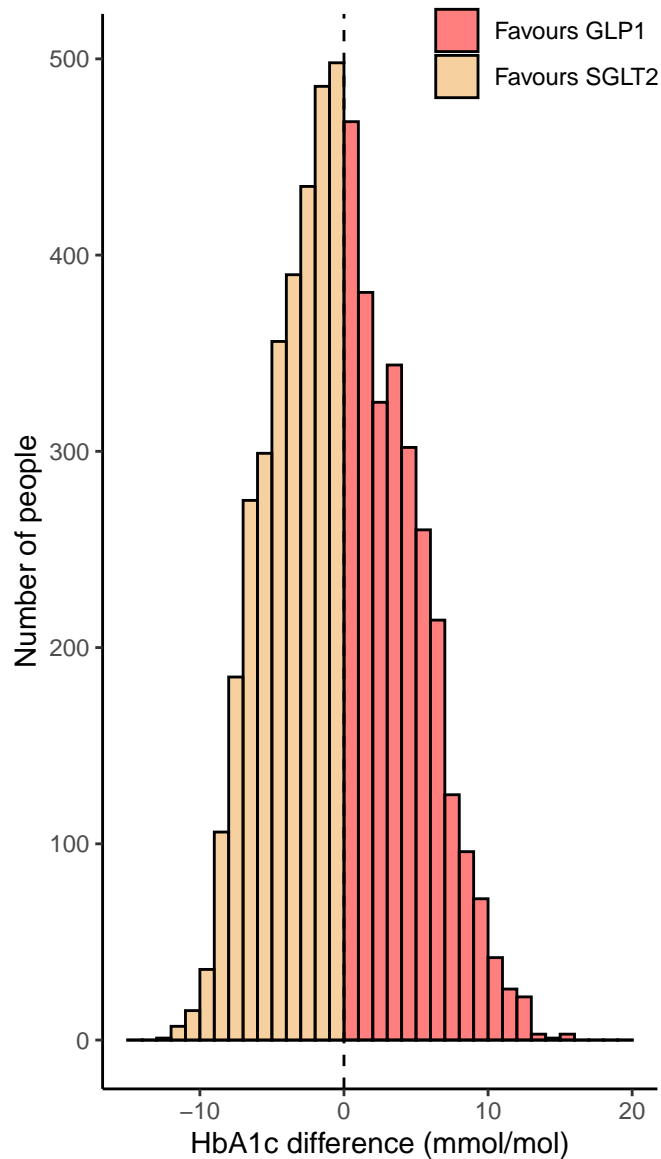
Effects validation prop score matching

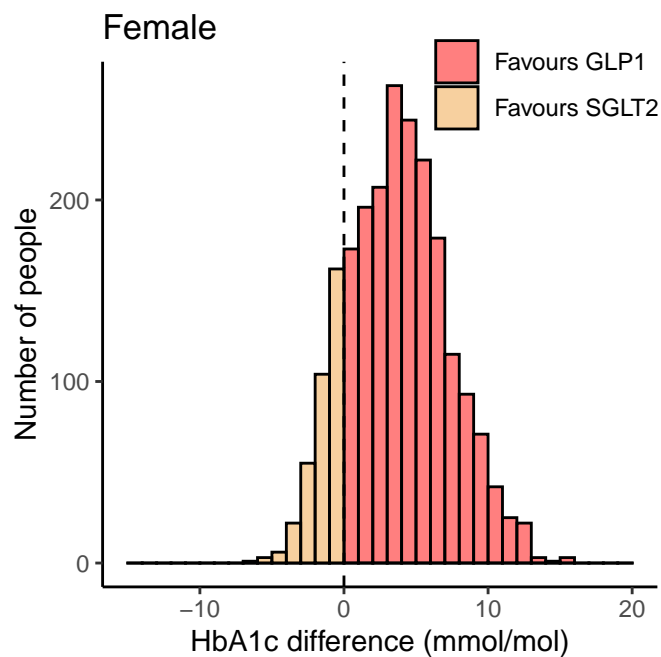
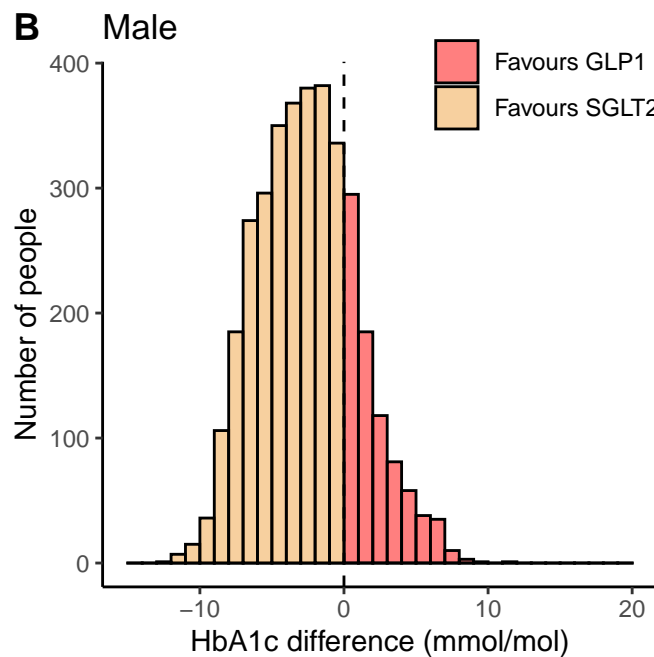
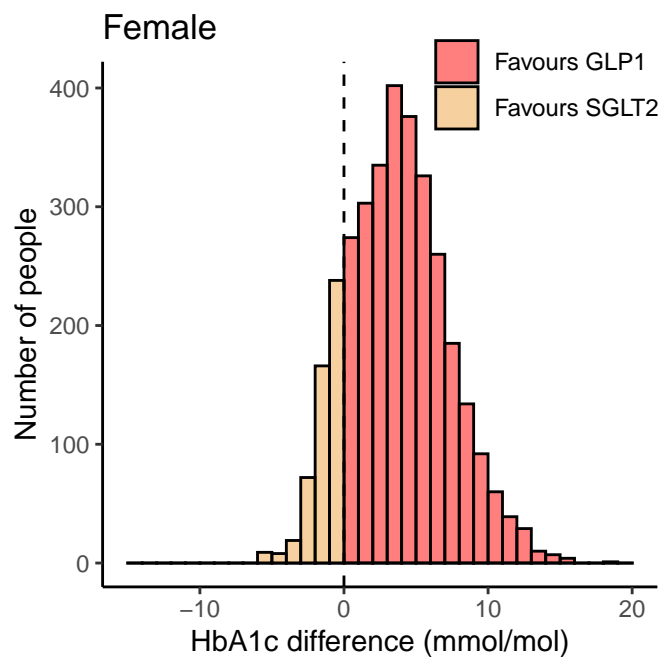
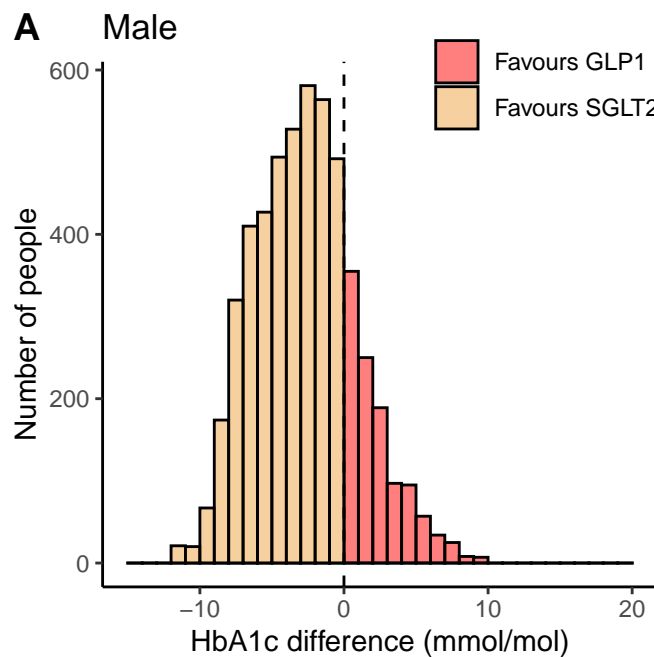
A**B**

Effects validation prop score inverse weighting

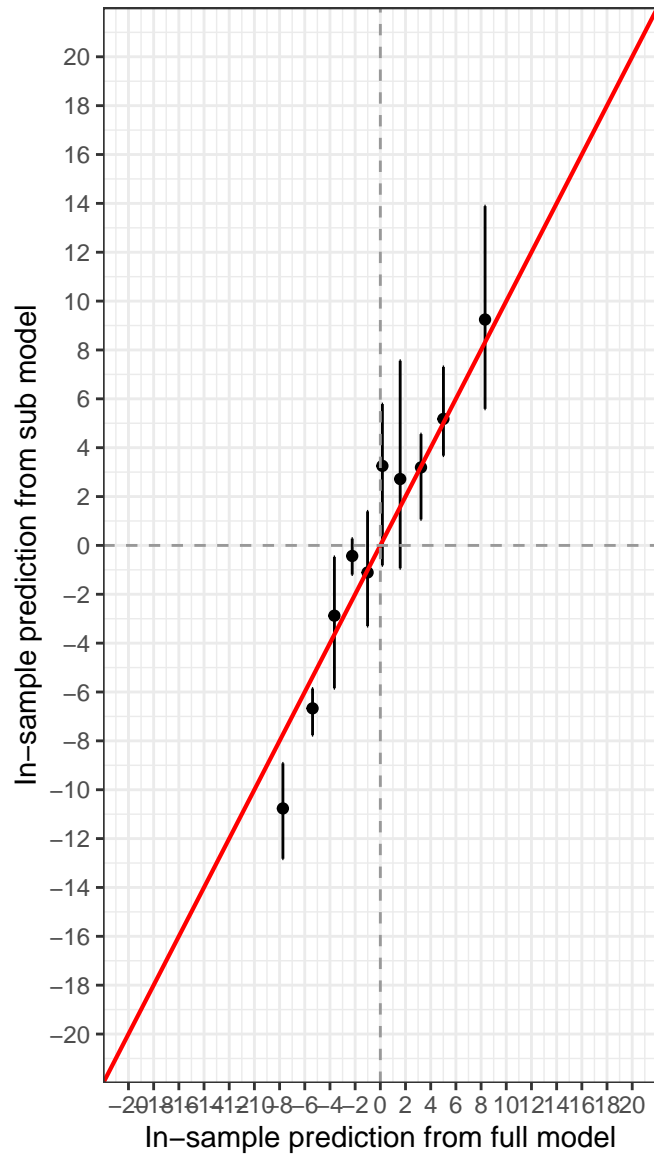
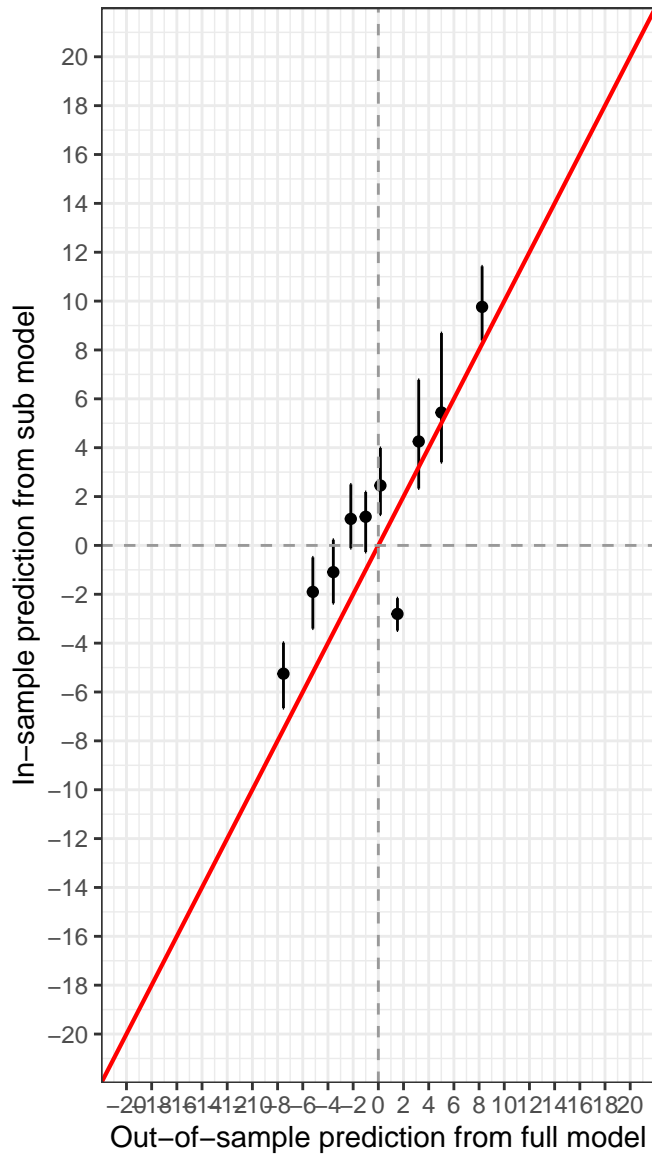
A**B**

Model fitting: Complete data (Routine/Propensity score)

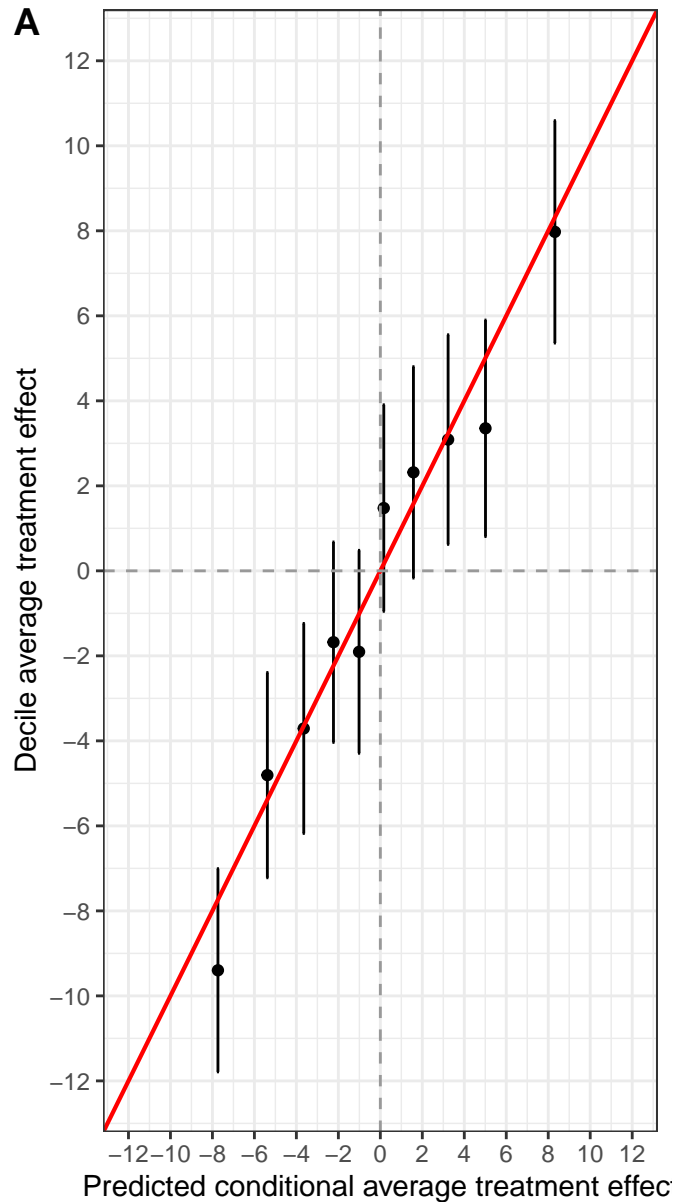
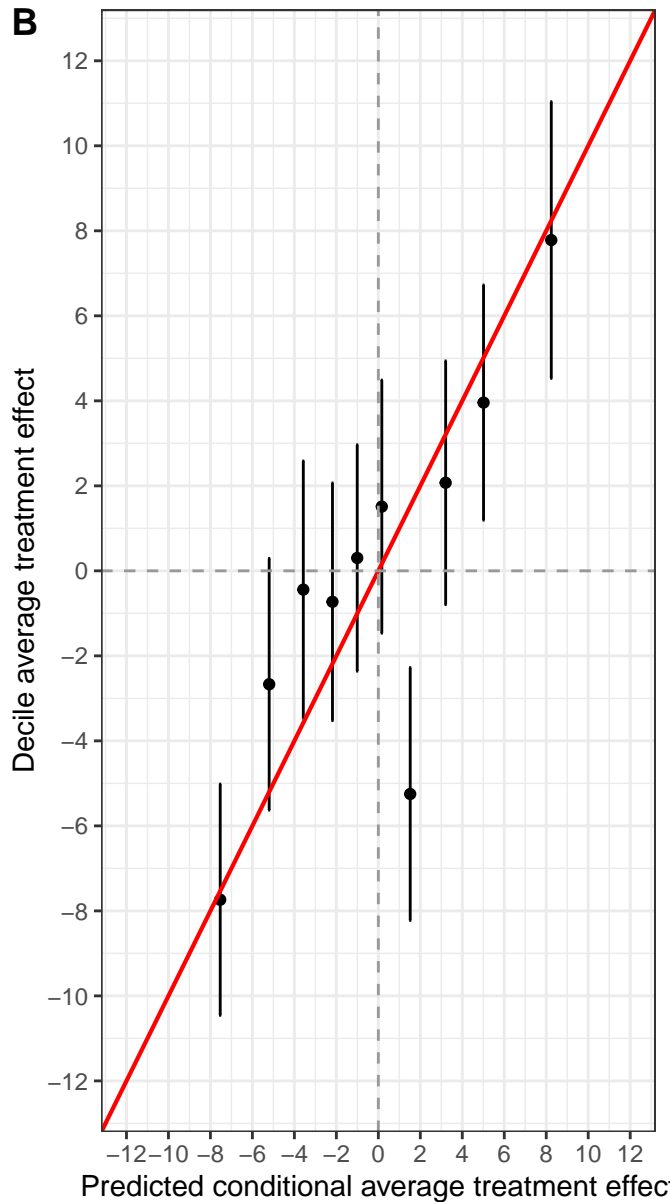
A**B**



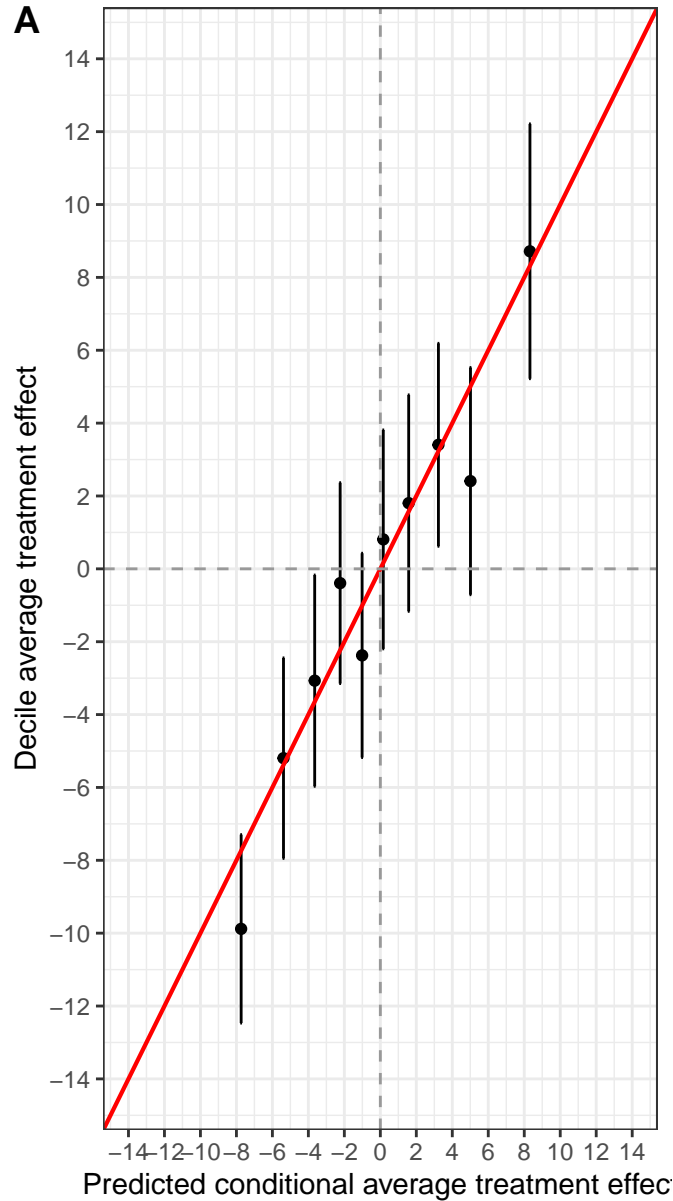
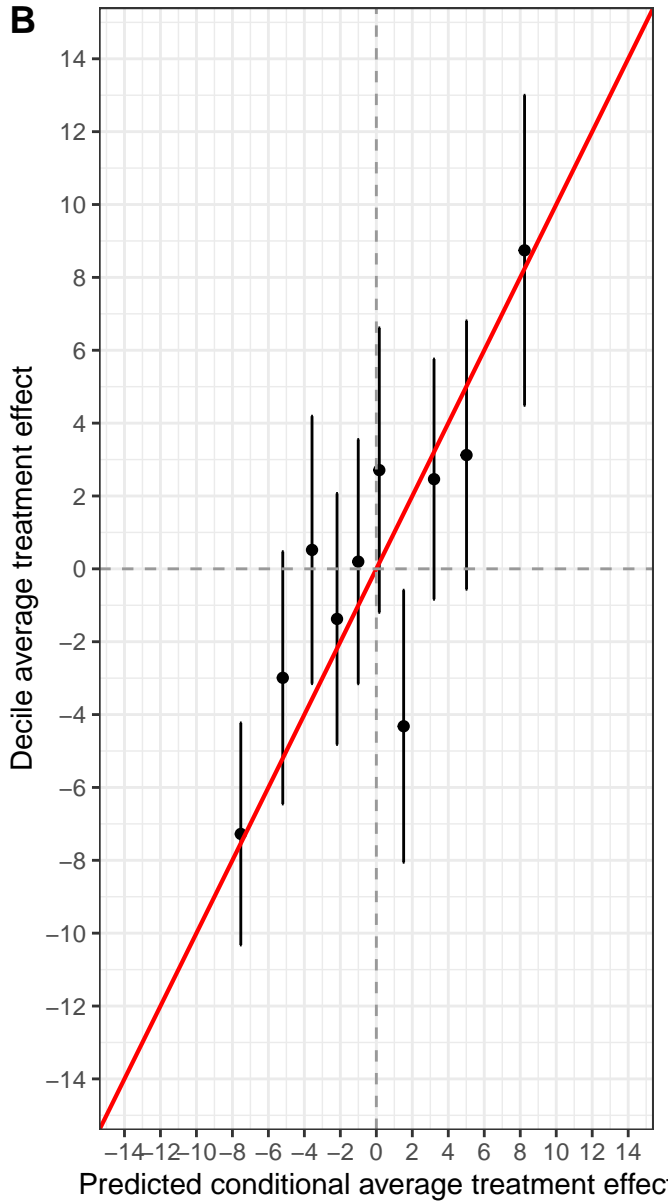
Effect submodels

A**B**

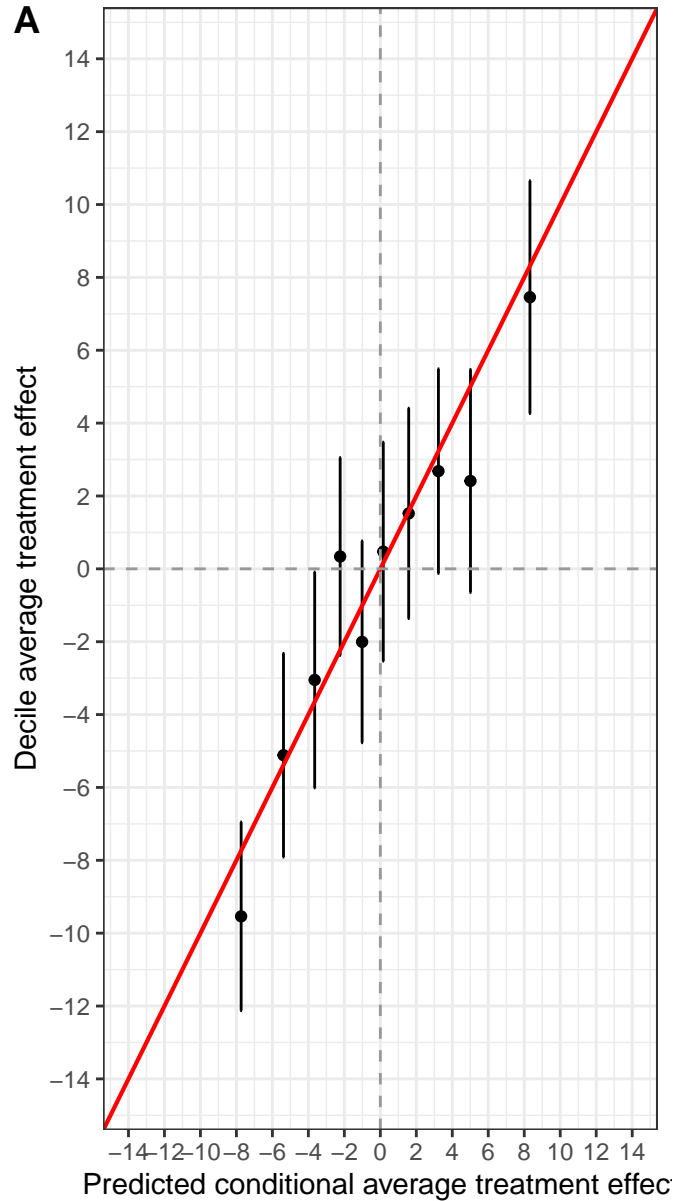
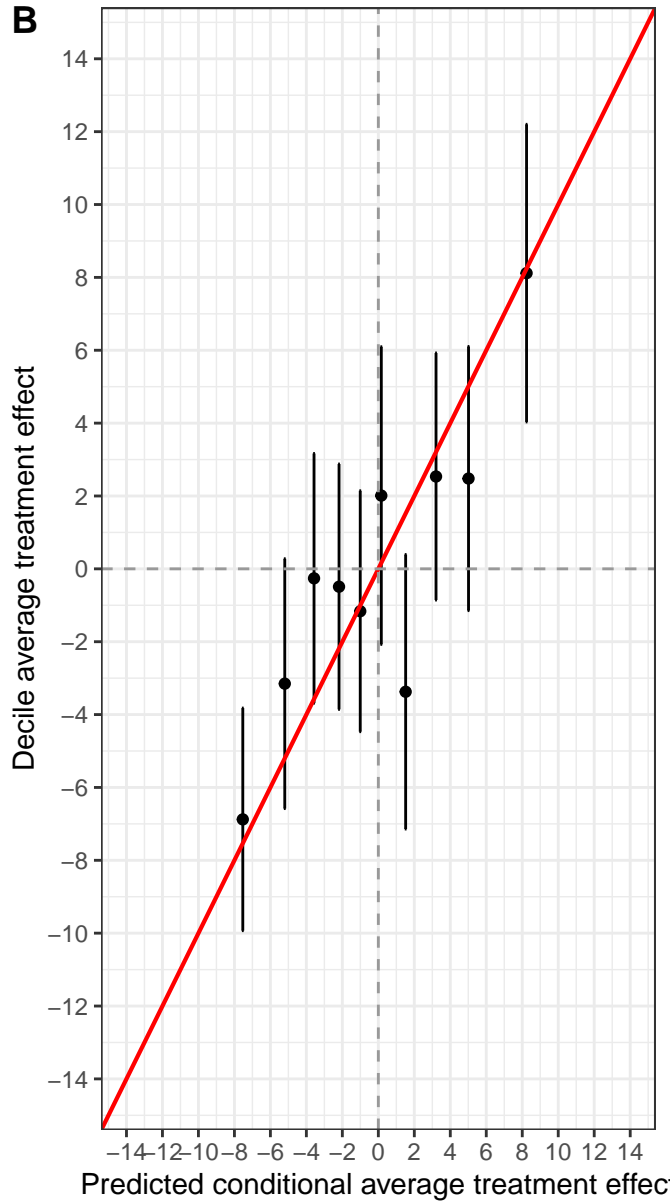
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

