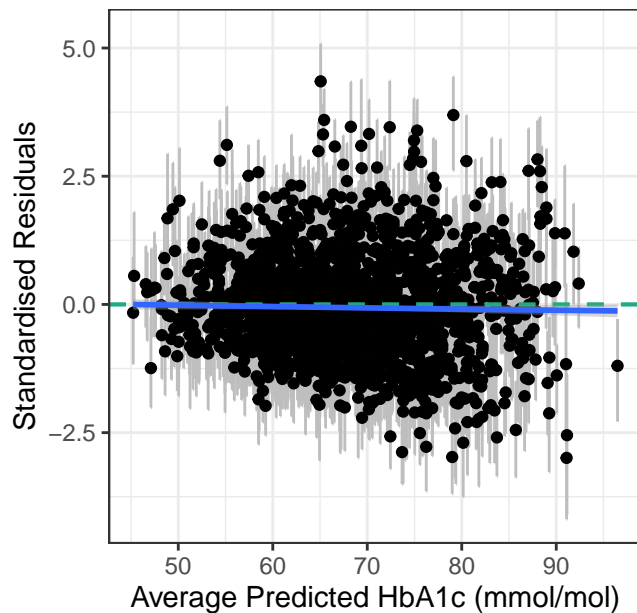
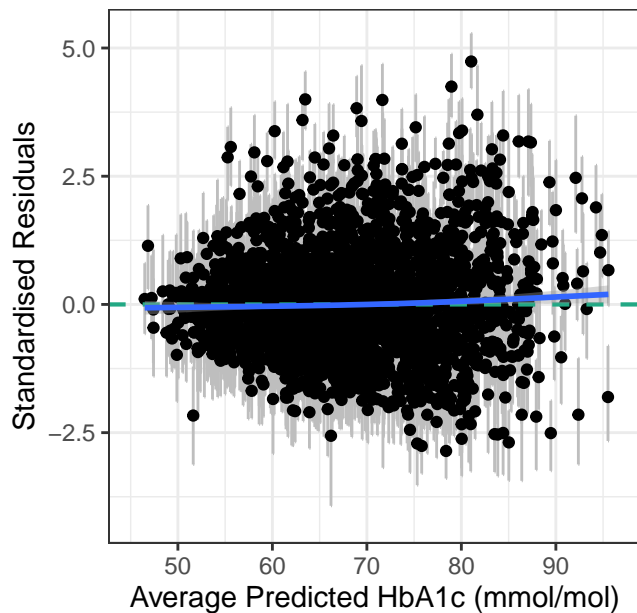
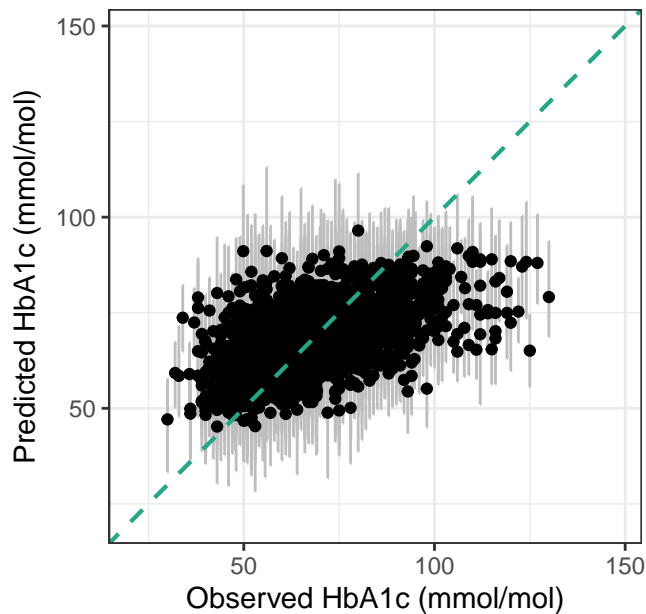
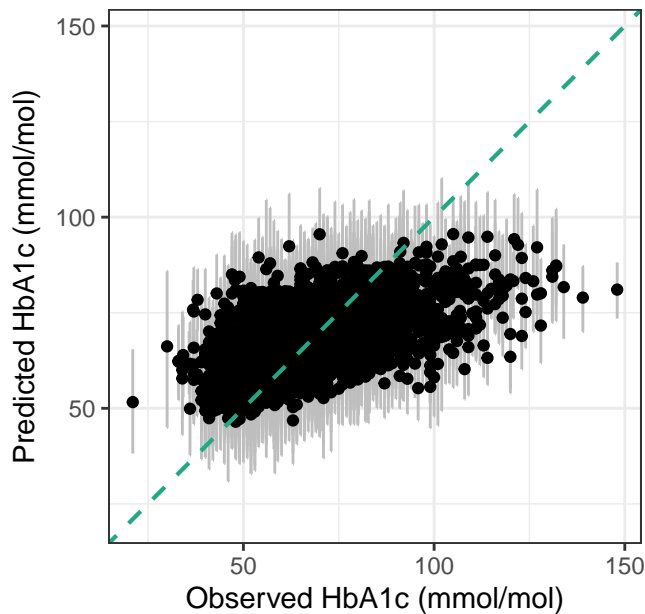