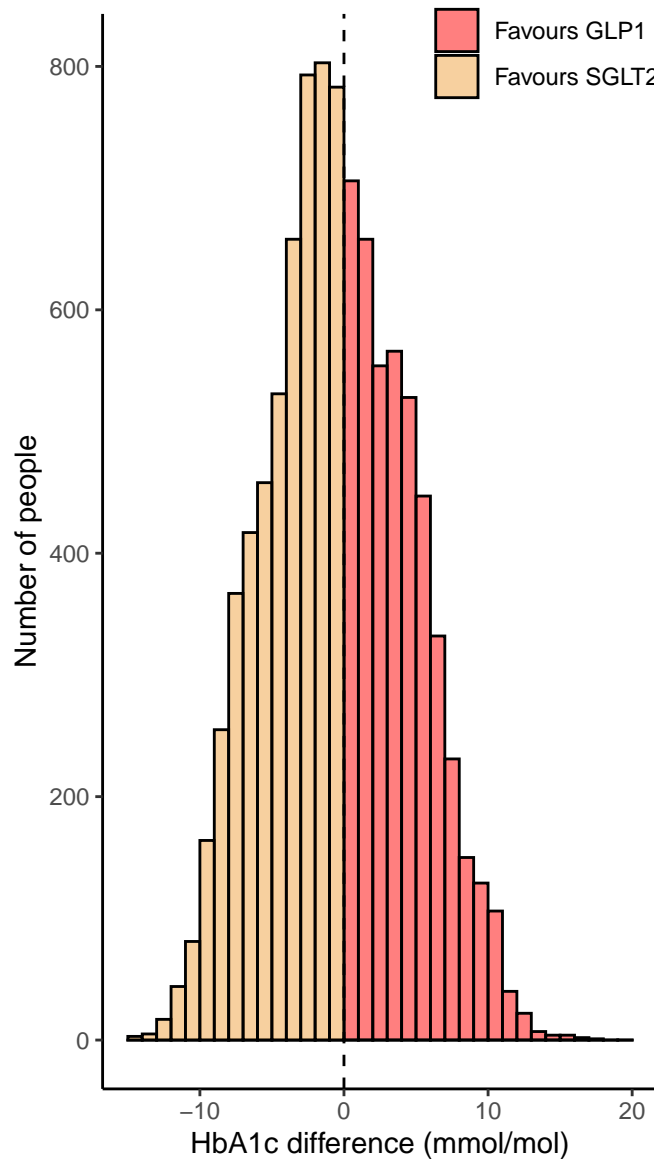
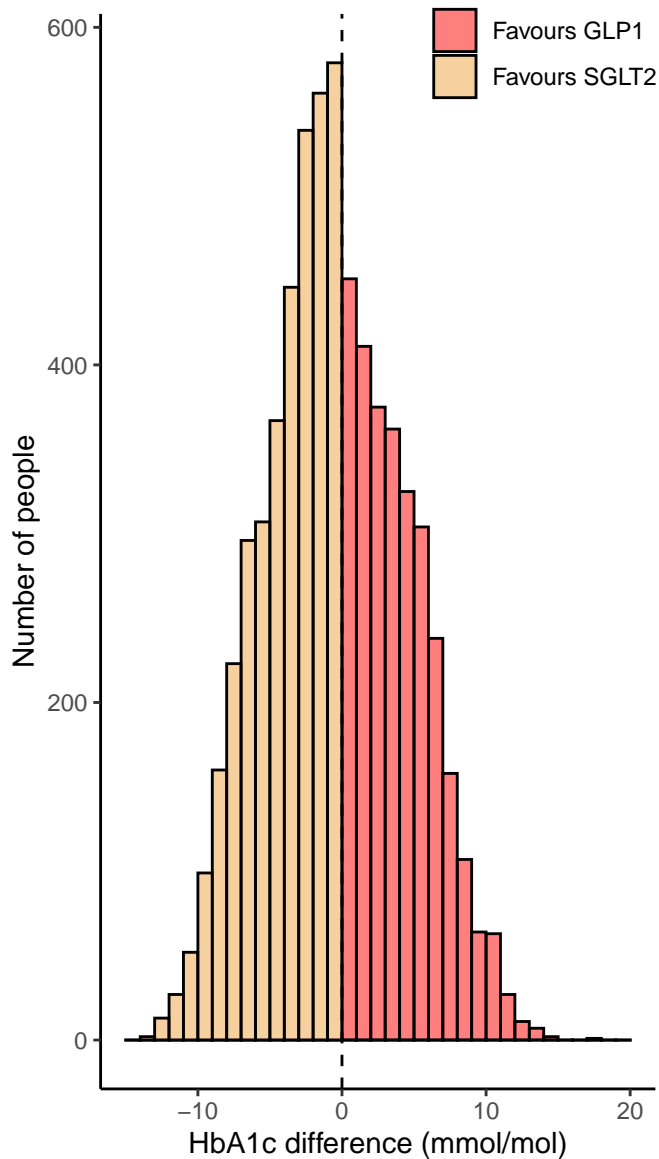
Effects validation prop score matching

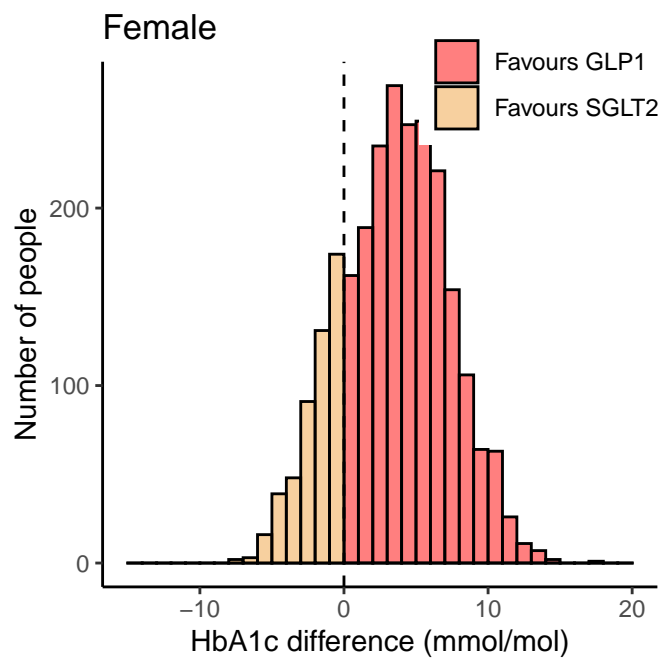
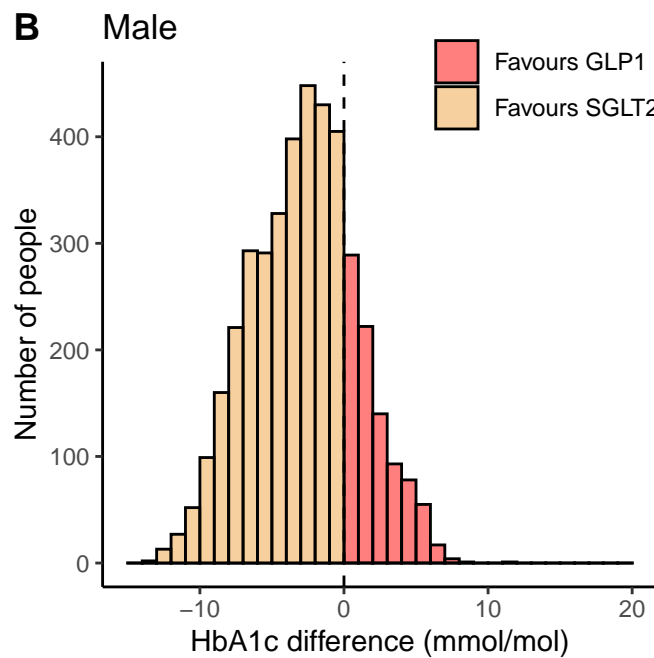
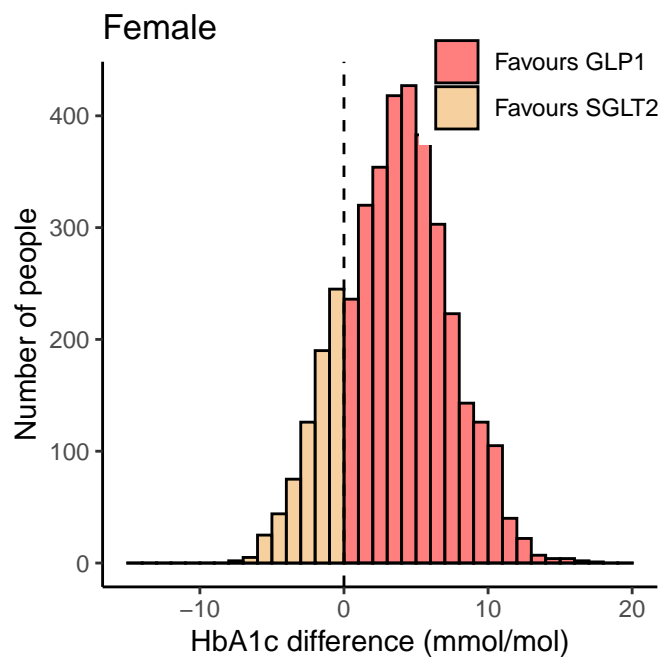
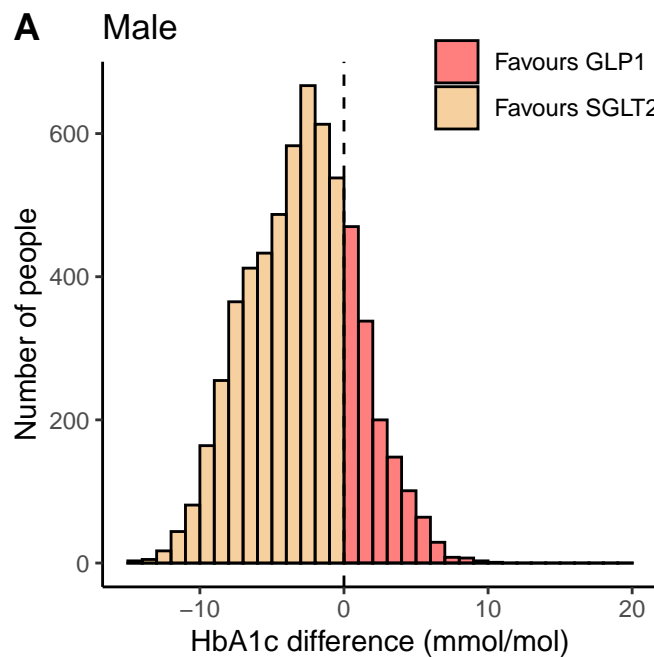
A**B**

Effects validation prop score inverse weighting

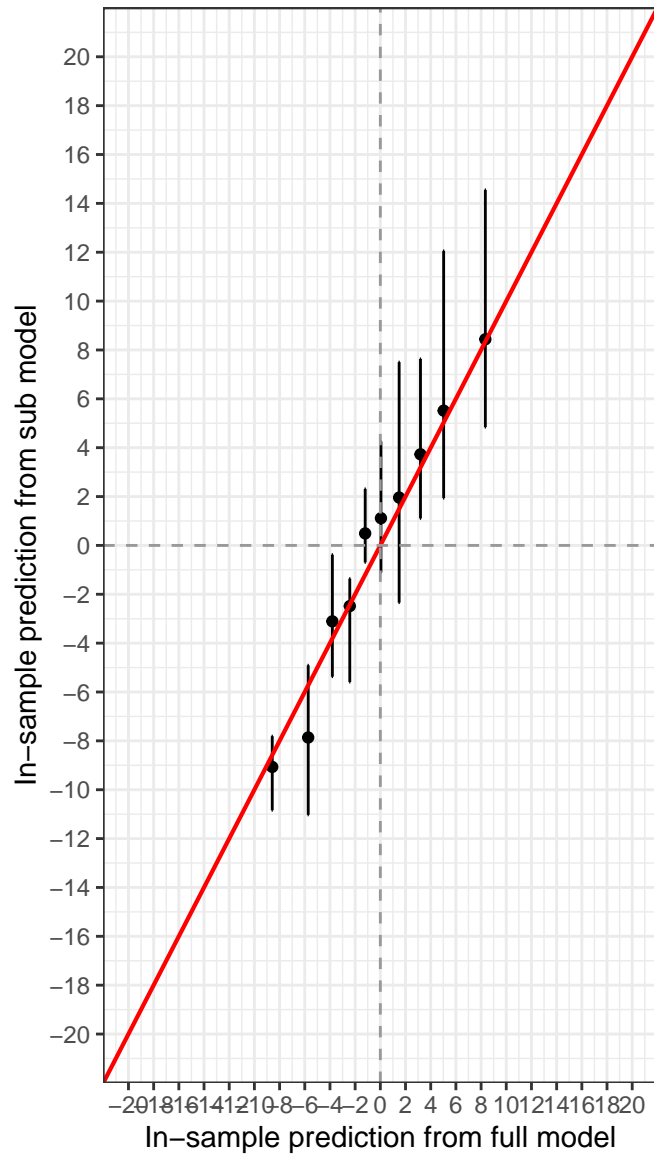
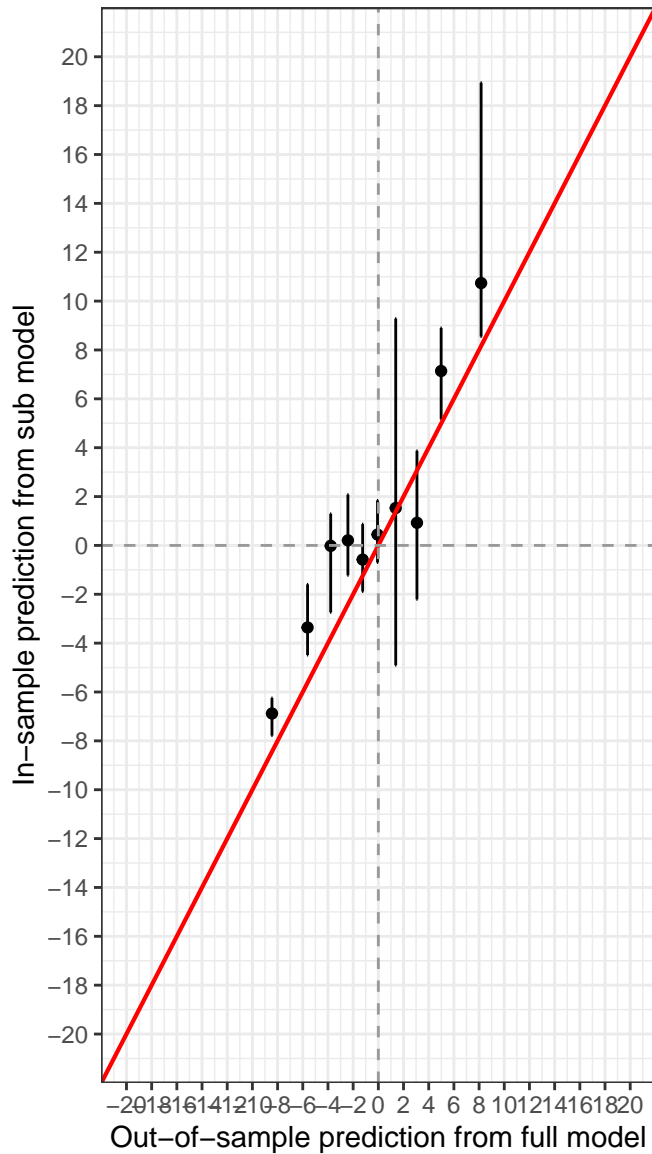
A**B**

Model fitting: Incomplete data (Routine/No propensity score)

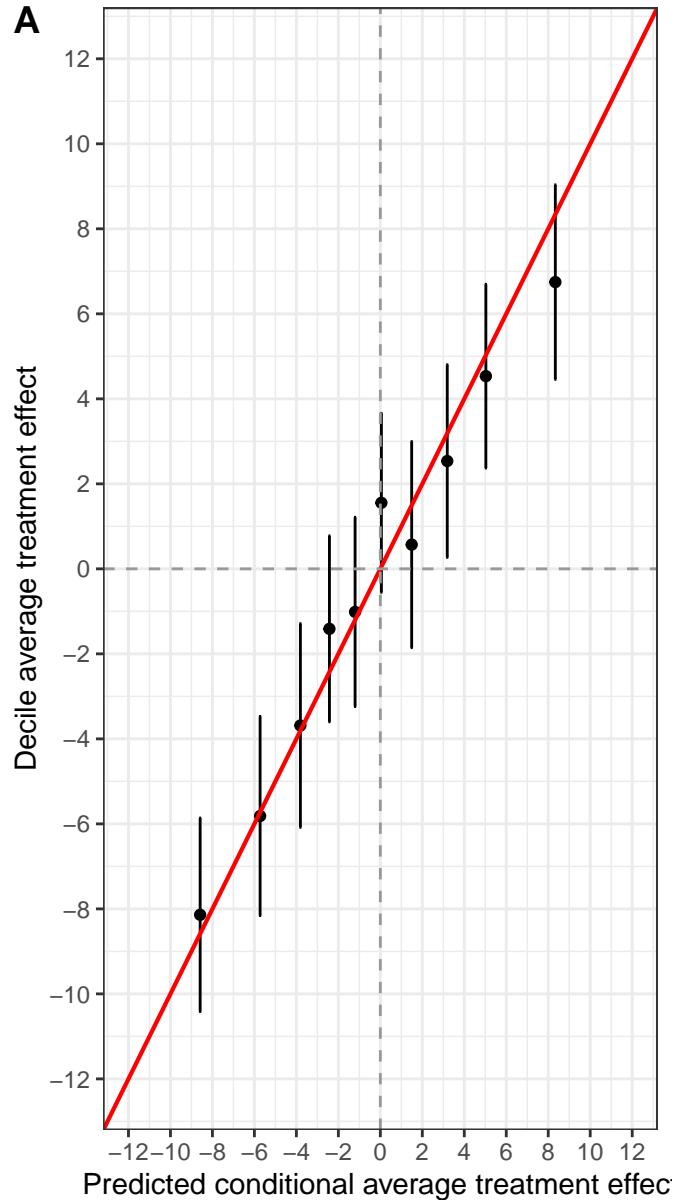
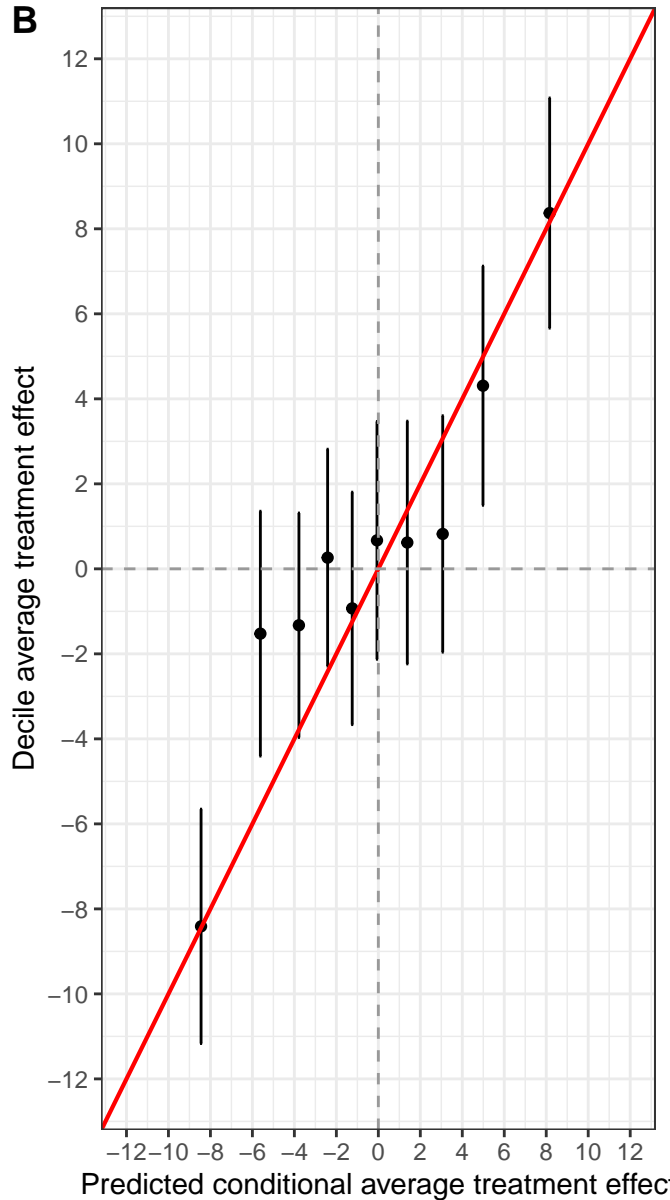
A**B**



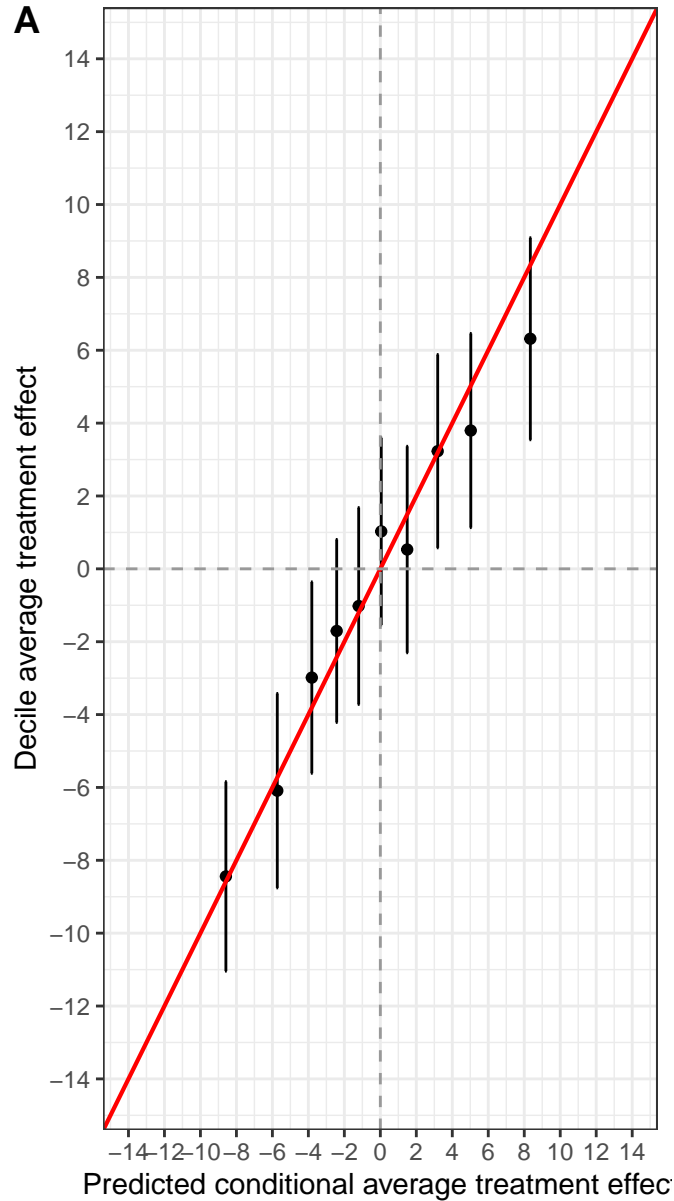
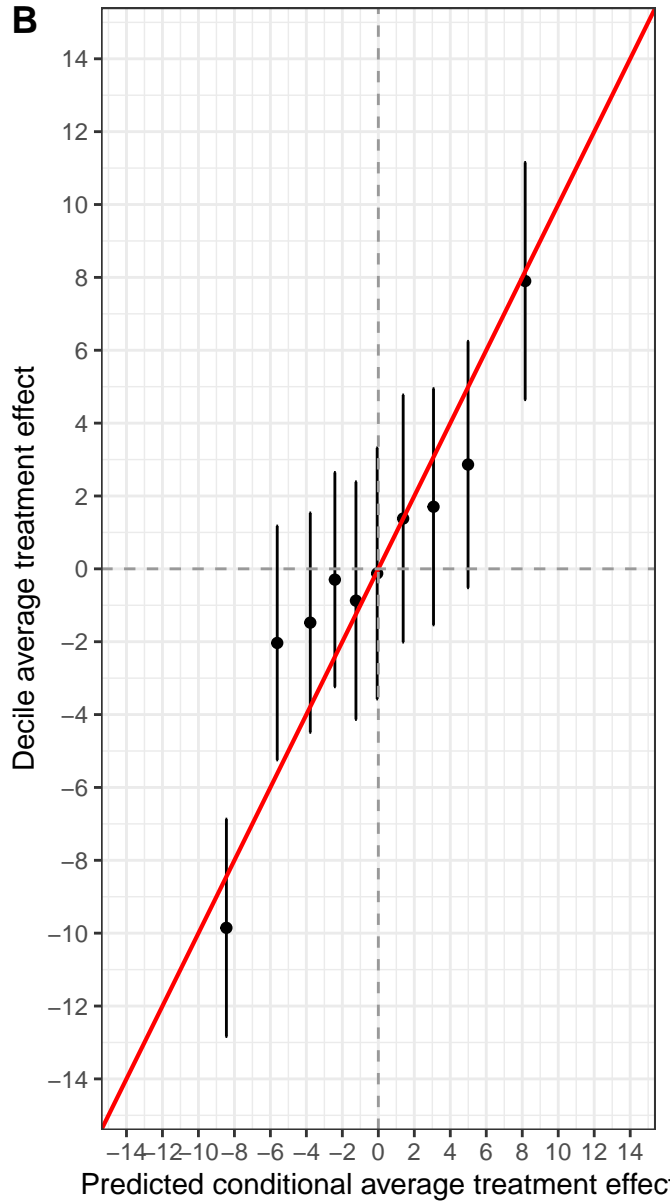
Effect submodels

A**B**

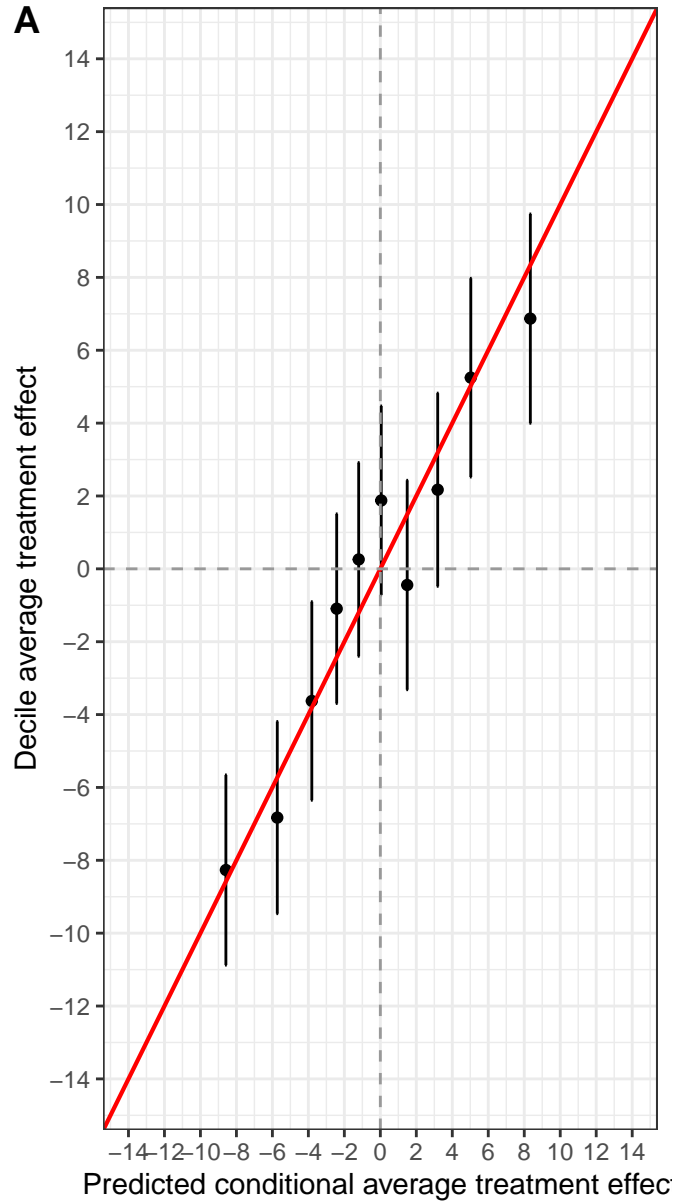
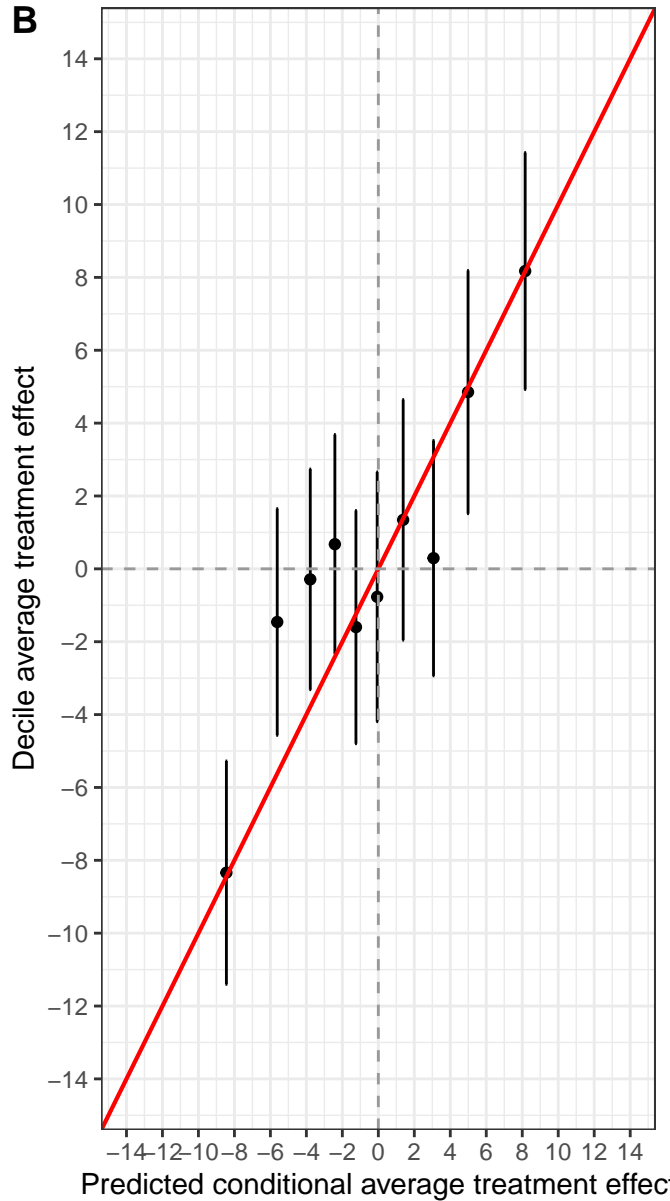
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

Effects validation prop score matching

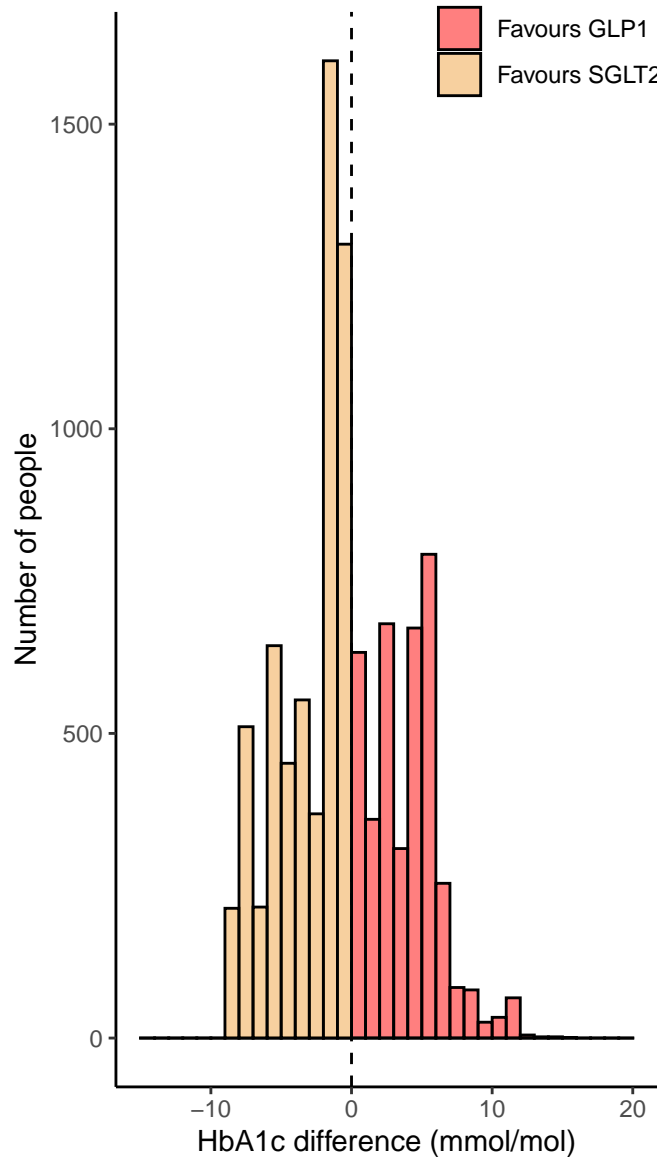
A**B**

Effects validation prop score inverse weighting

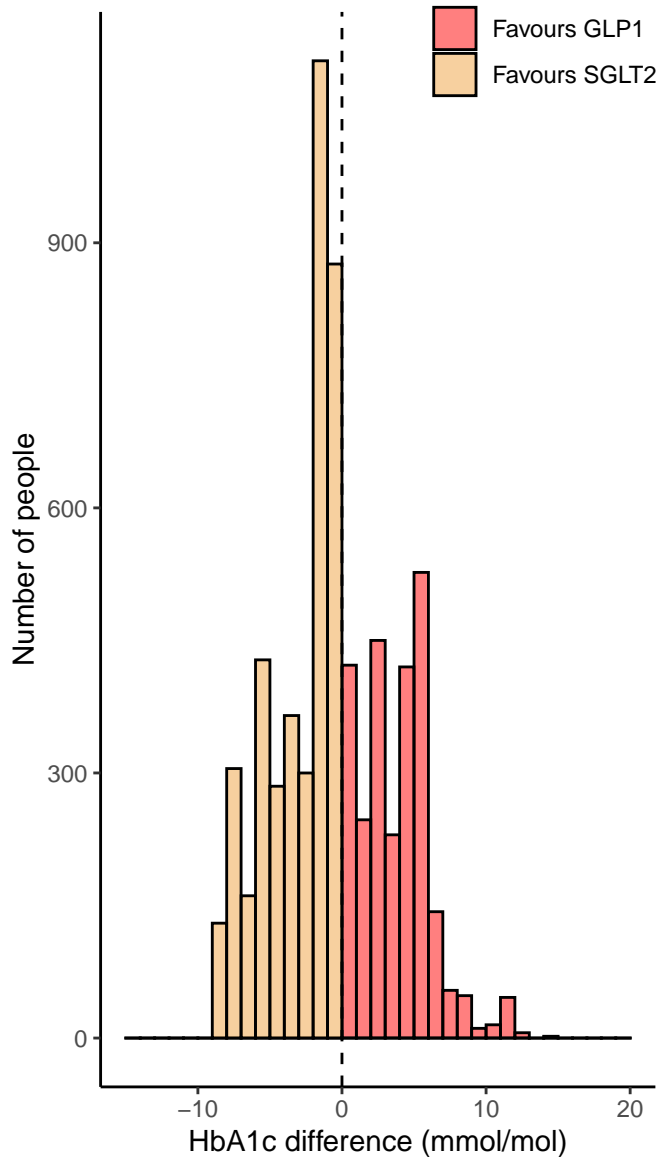
A**B**

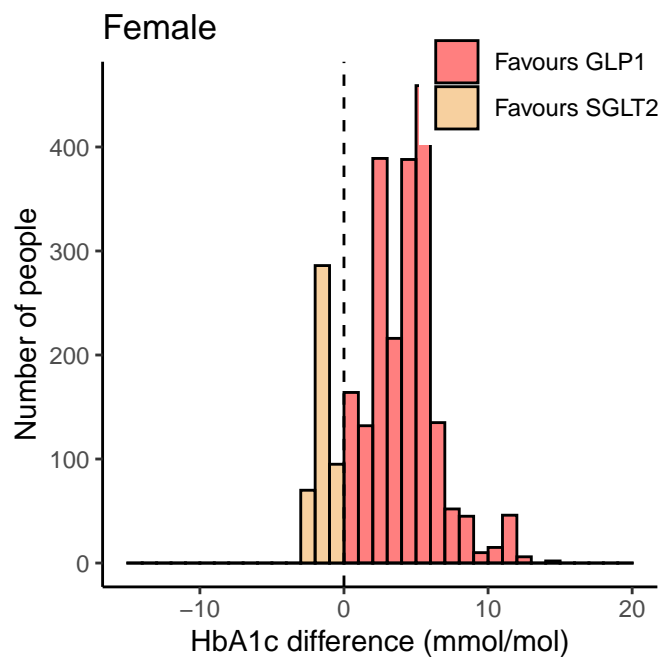
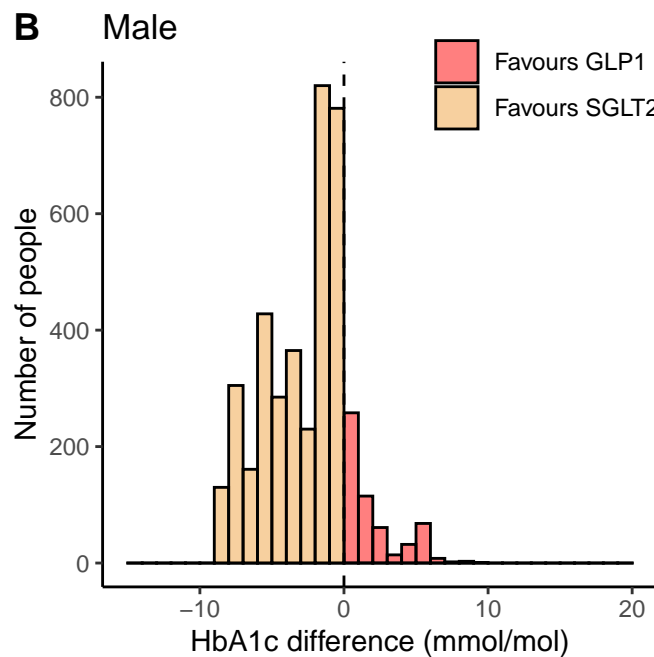
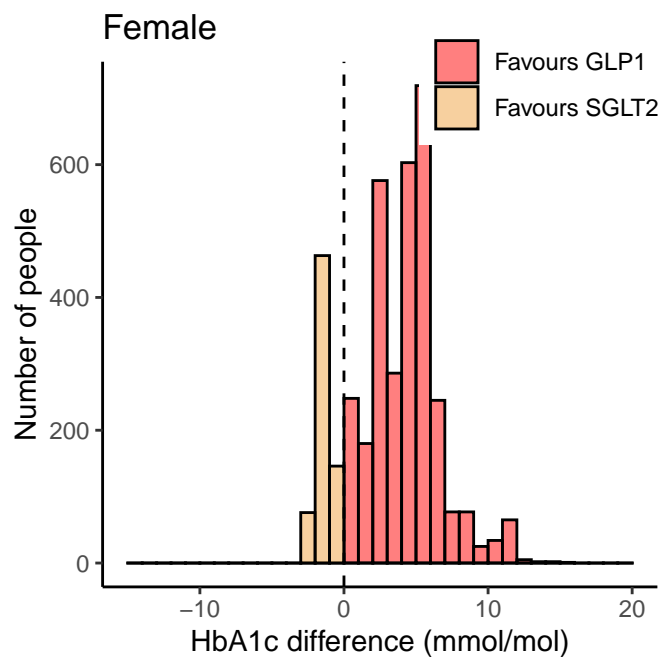
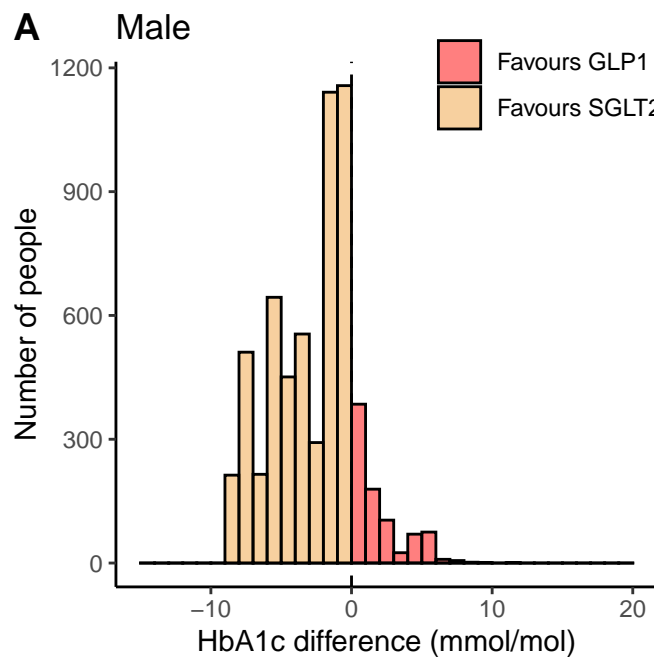
Model fitting: Incomplete data (No propensity score)

A

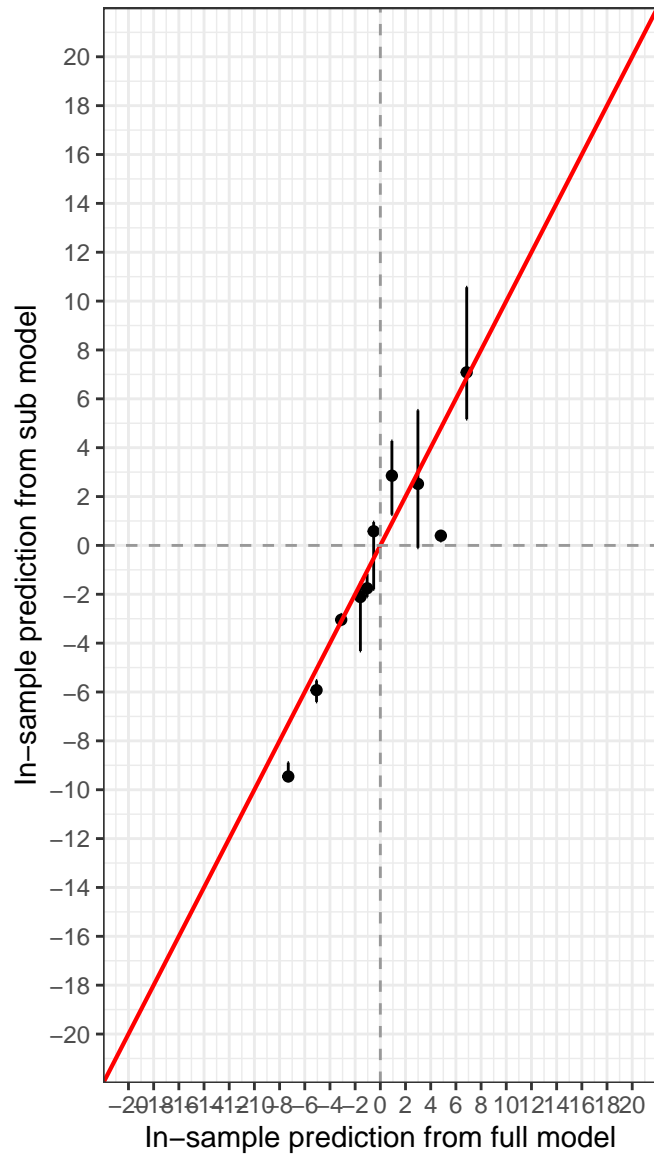
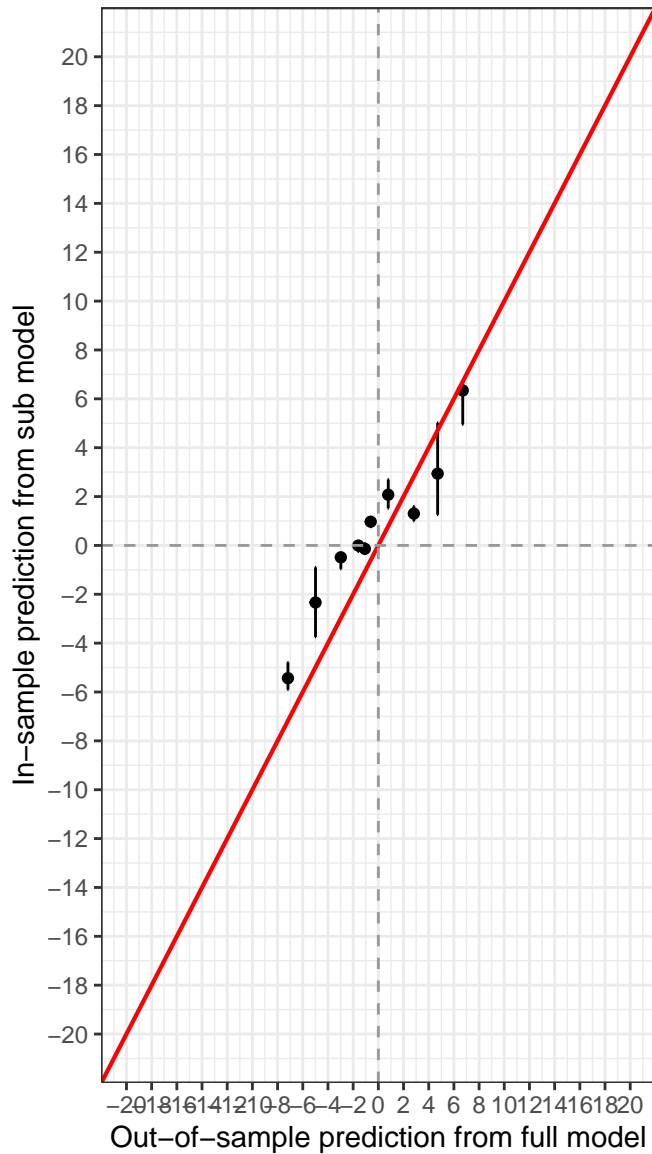


B

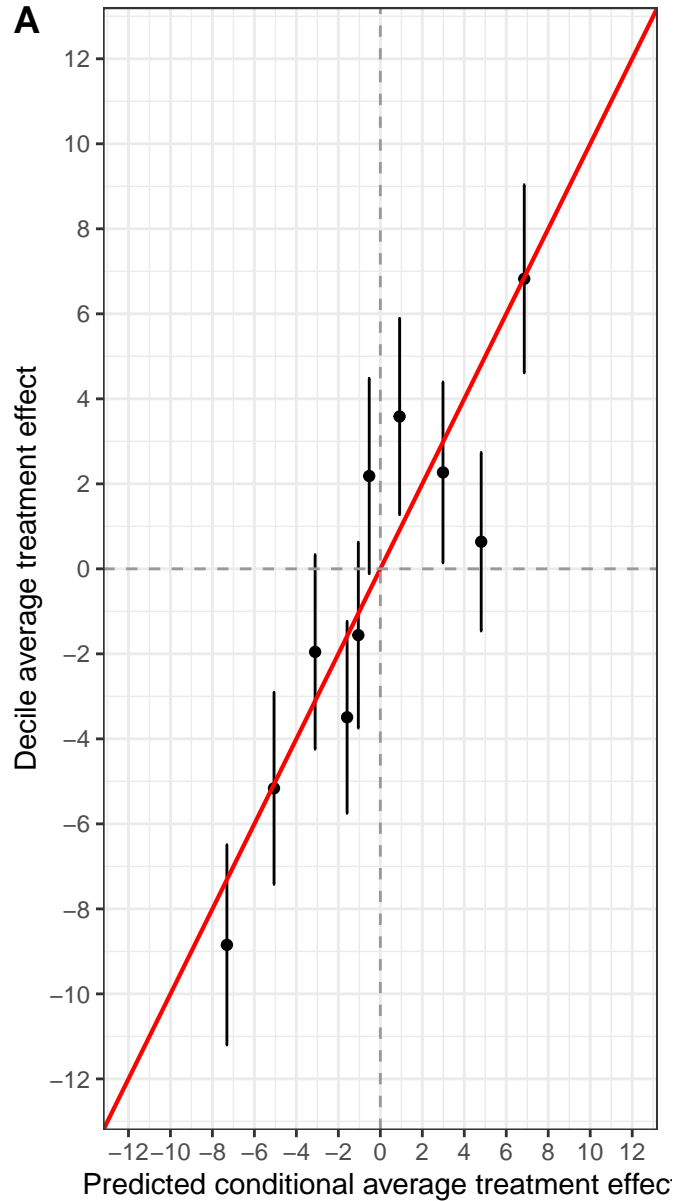
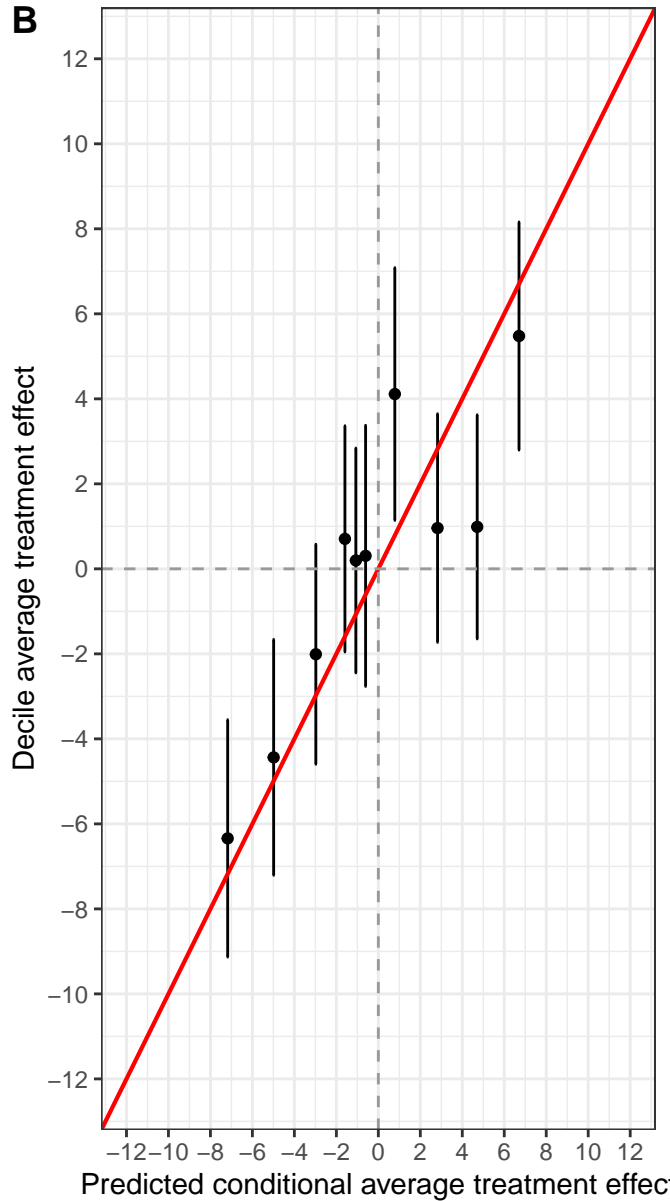