## Residuals of Model 6



Model

R2 (bigger is better)

0.23

0.24

0.25

0.26

0.27

0.28

Model

RMSE (smaller is better)

13.6

13.8

14.0

14.2

Model

RSS (smaller is better)

350000

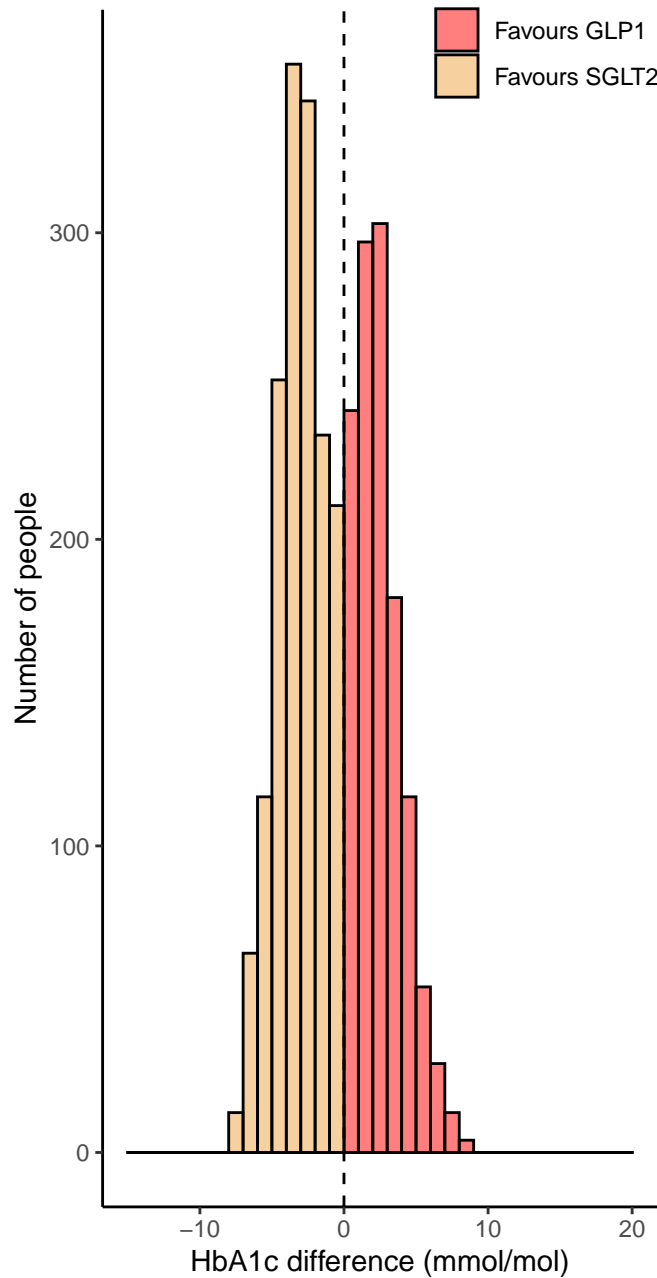
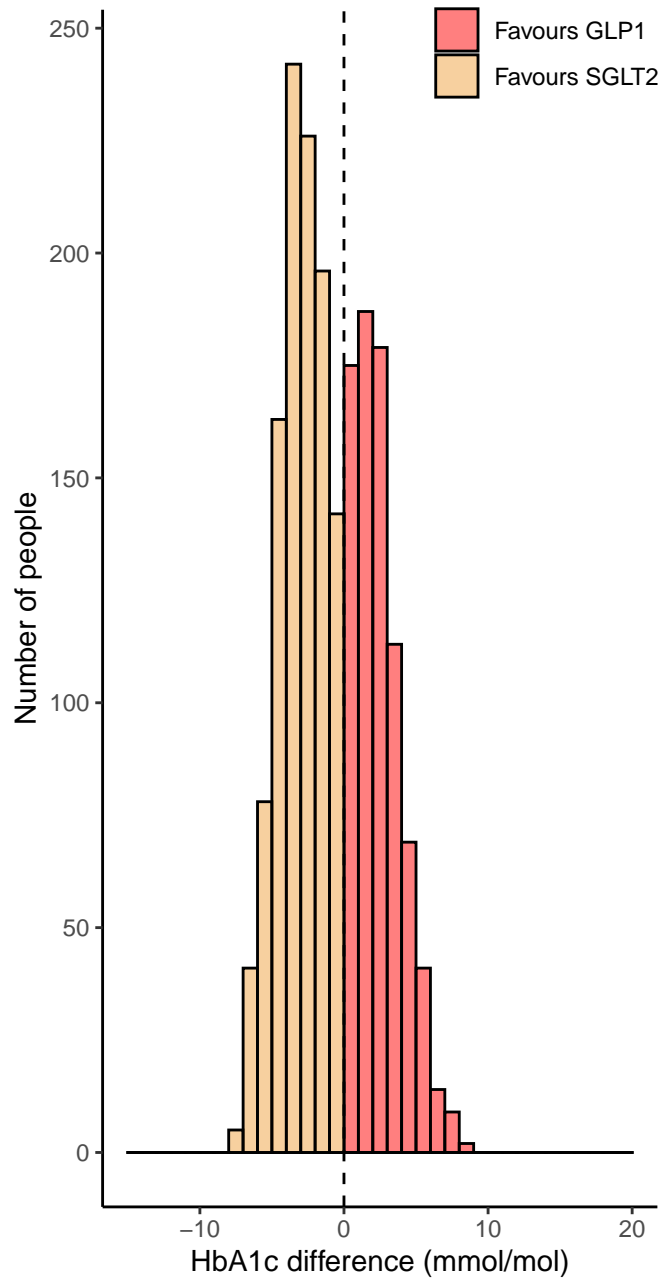
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