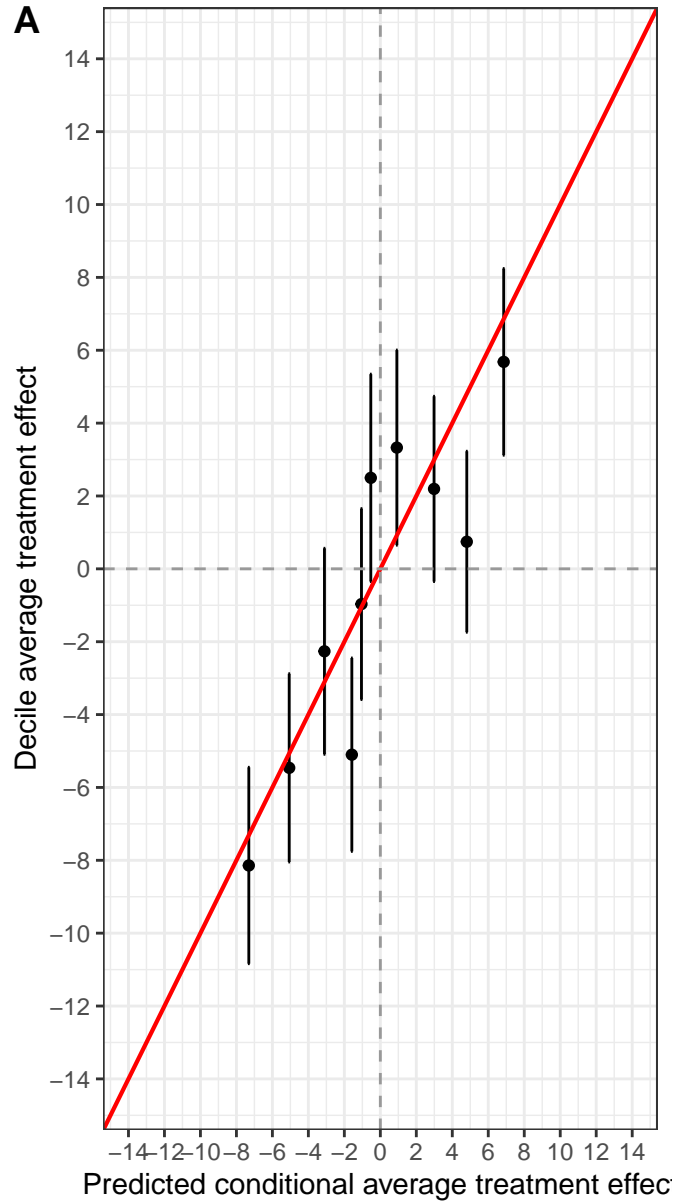
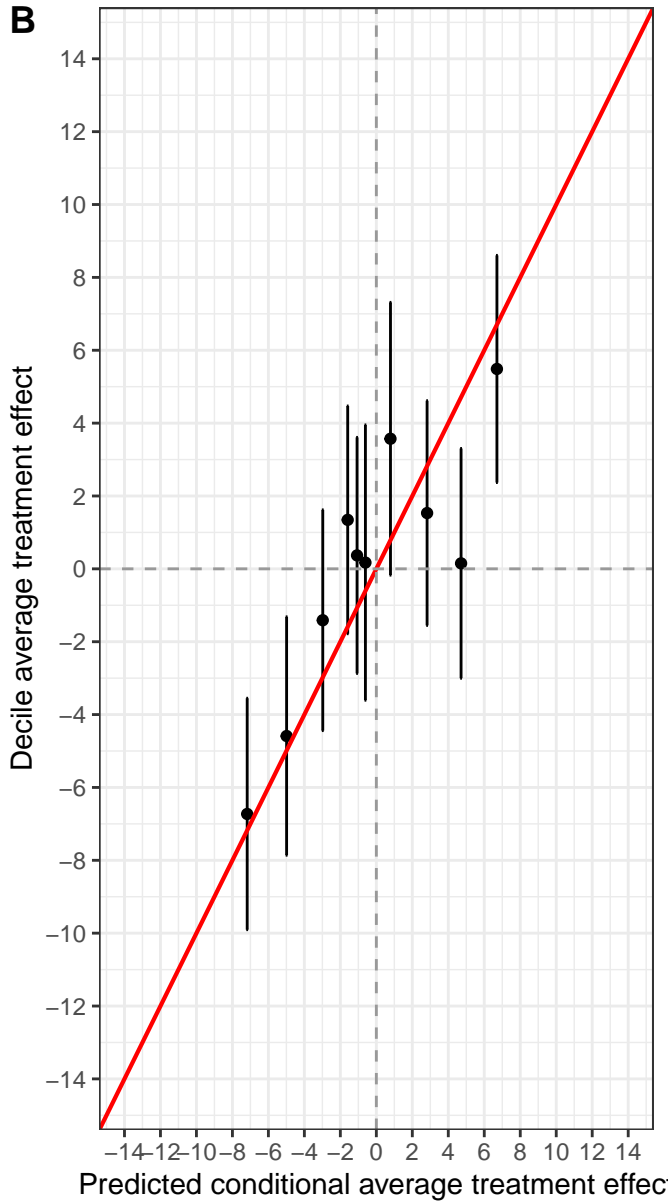
Effect submodels

A**B**

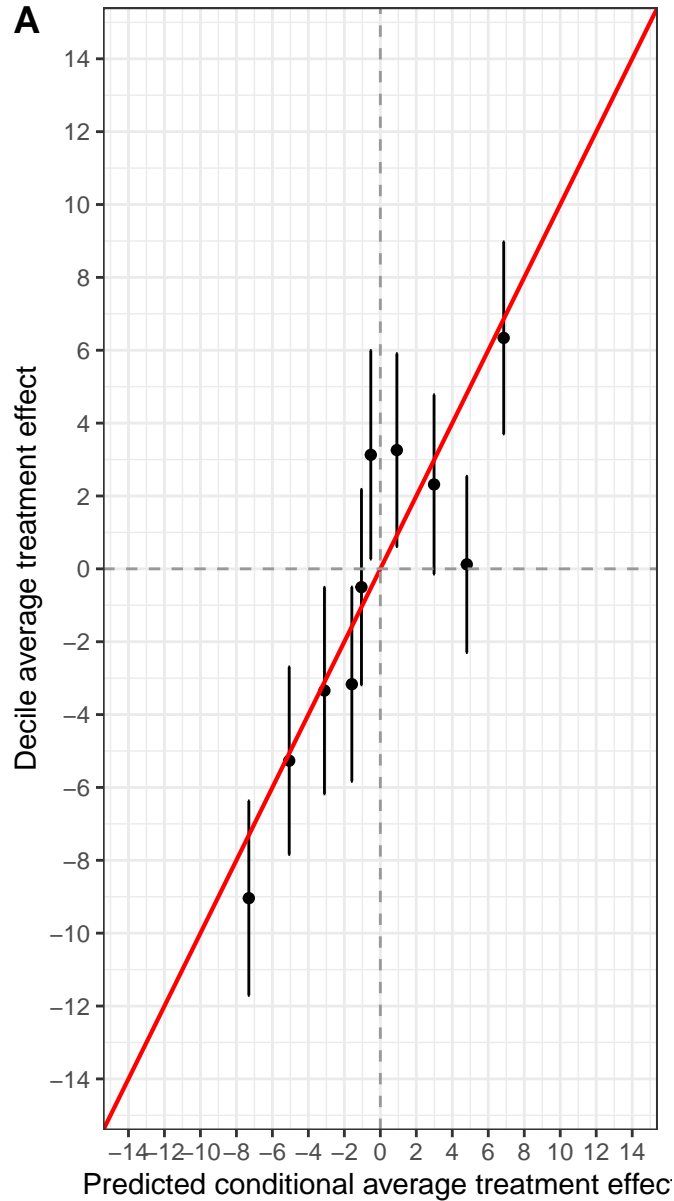
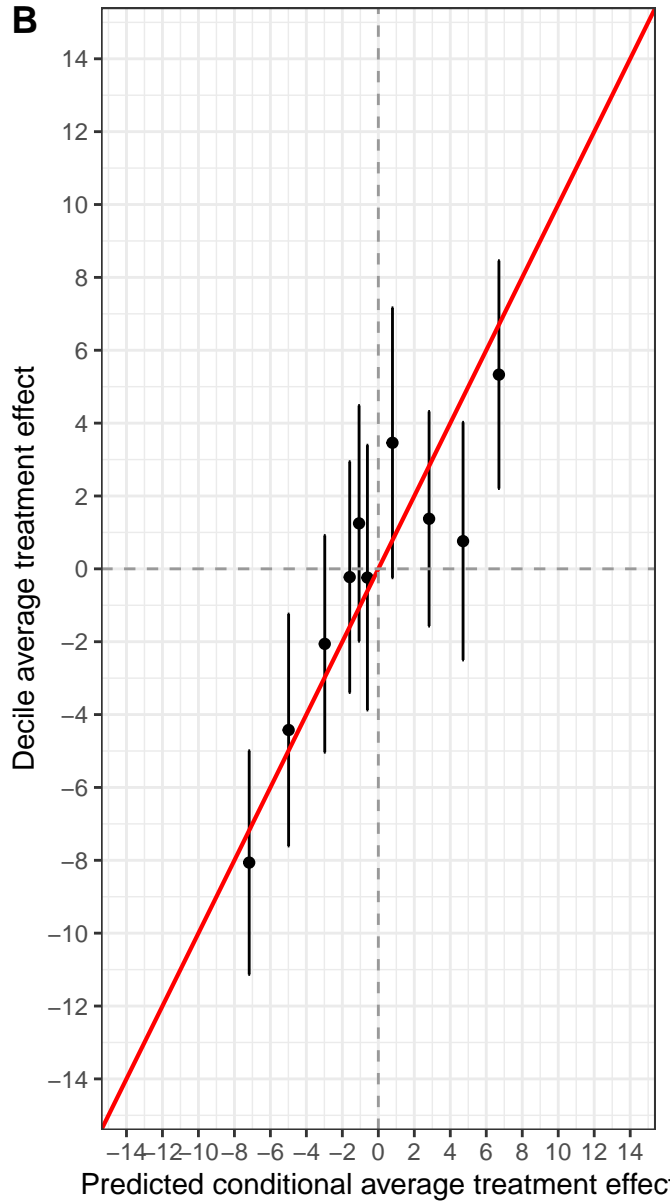
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

Effects validation prop score matching

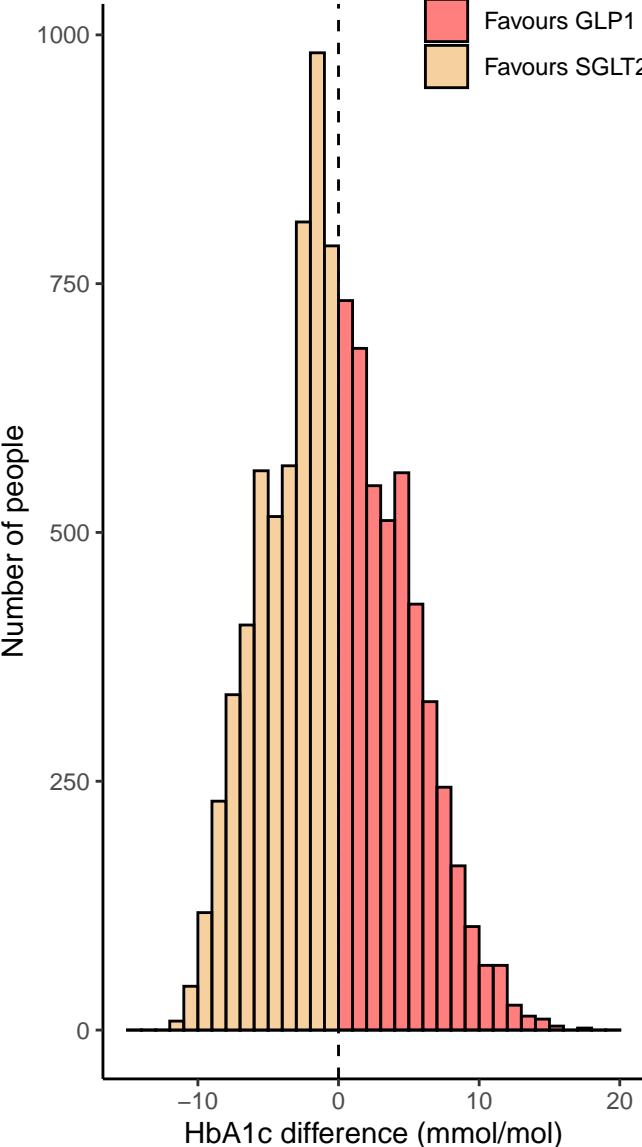
A**B**

Effects validation prop score inverse weighting

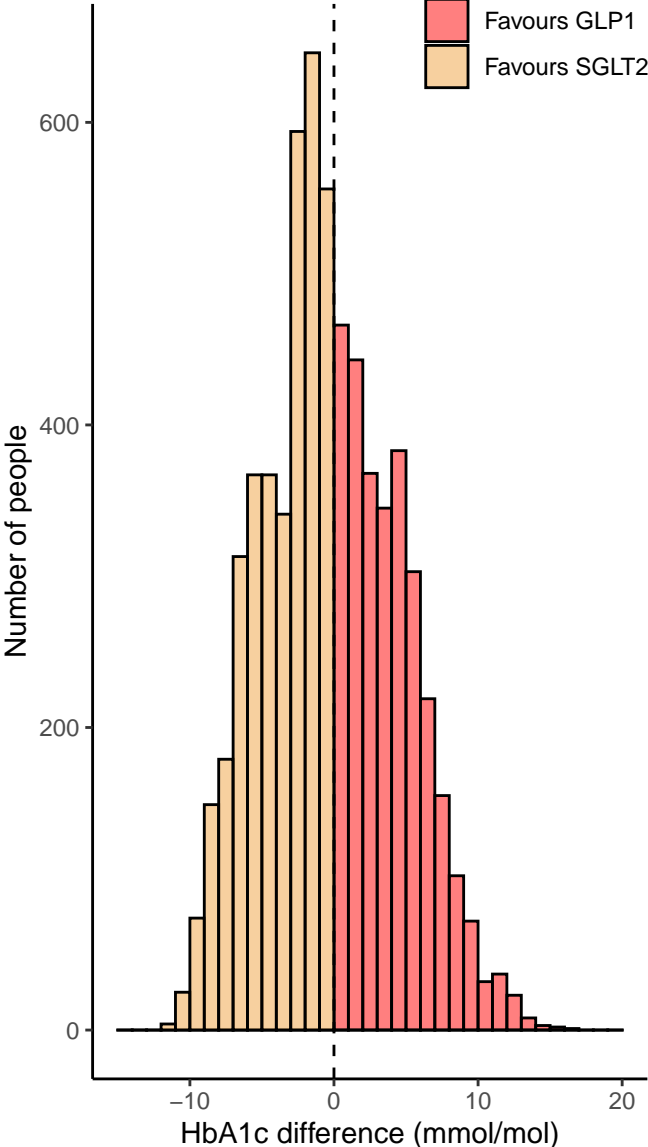
A**B**

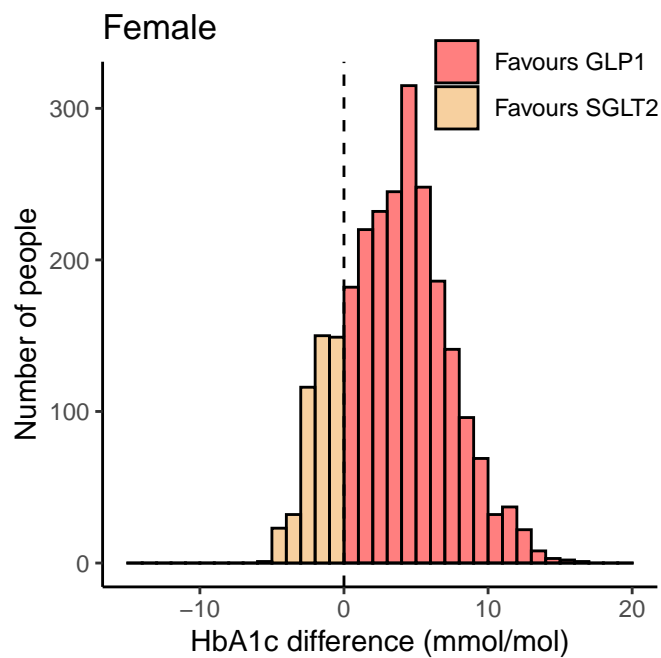
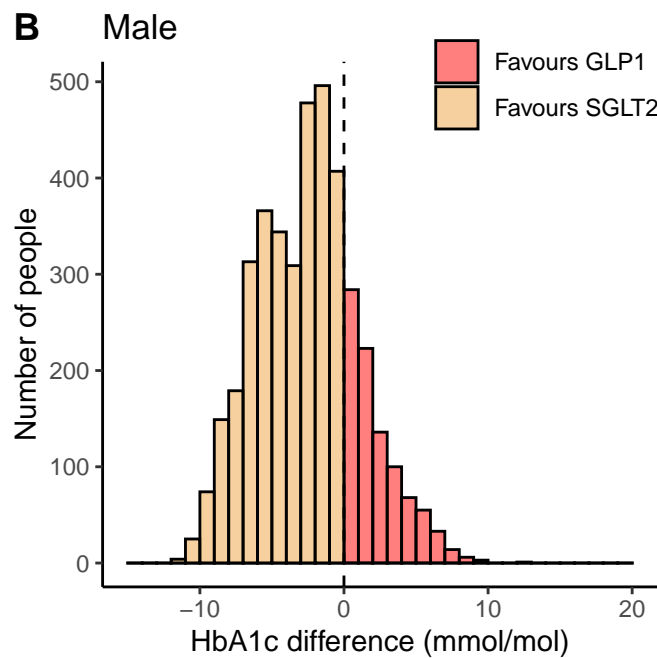
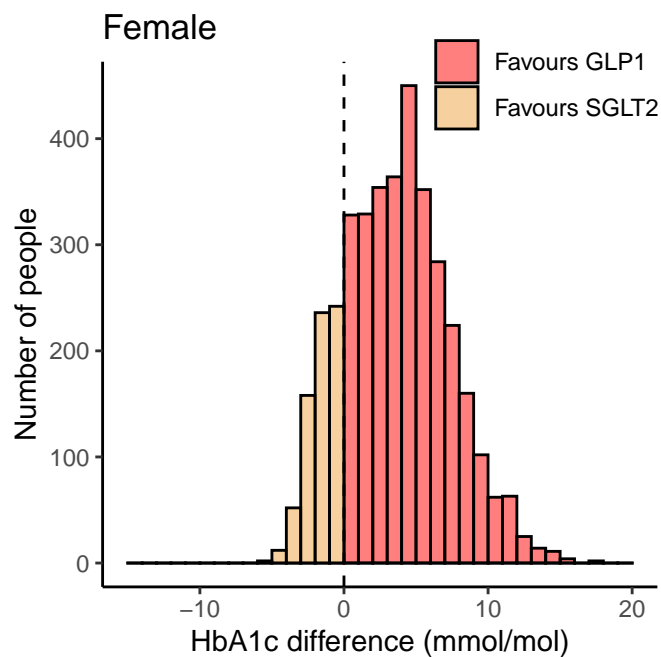
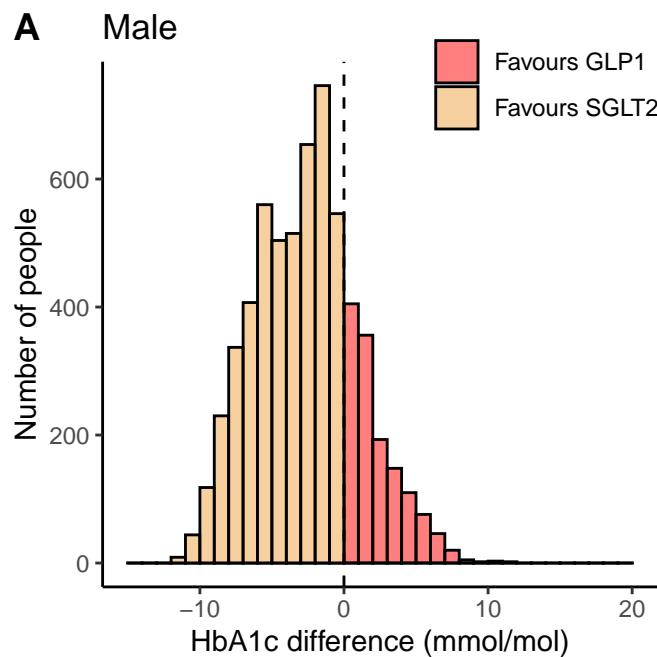
Model fitting: Variable Selection 1, Incomplete data (No propensity score)

A

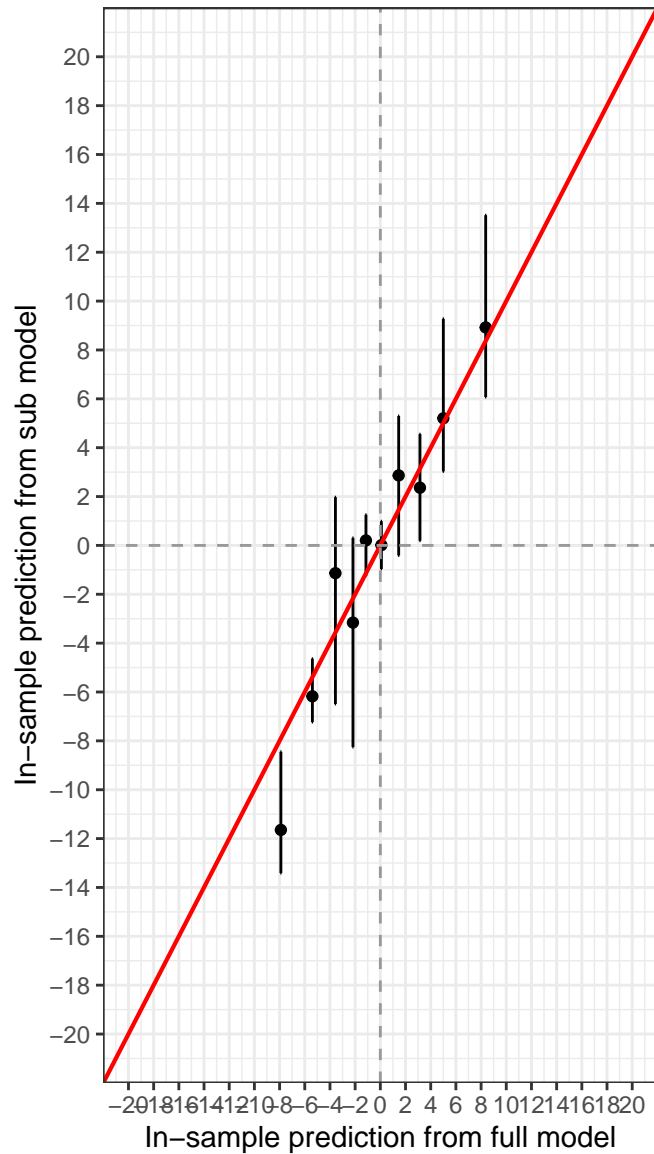
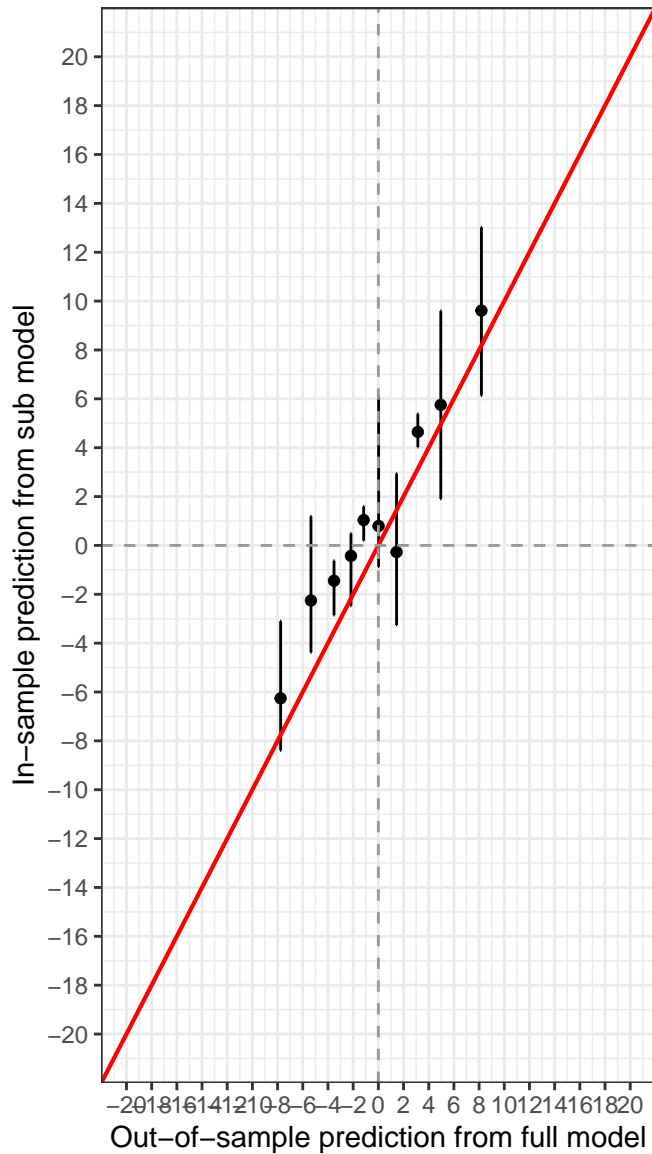


B

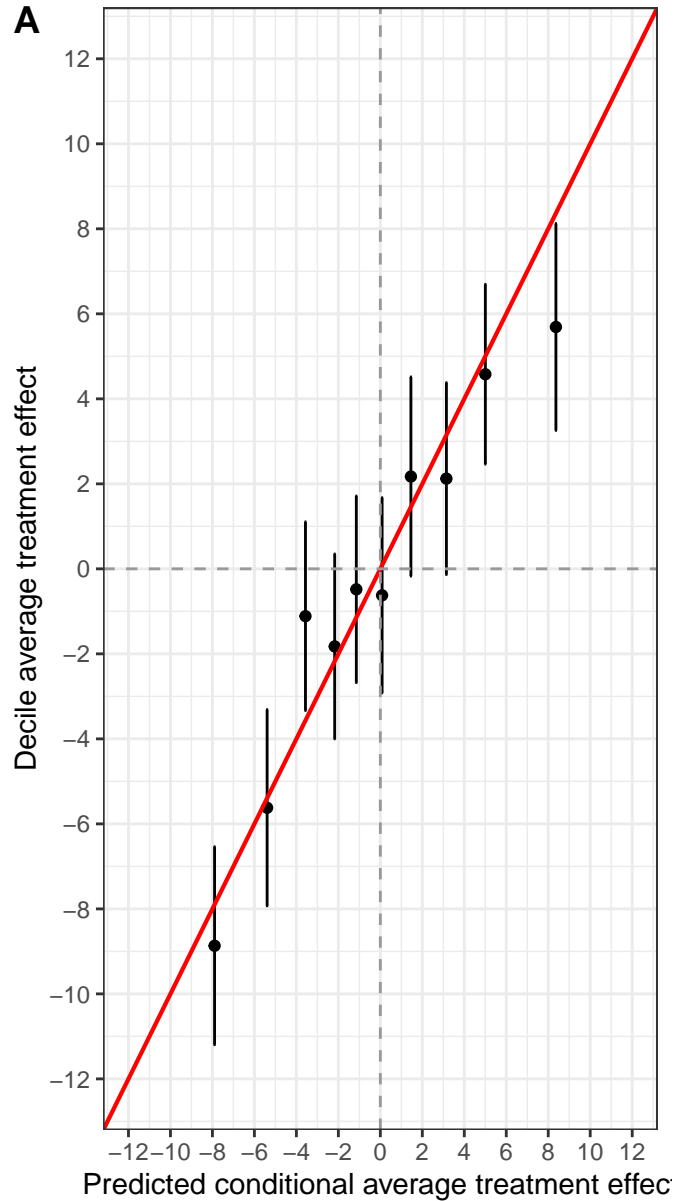
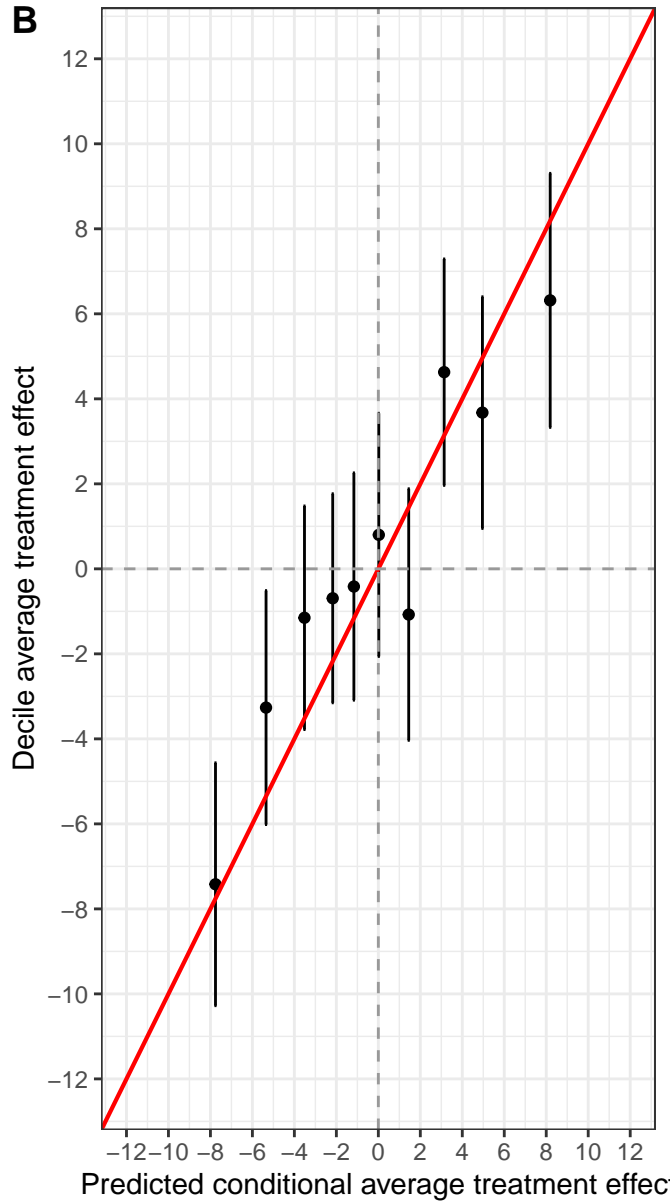




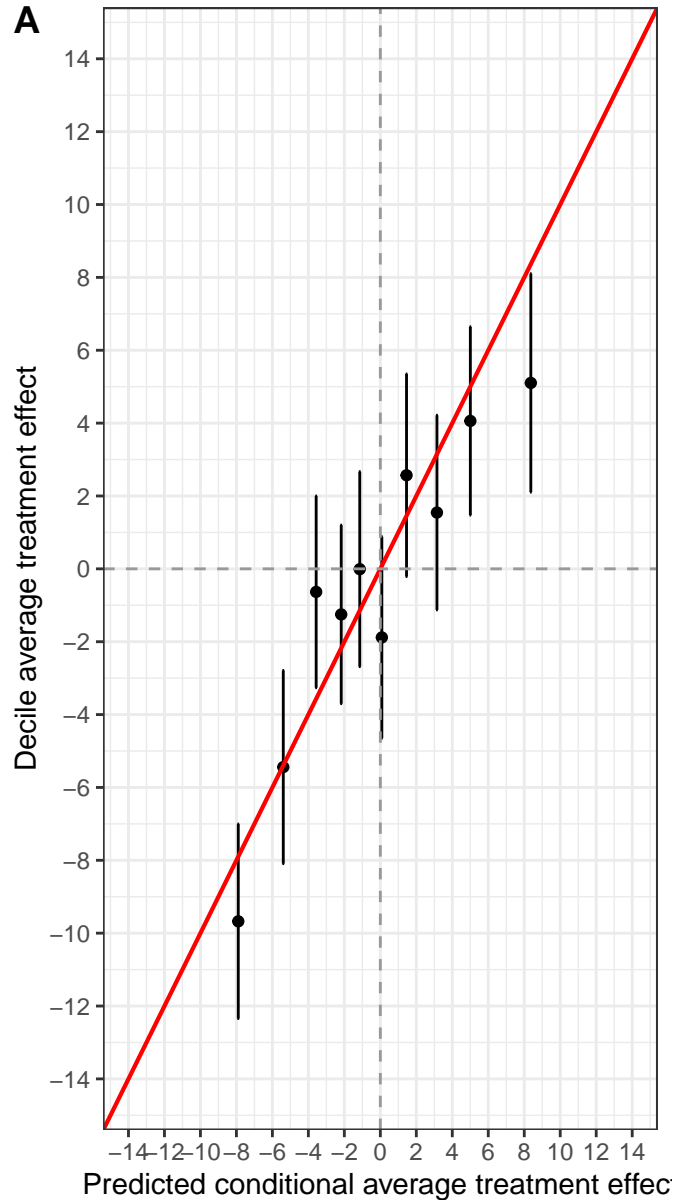
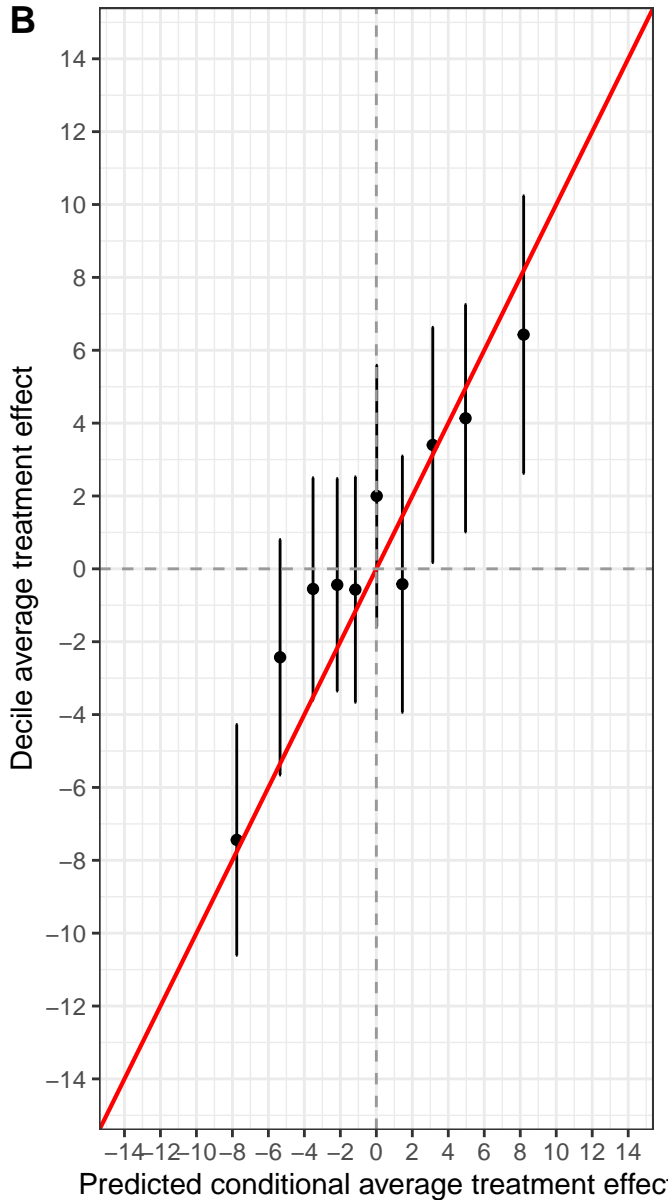
Effect submodels

A**B**

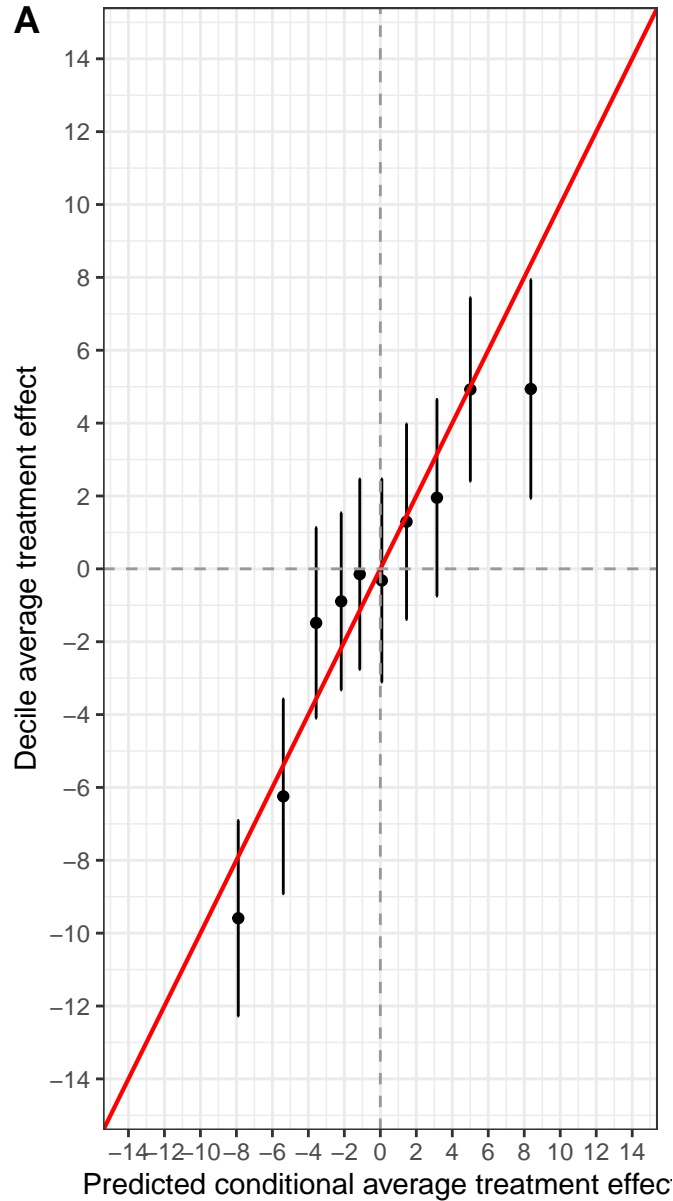
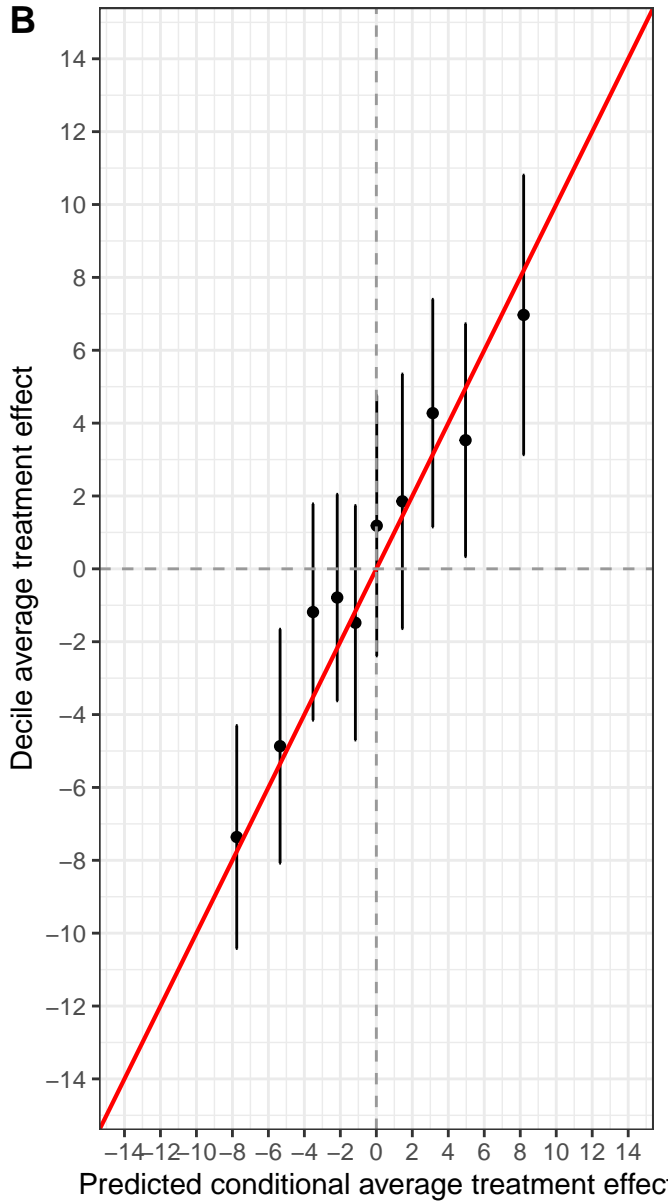
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

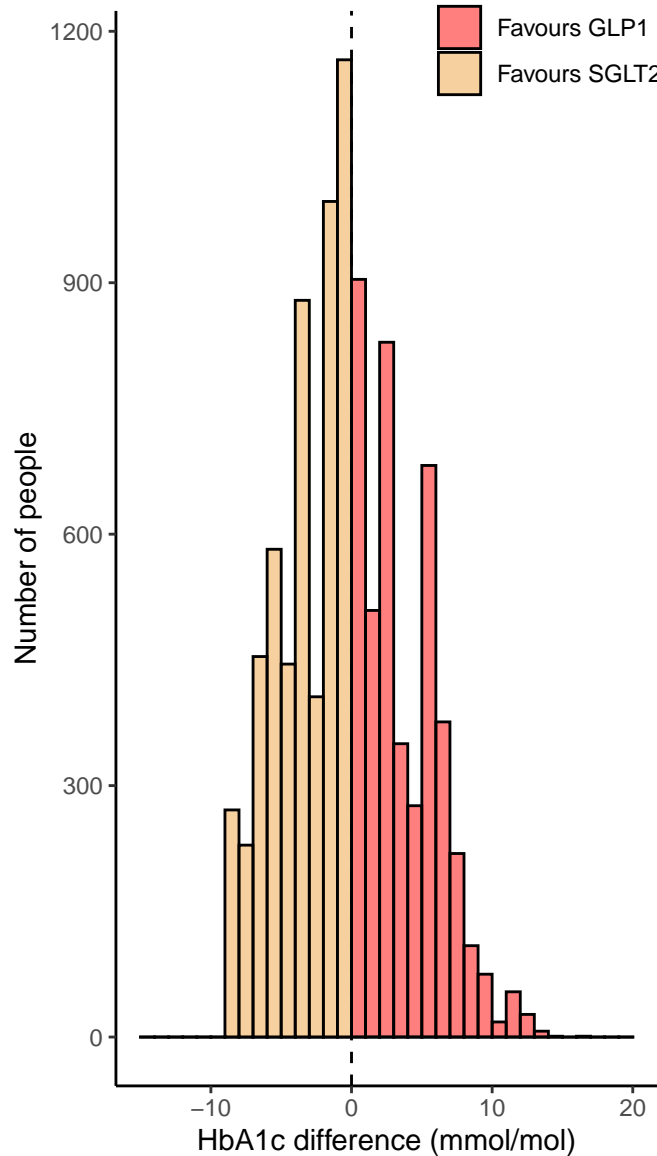
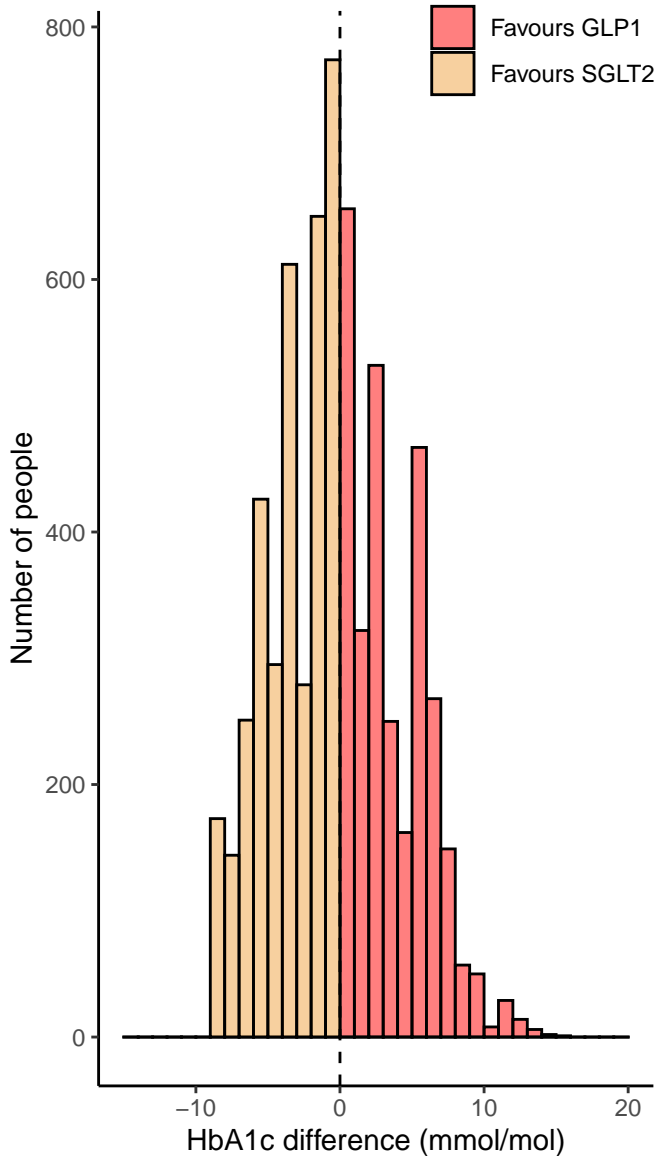
Effects validation prop score matching

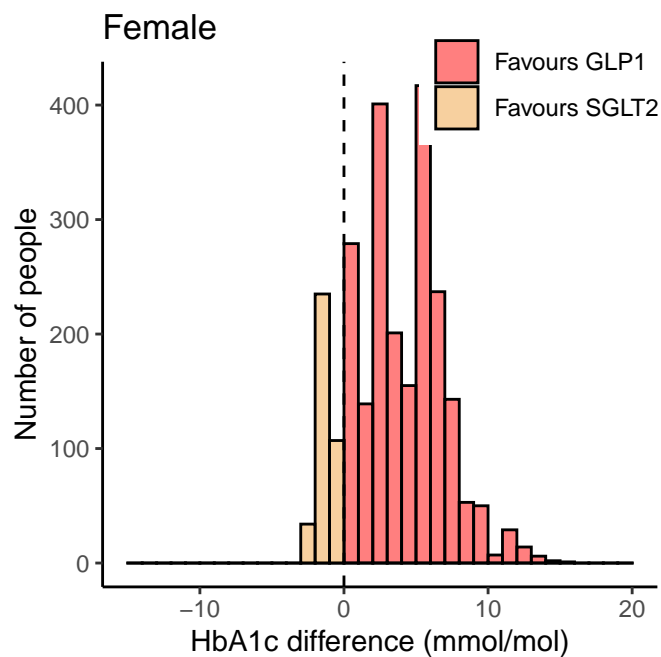
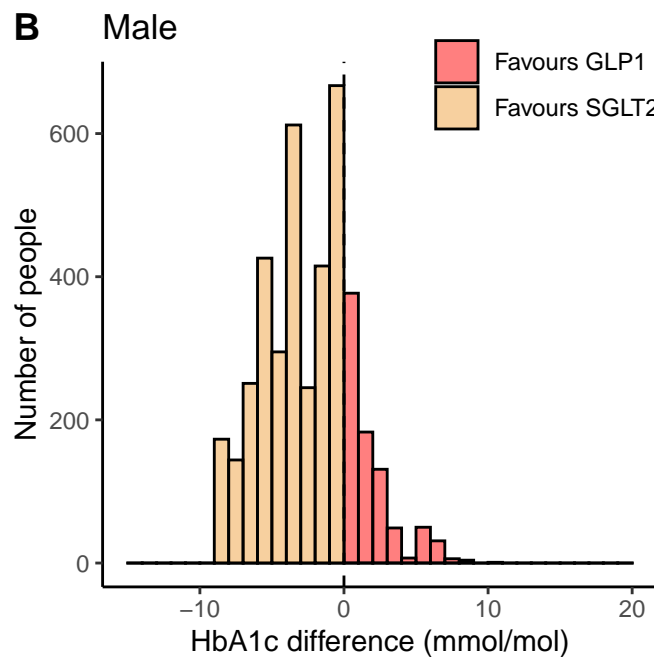
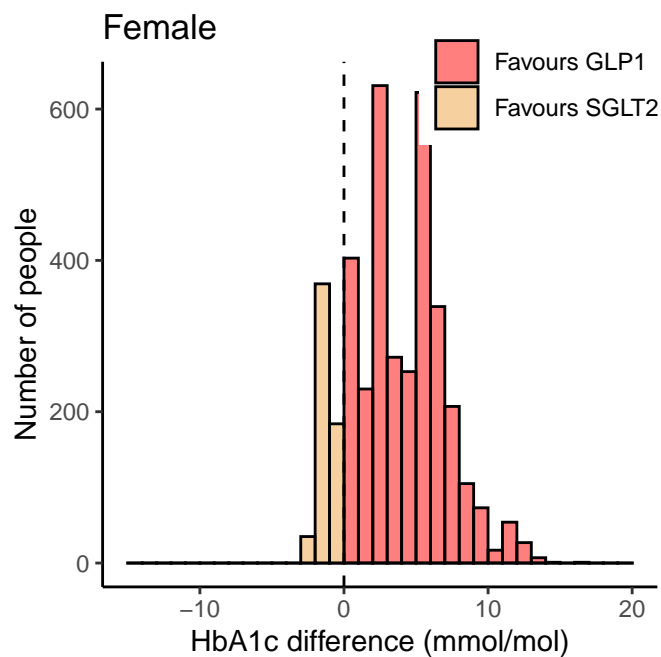
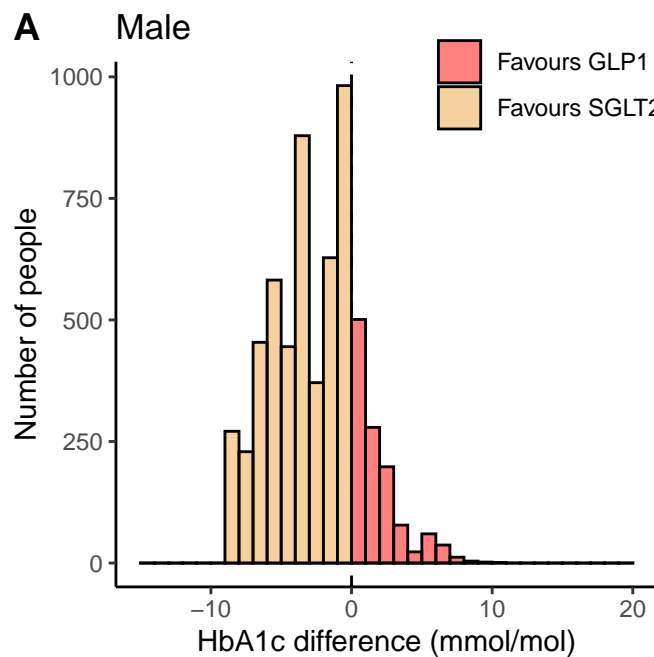
A**B**