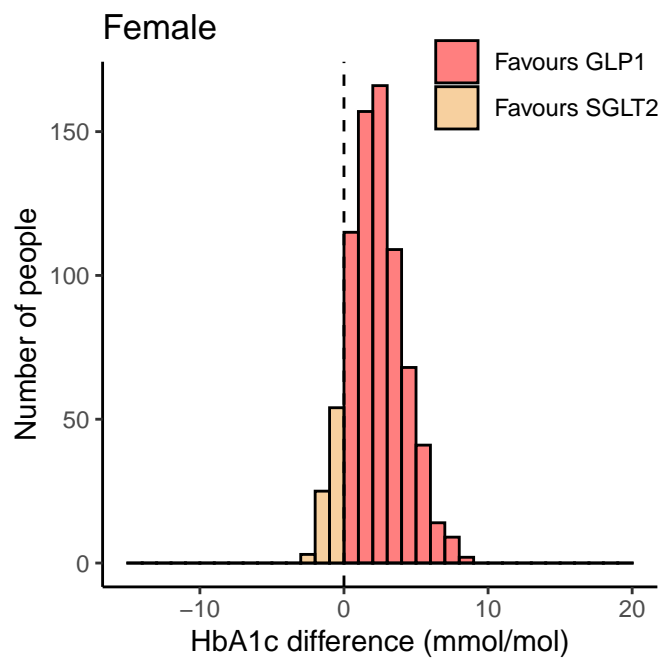
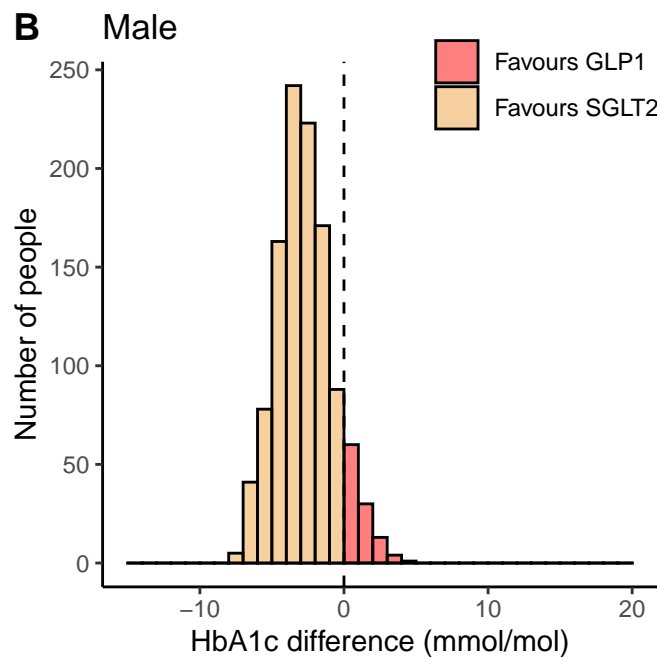
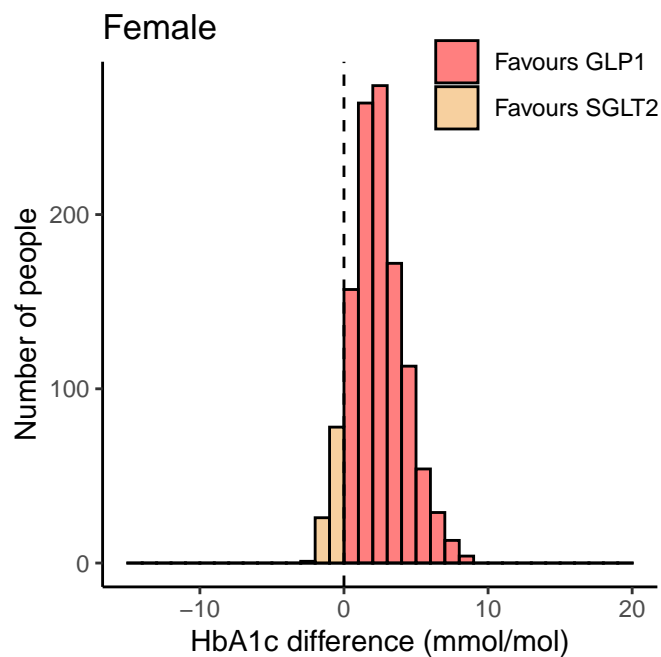
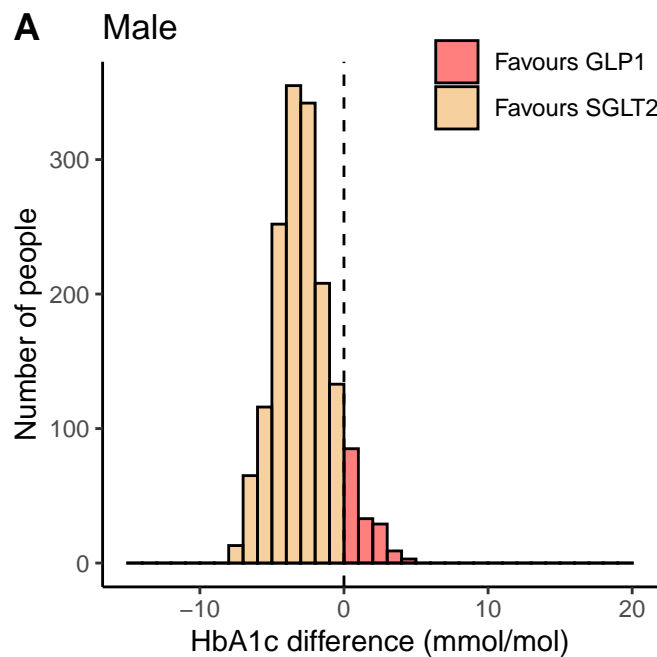
450000