Effects validation prop score inverse weighting

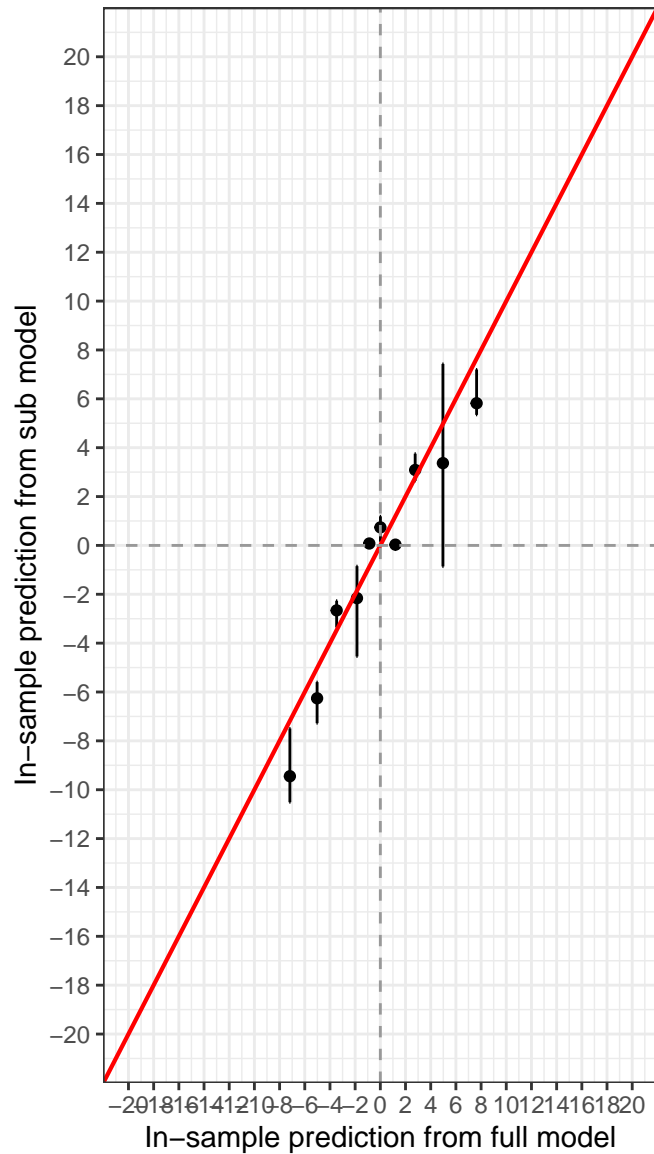
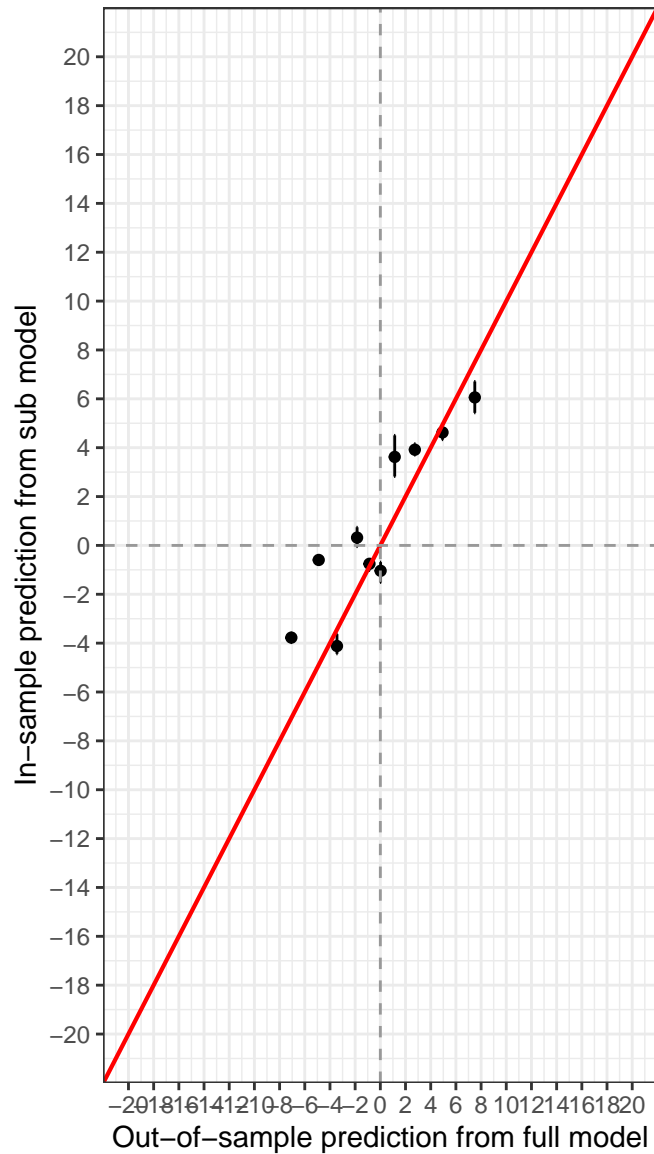
A**B**

Model fitting: Incomplete data (Propensity score)

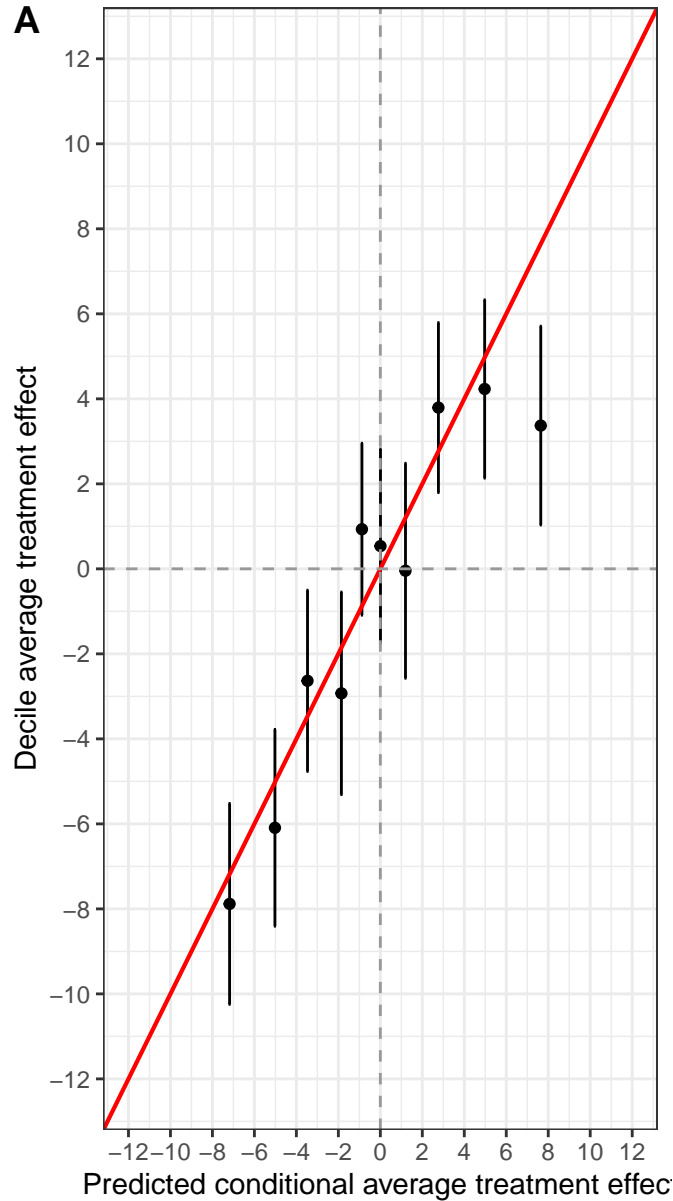
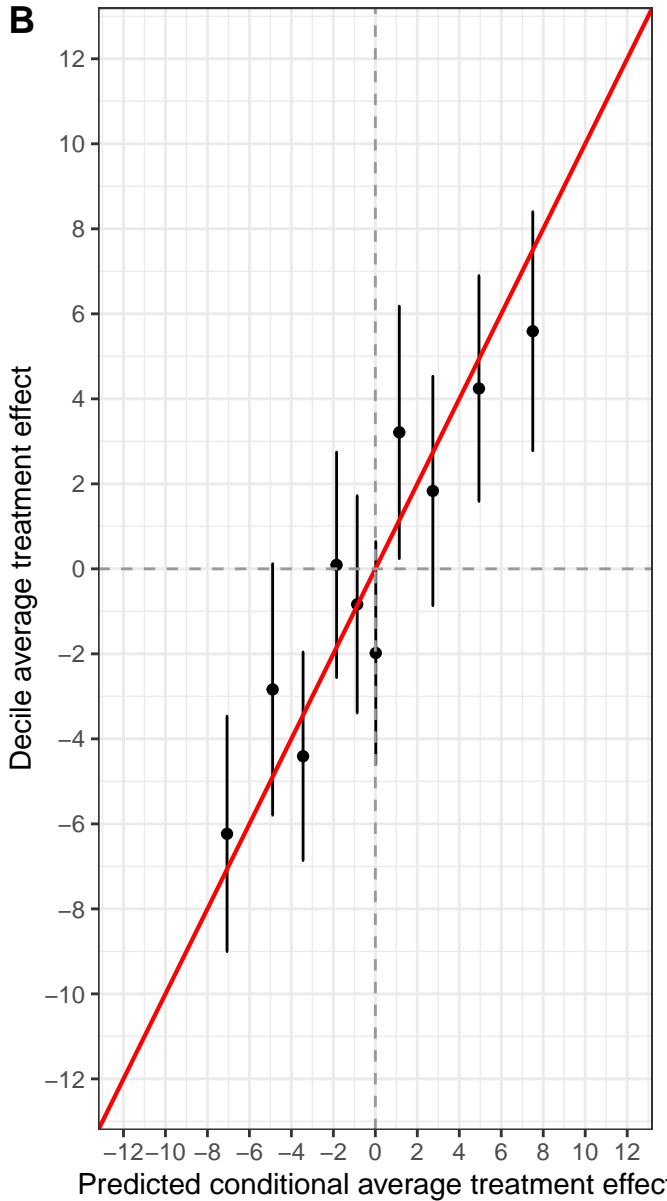
A**B**



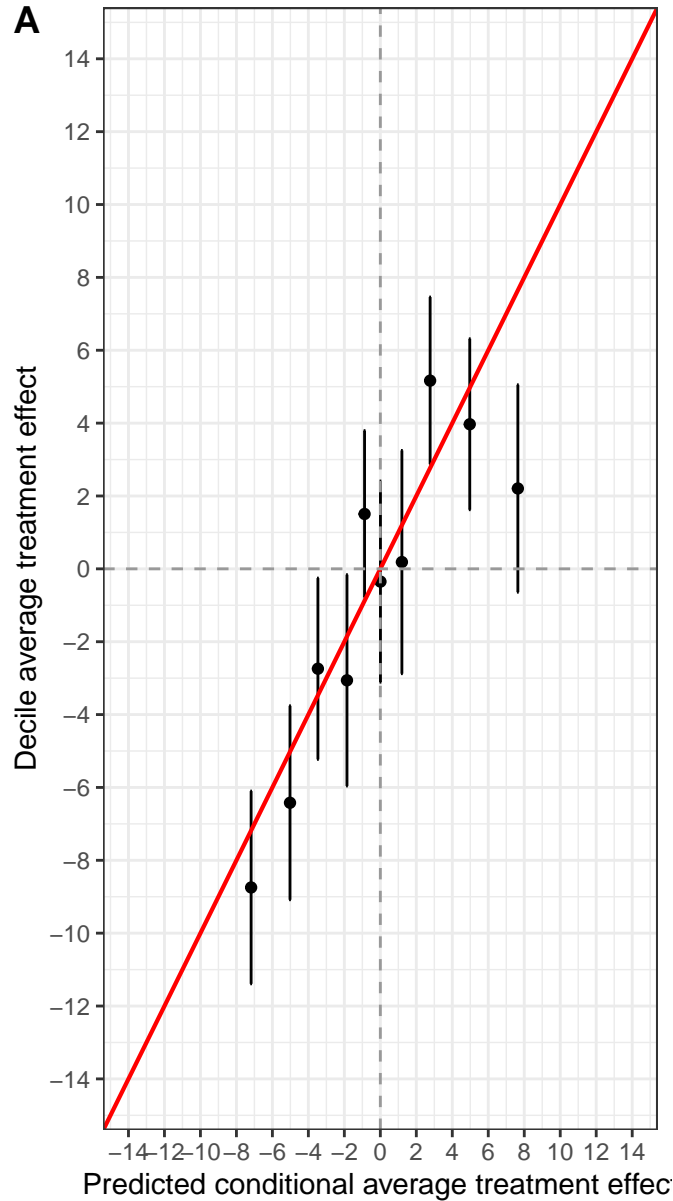
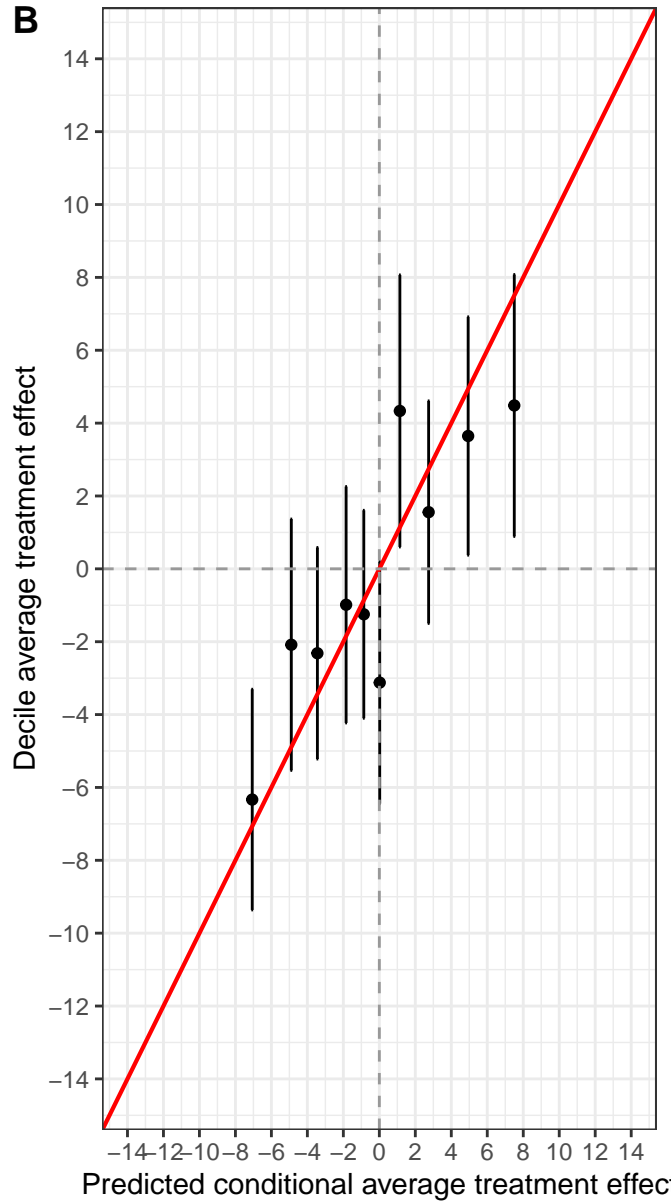
Effect submodels

A**B**

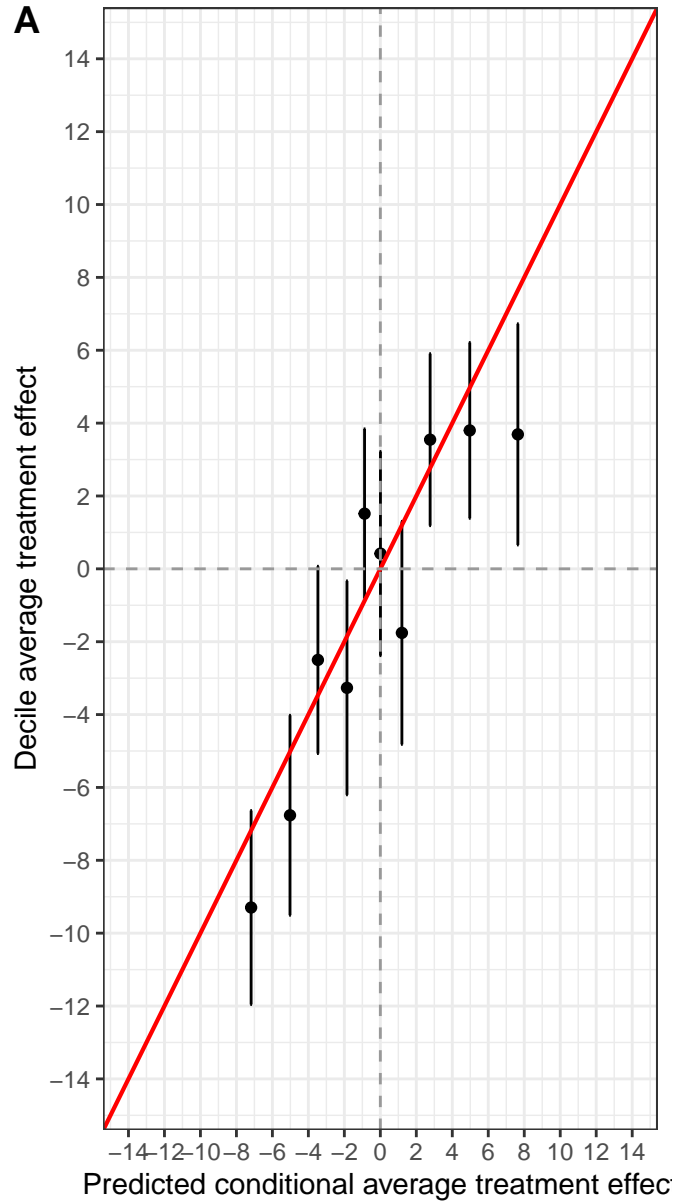
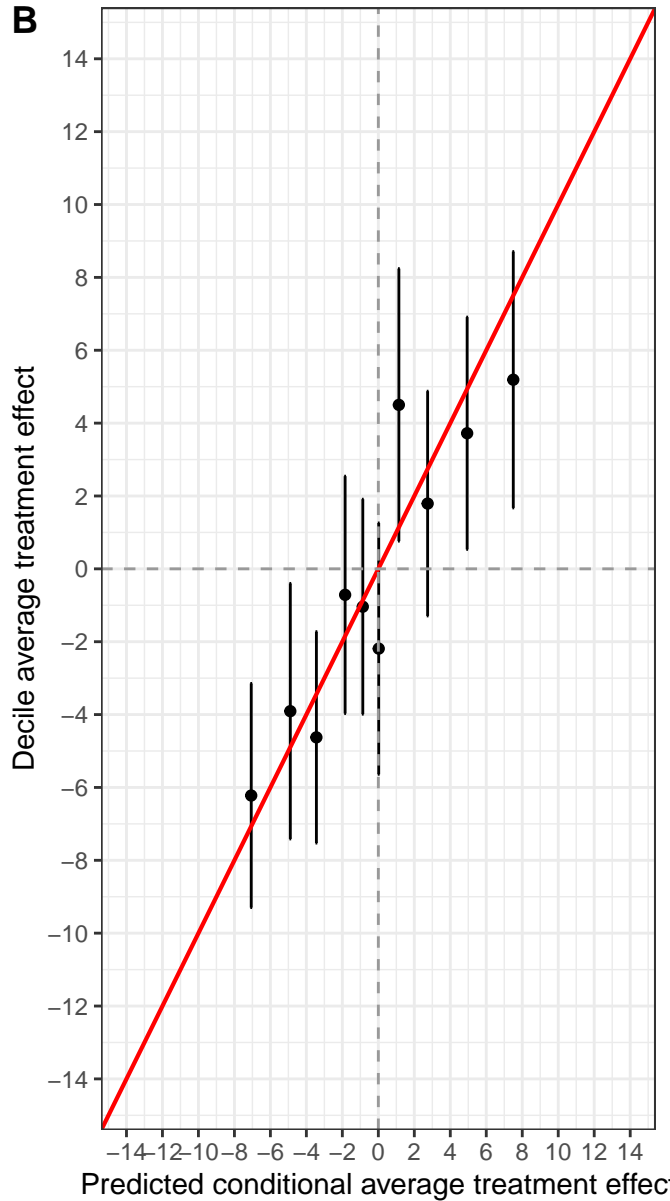
Effects validation Im(hba1c~drugclass+prop_score)