500000

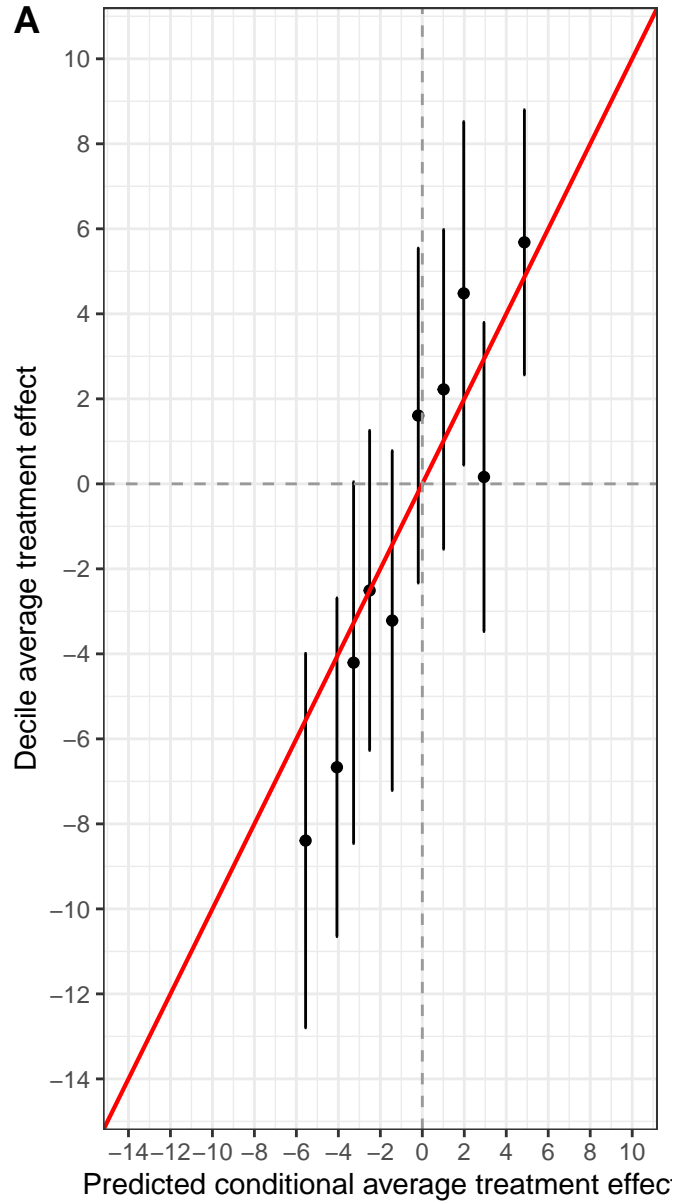
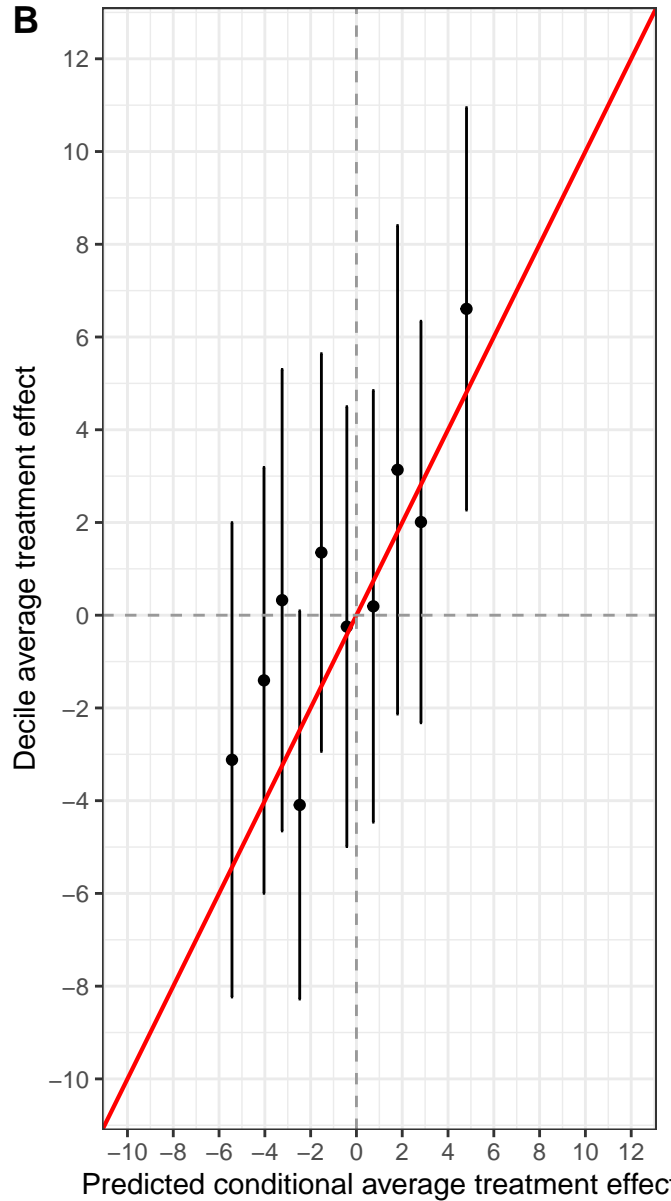
550000