A**B**

Effects validation prop score matching

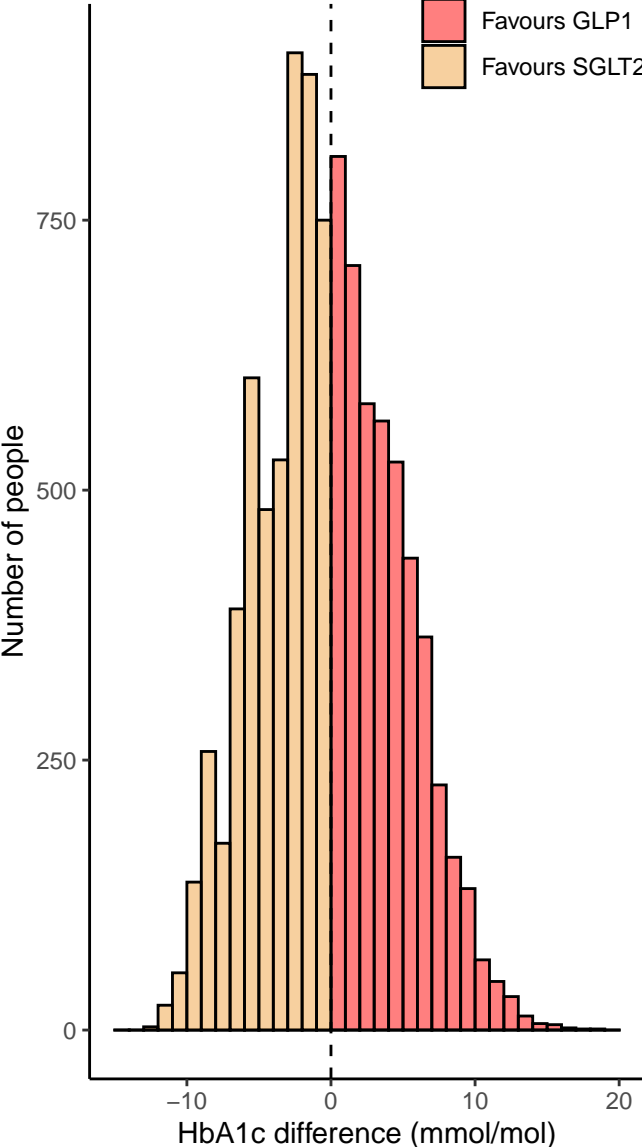
A**B**

Effects validation prop score inverse weighting

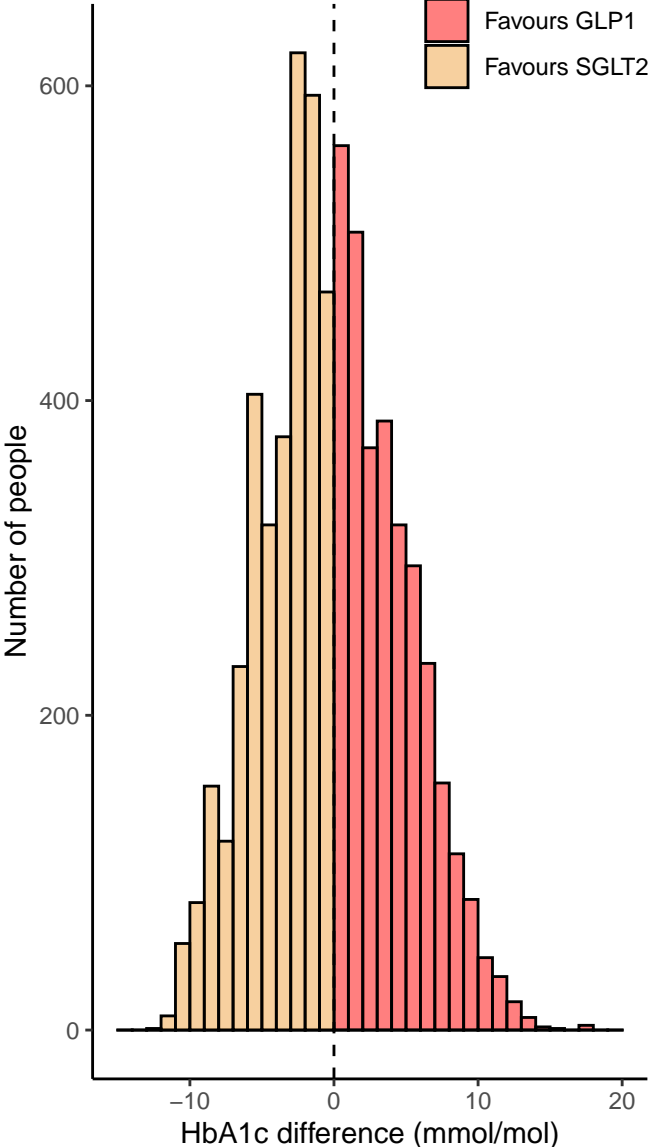
A**B**

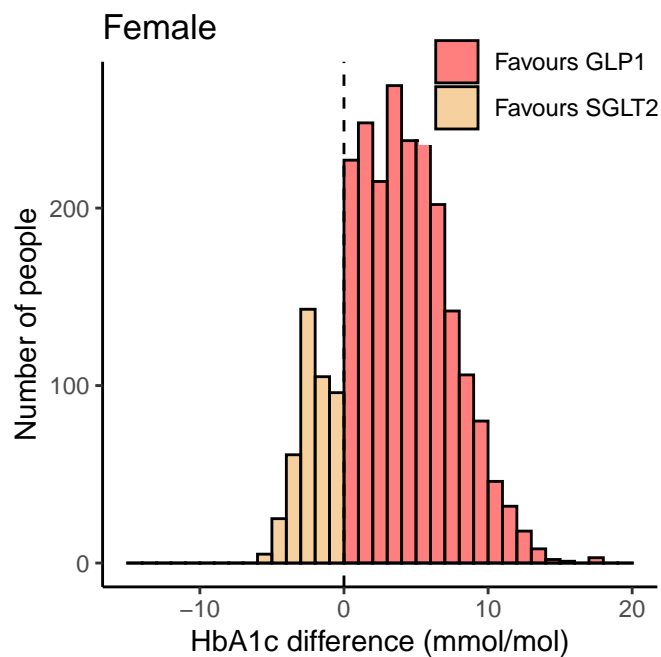
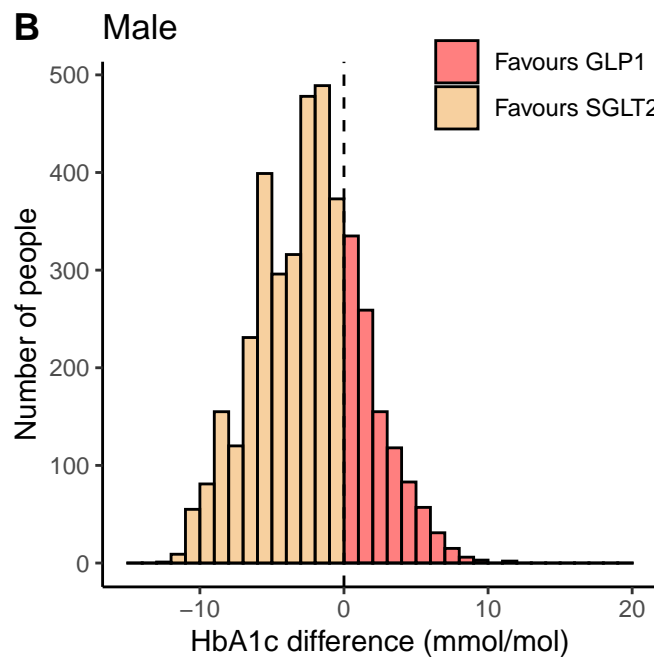
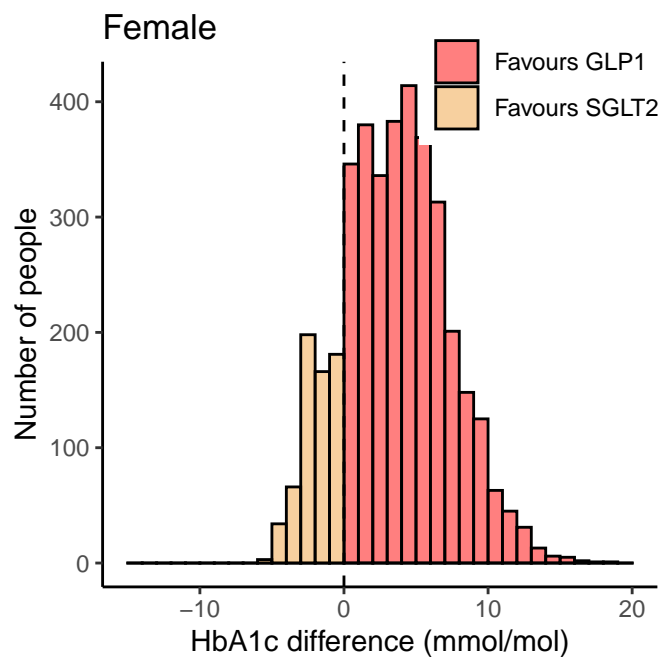
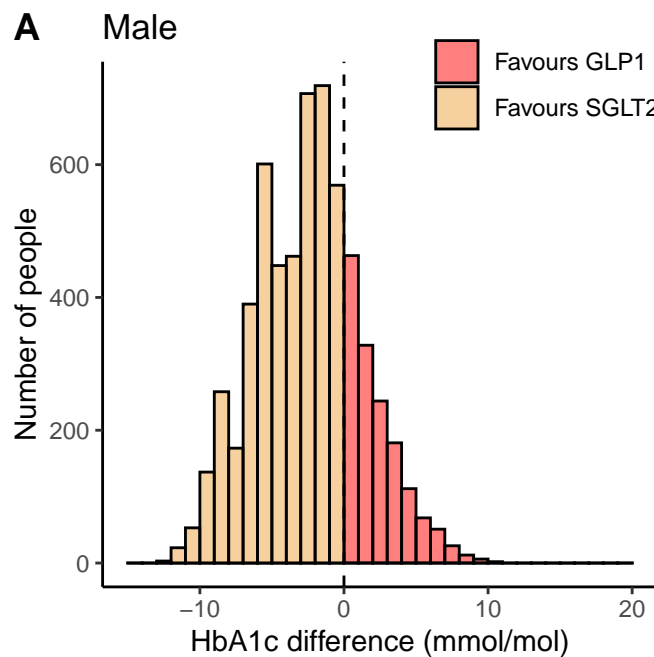
Model fitting: Variable Selection 1, Incomplete data (Propensity score)

A

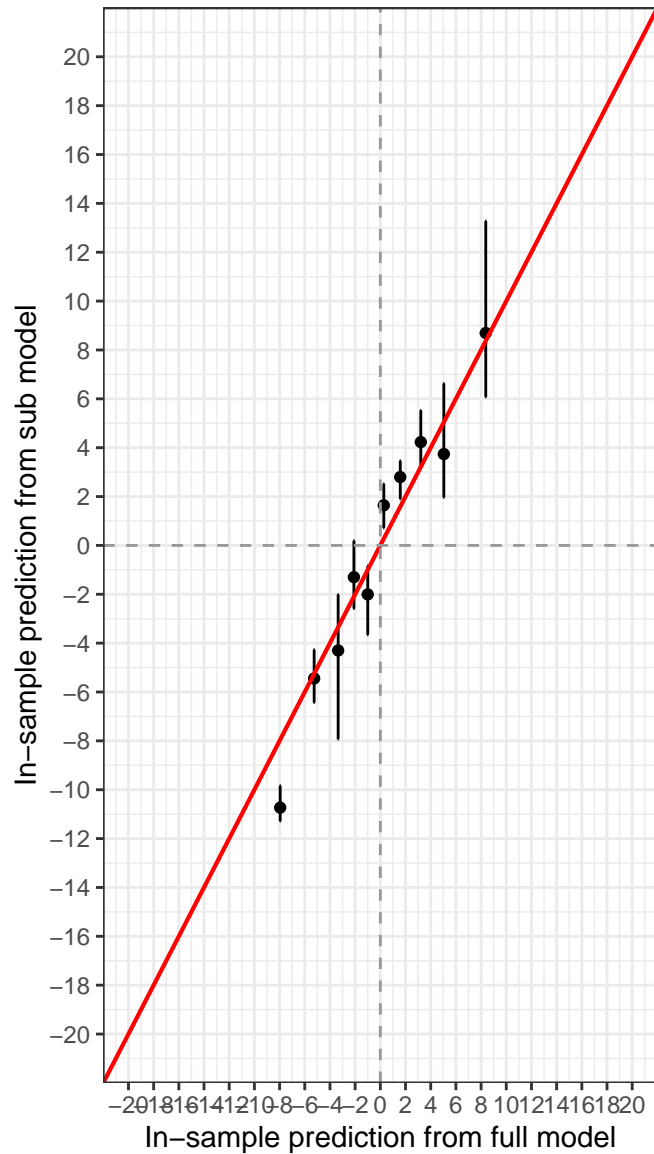
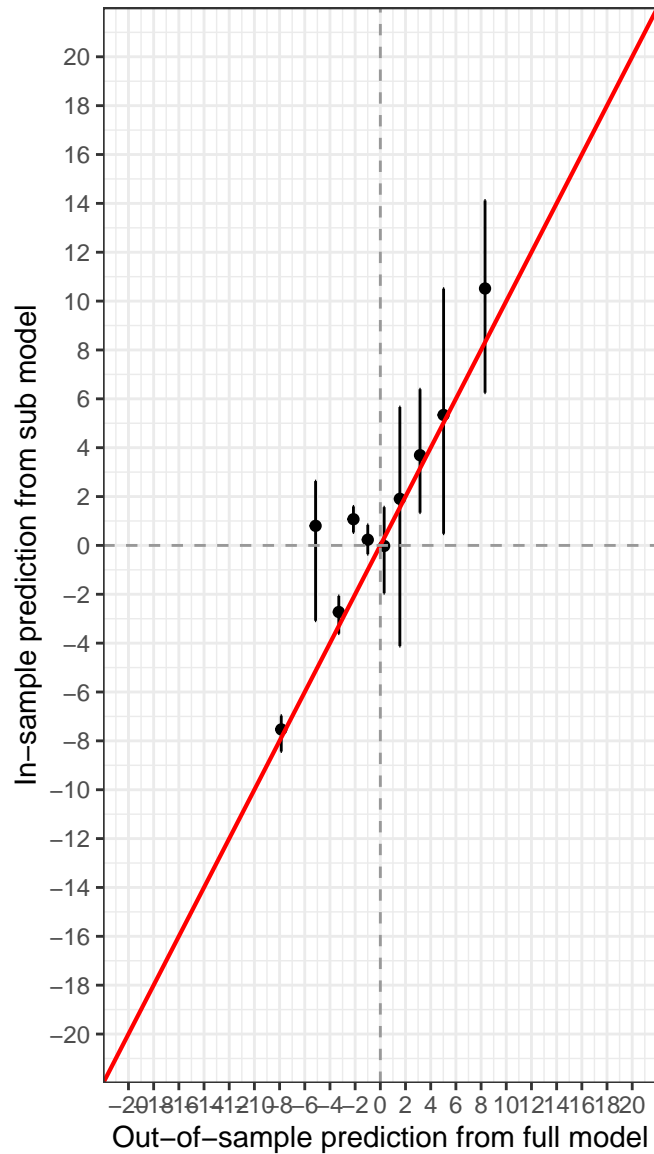


B

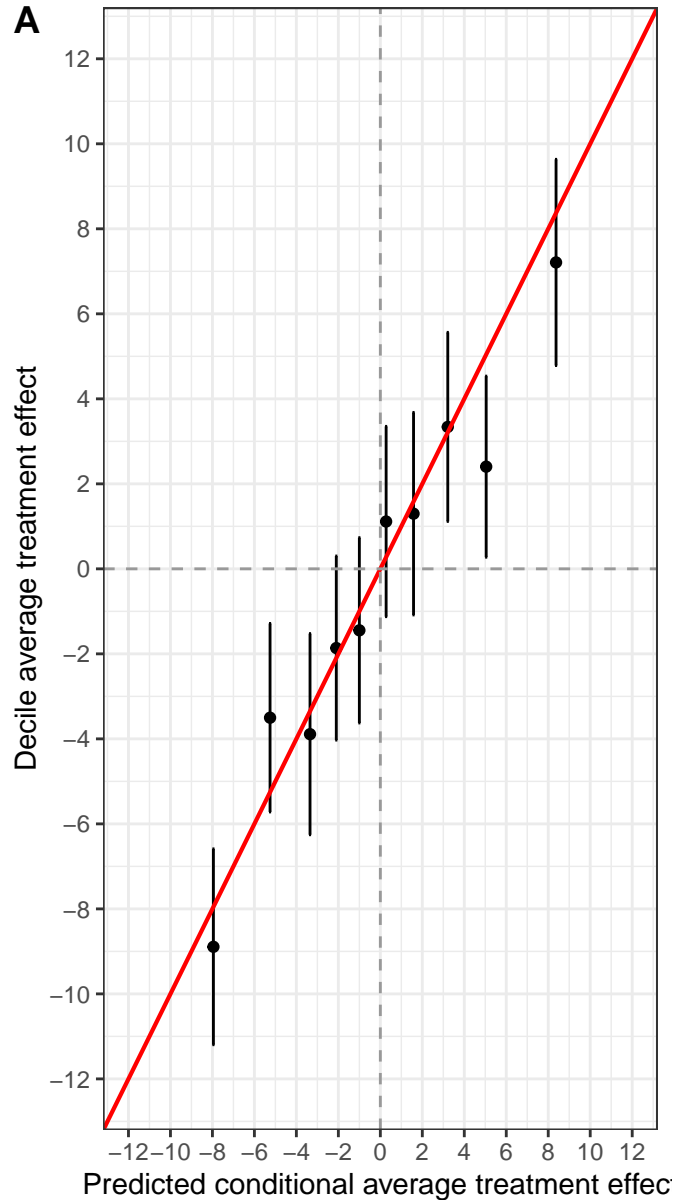
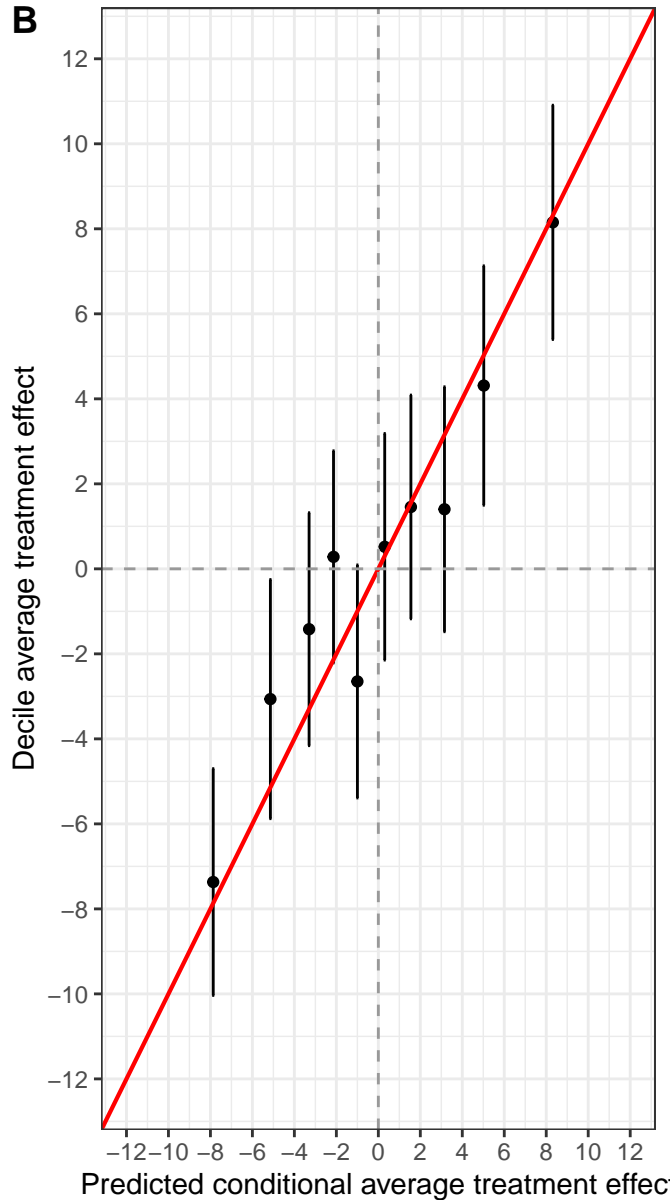




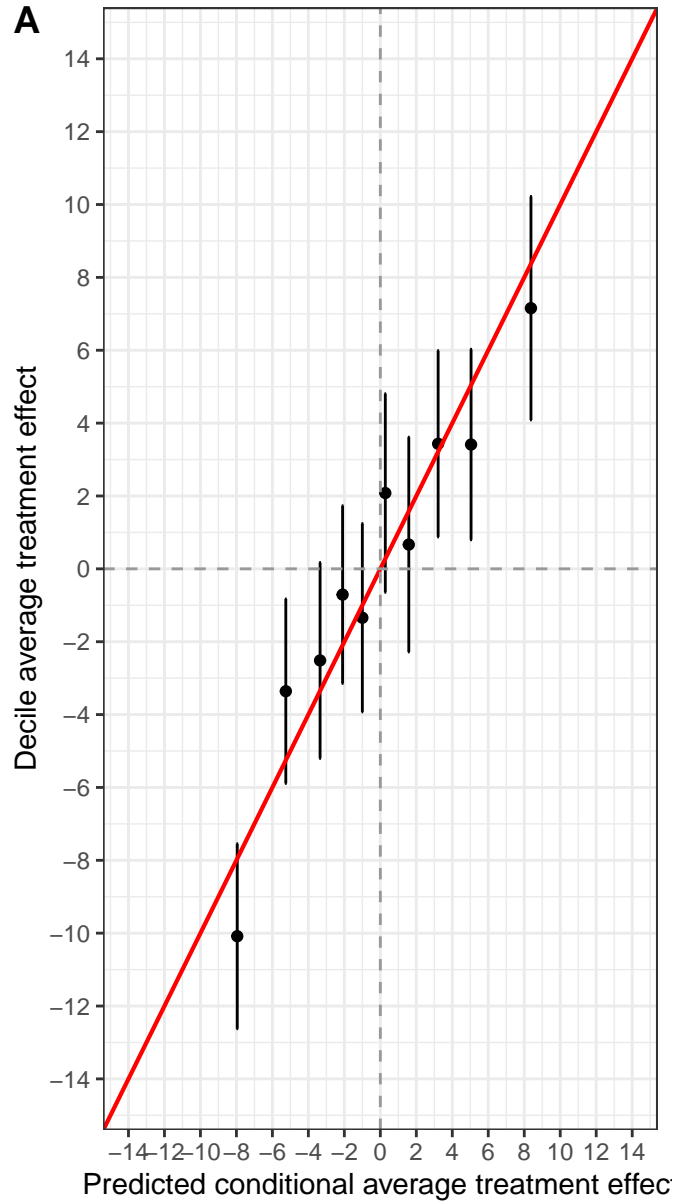
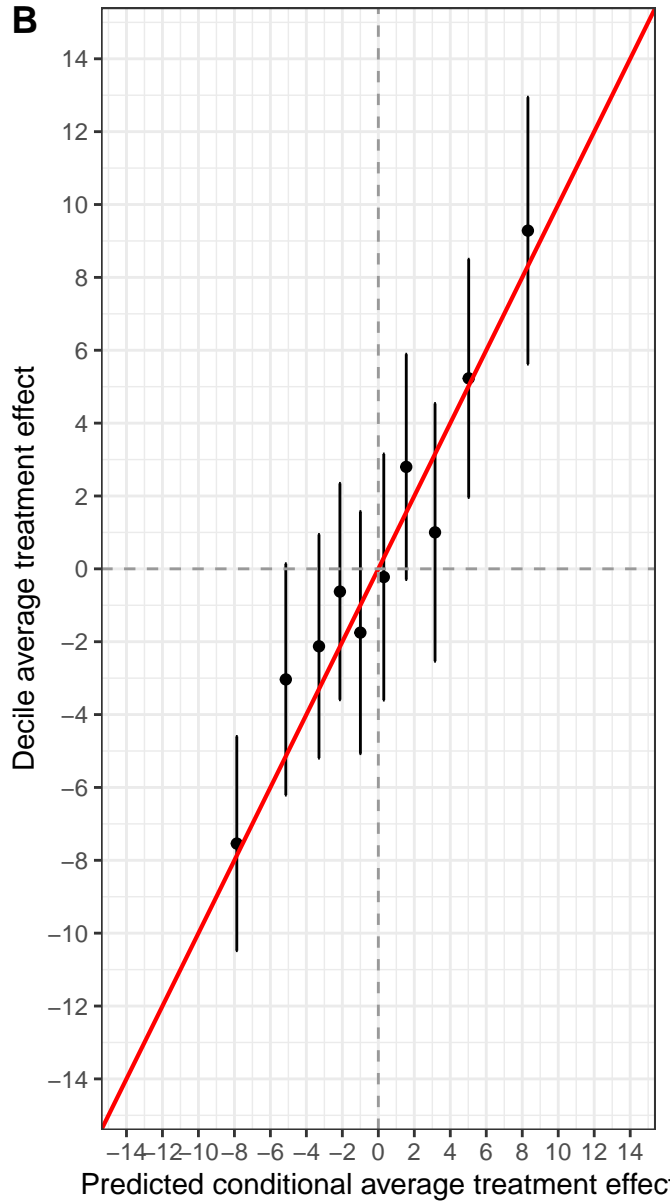
Effect submodels

A**B**

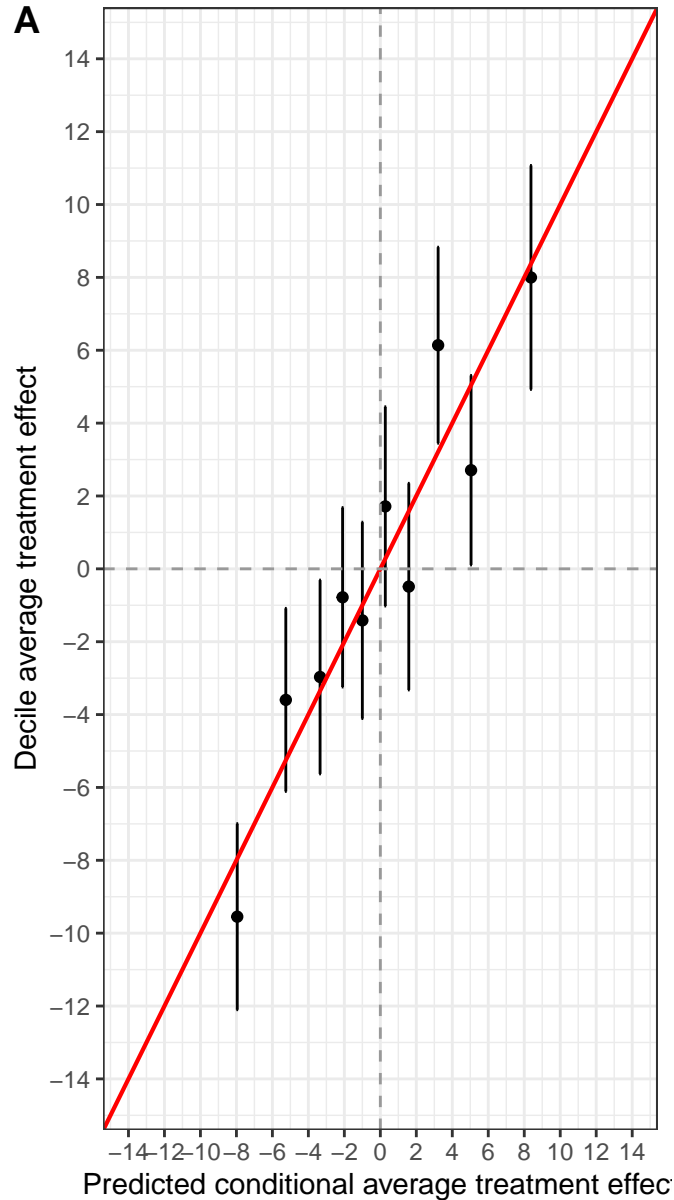
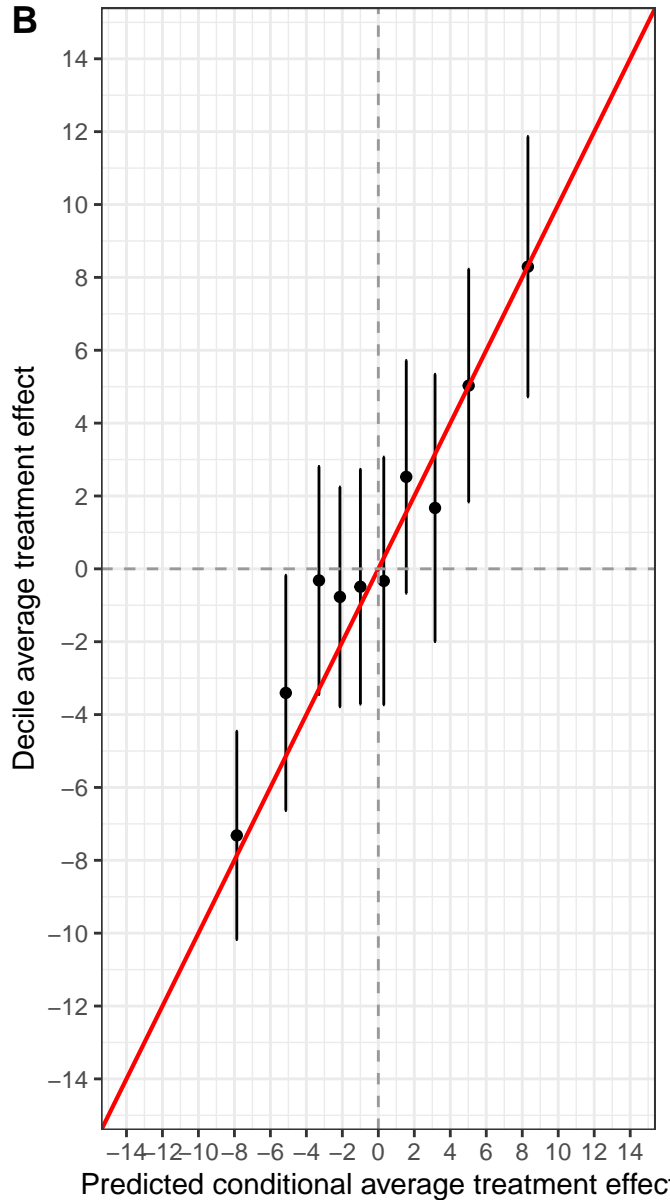
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