Dataset ● Development ▲ Validation

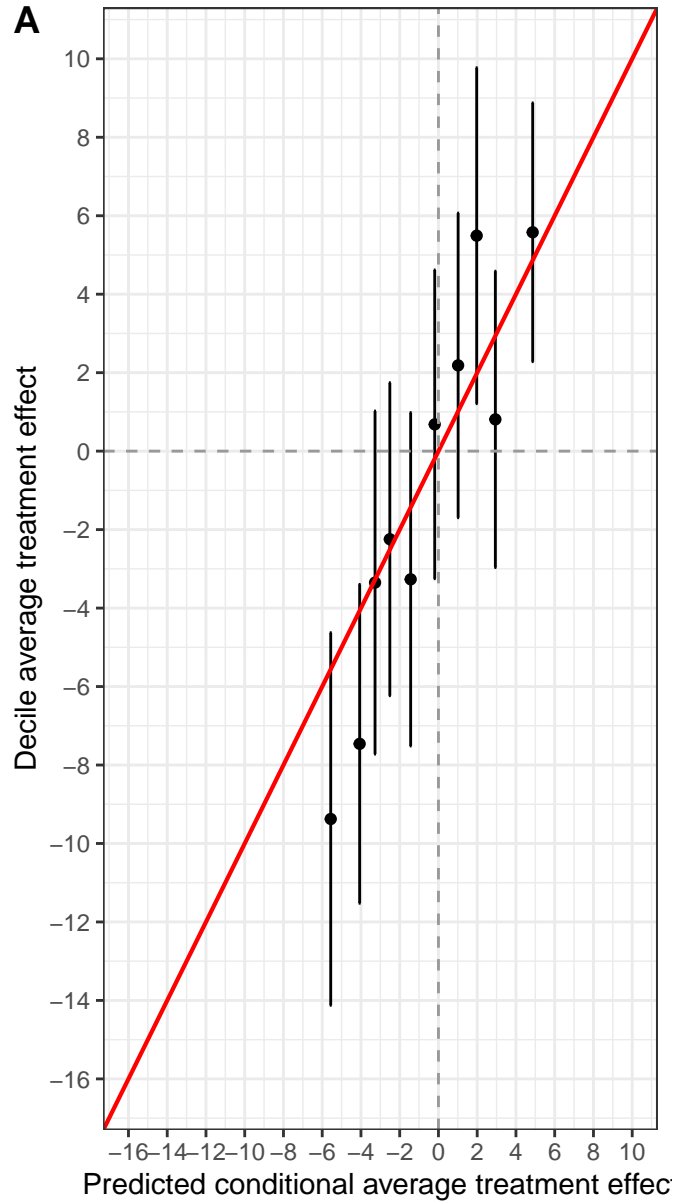
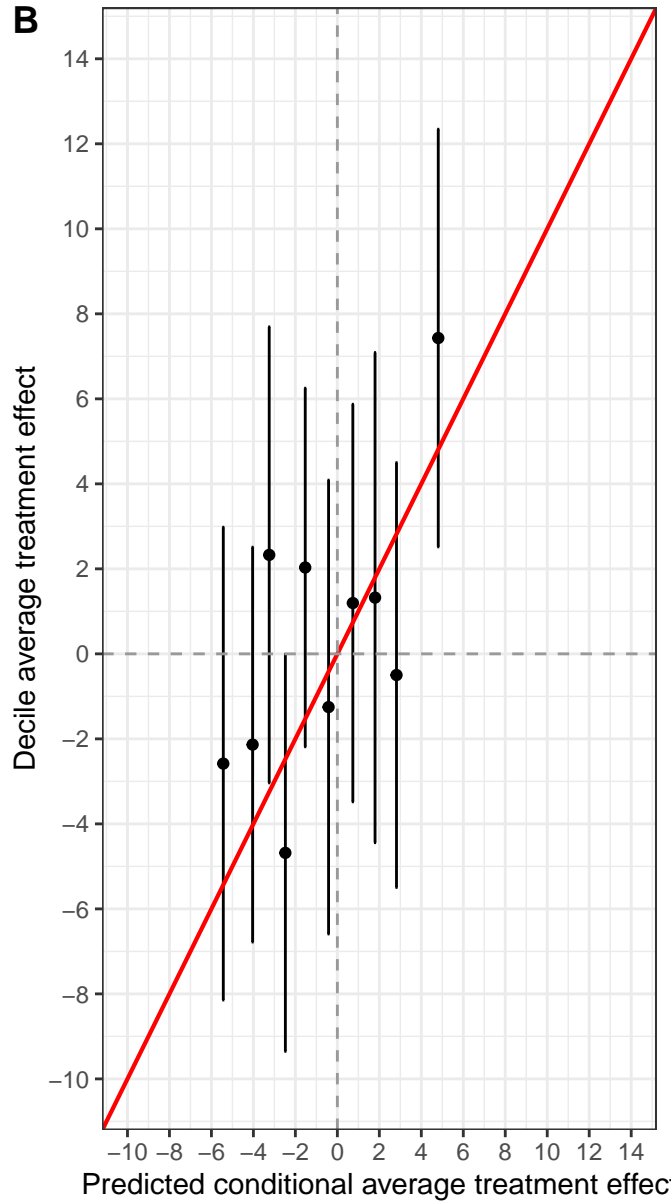
**A****B**



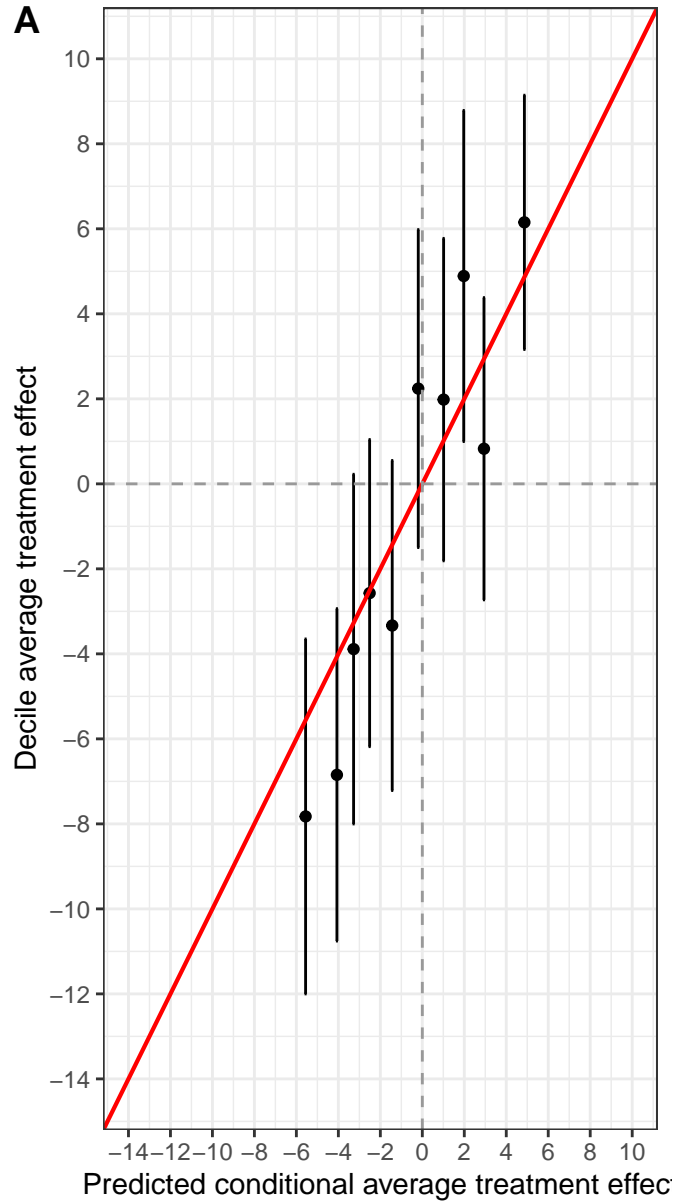
# Effects validation $\text{lm}(\text{hba1c} \sim \text{drugclass} + \text{prop\_score})$

**A****B**

# Effects validation prop score matching

**A****B**

# Effects validation prop score inverse weighting

**A****B**