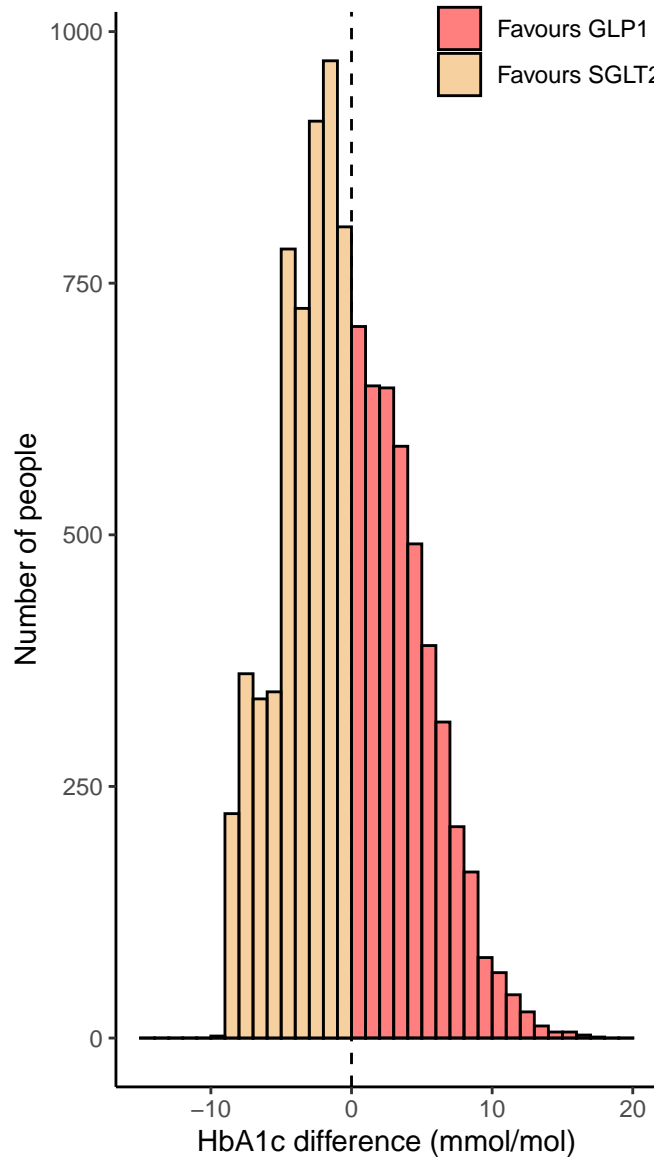
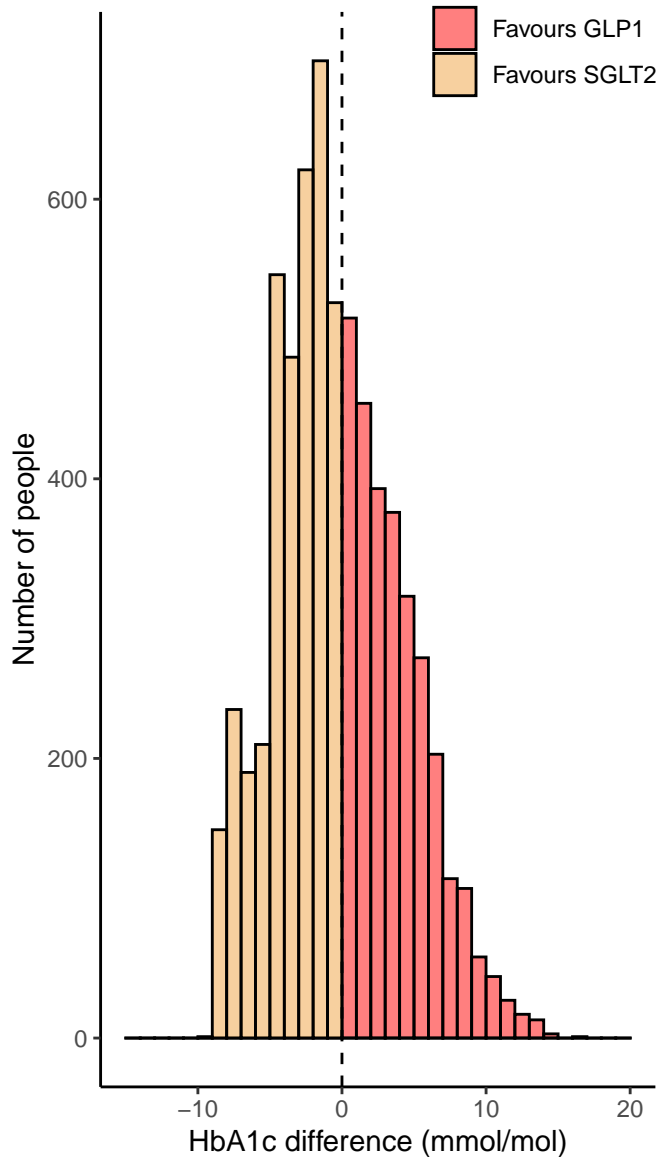
Effects validation prop score matching

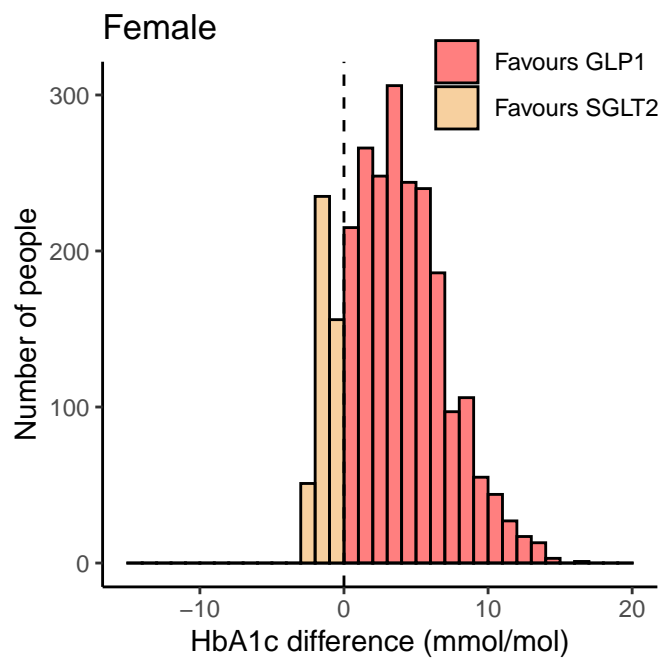
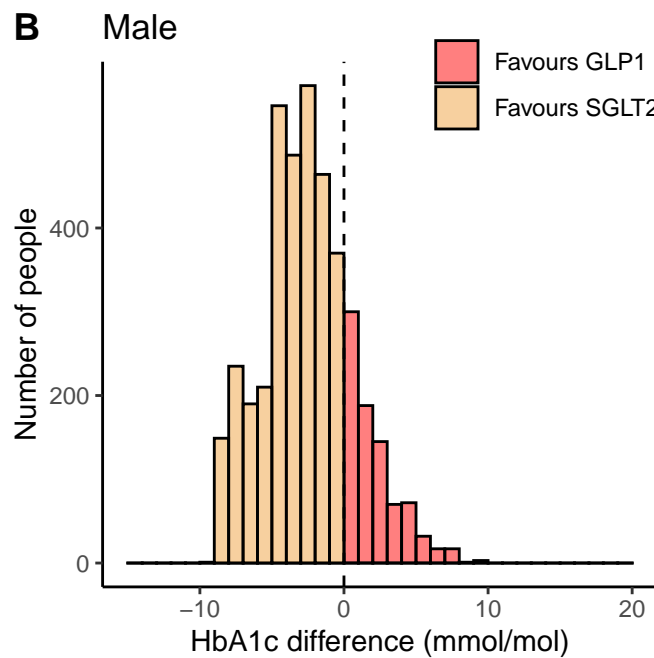
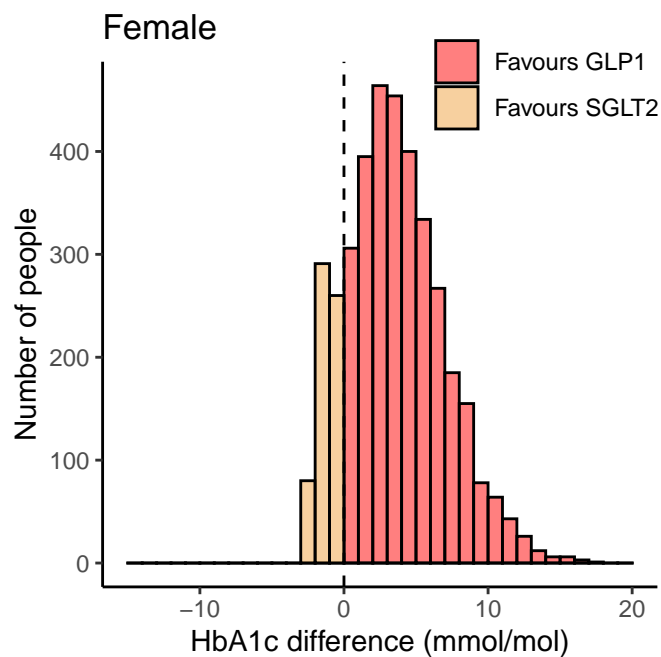
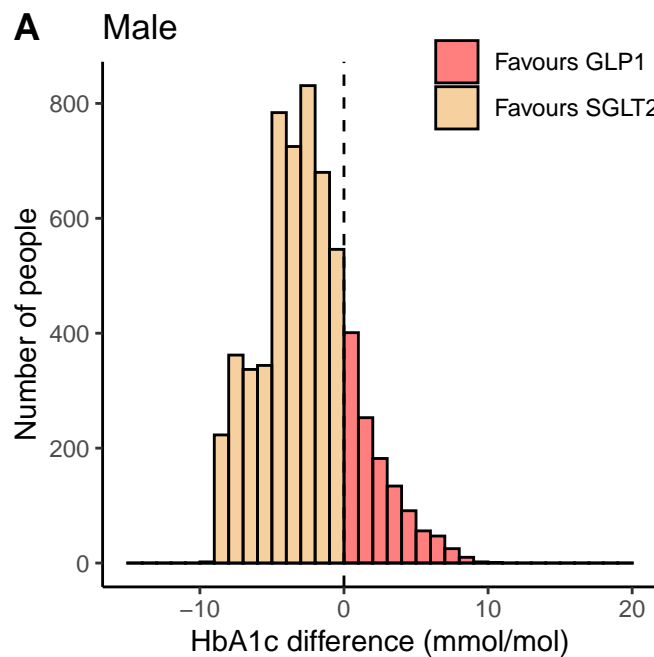
A**B**

Effects validation prop score inverse weighting

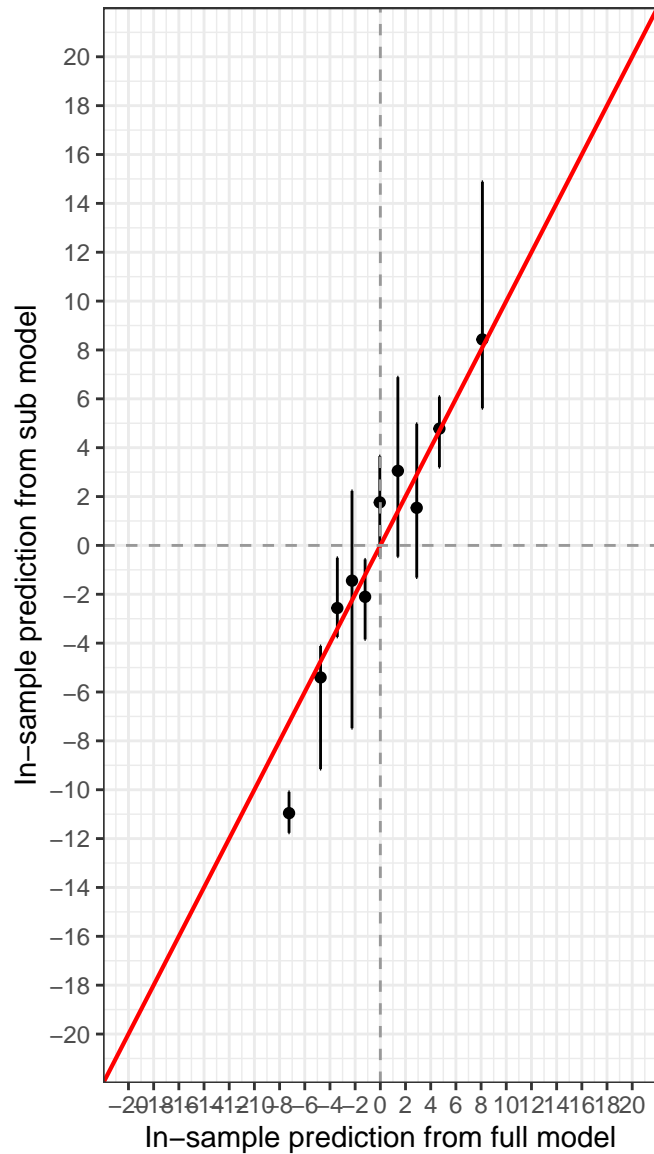
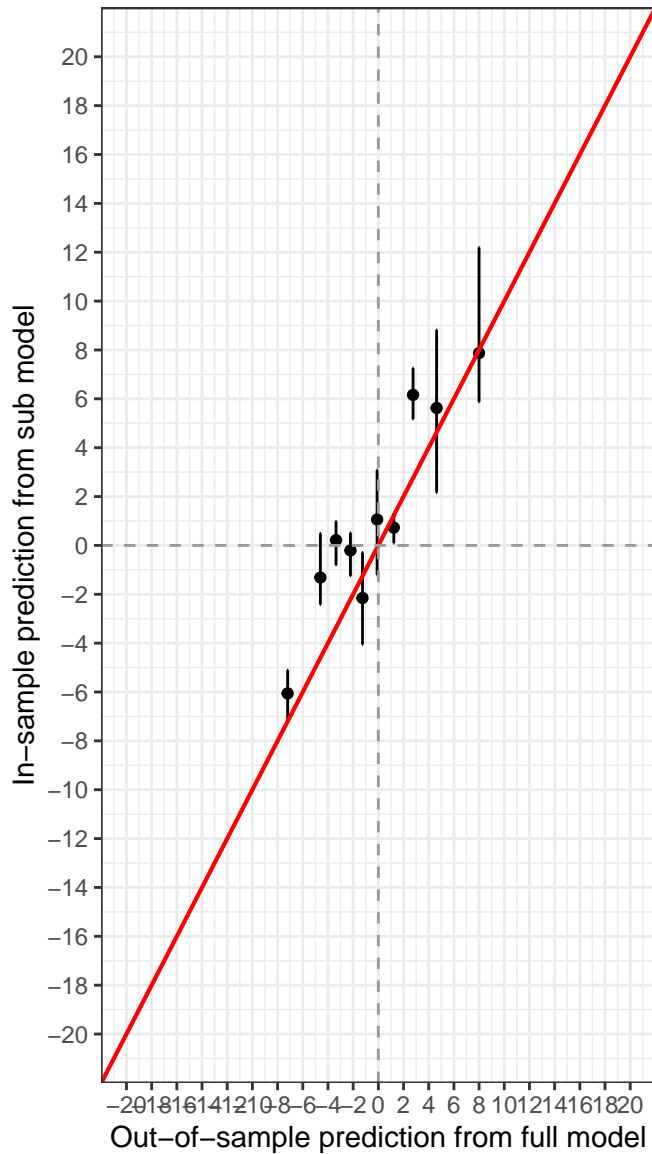
A**B**

Model fitting: Variable Selection 2, Incomplete data (Propensity score)

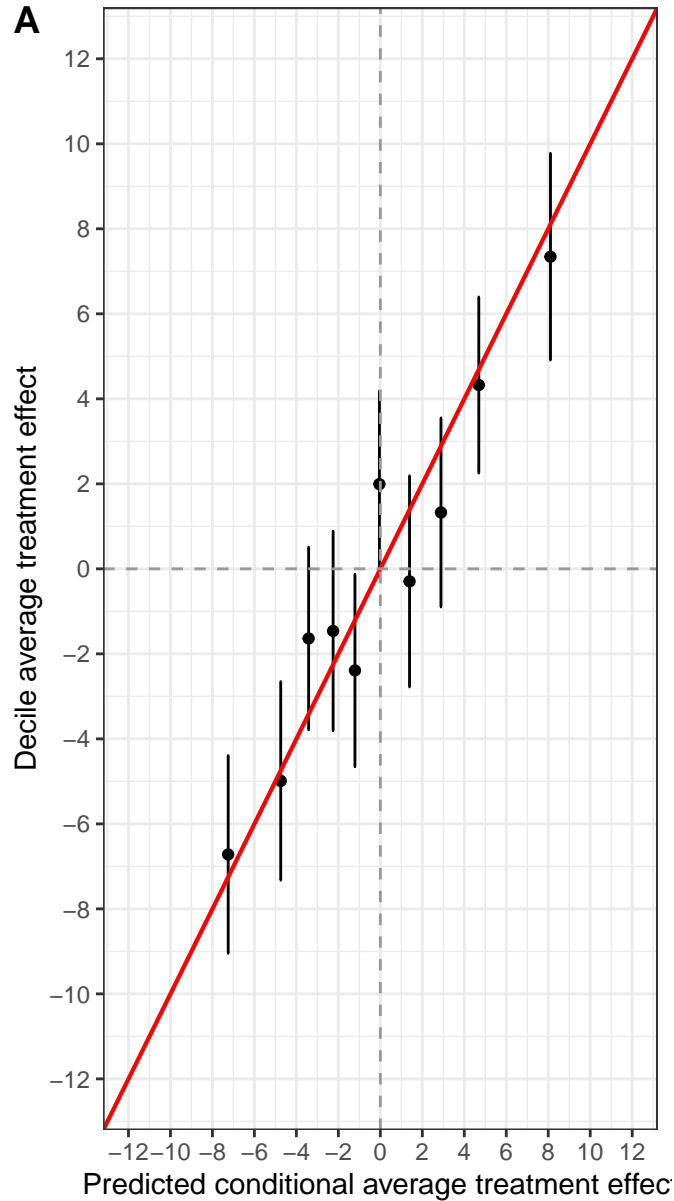
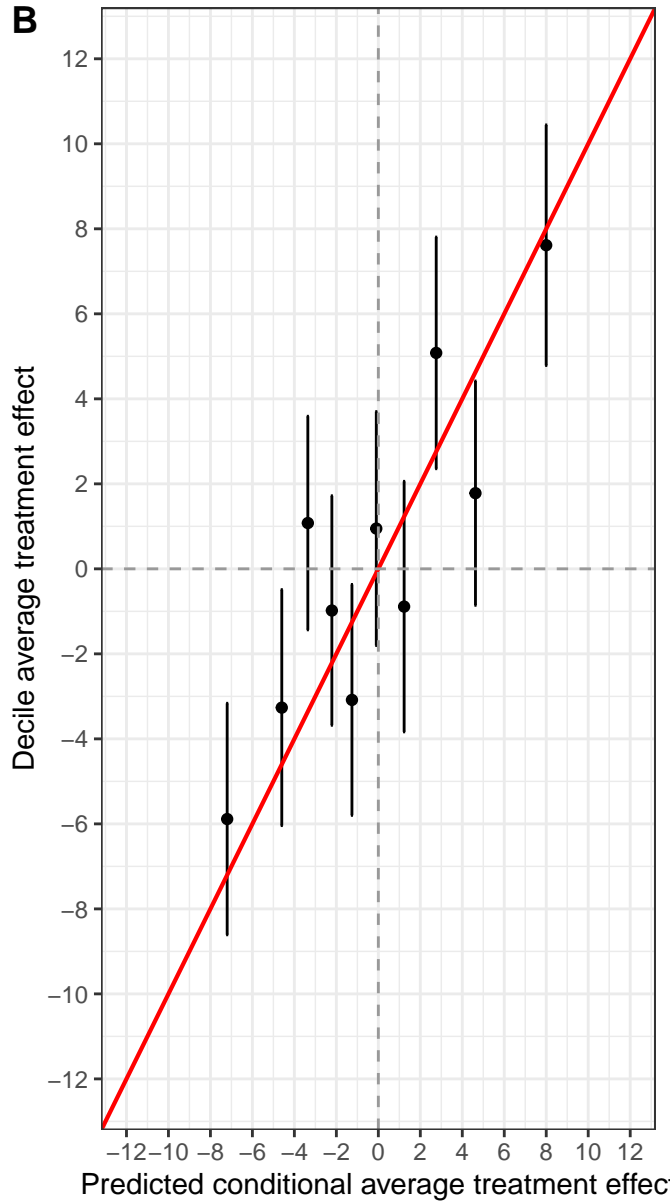
A**B**



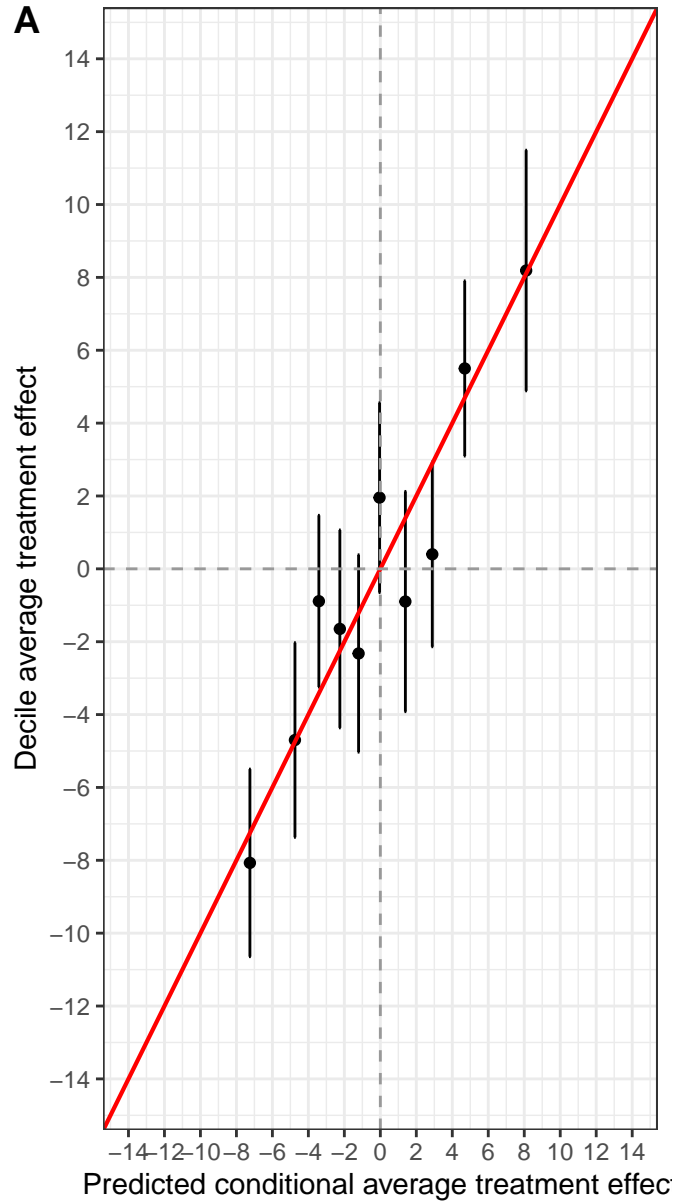
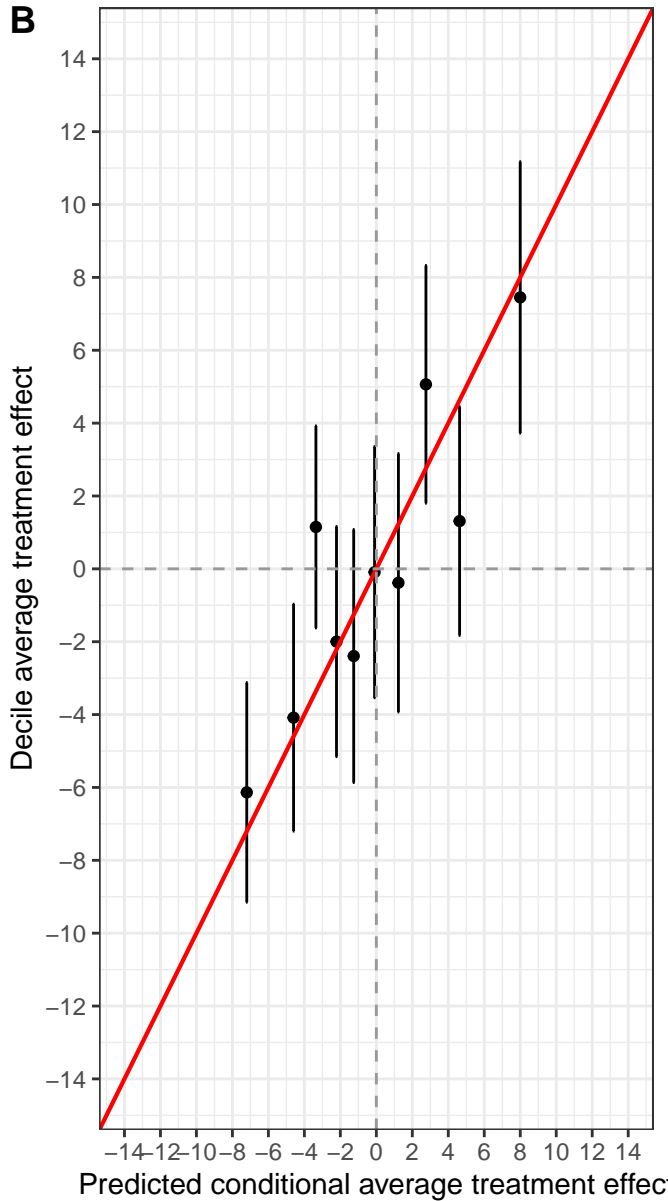
Effect submodels

A**B**

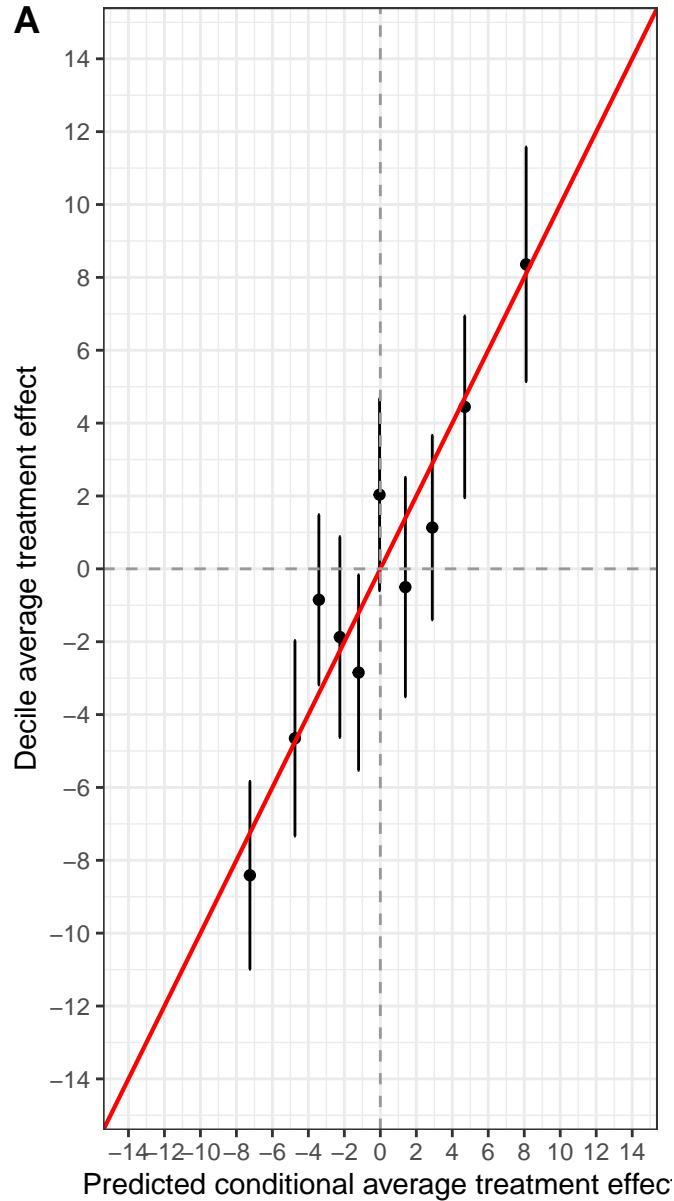
Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop_score})$

A**B**

Effects validation prop score matching

A**B**

Effects validation prop score inverse weighting

A**B**