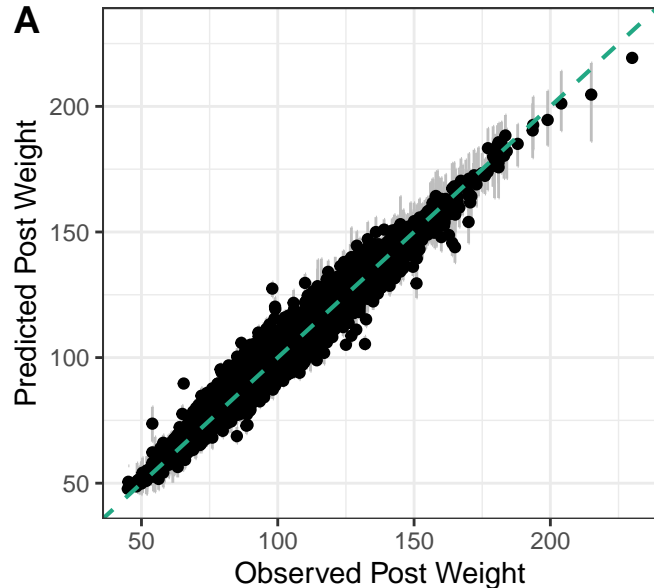
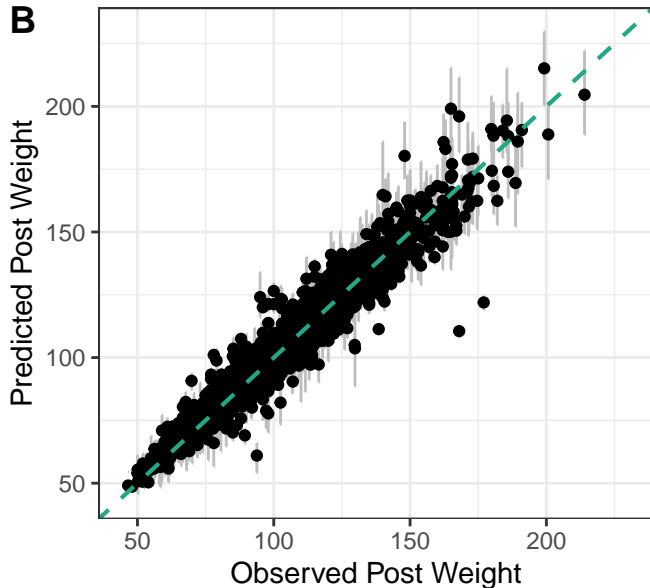


Residuals of Model 4

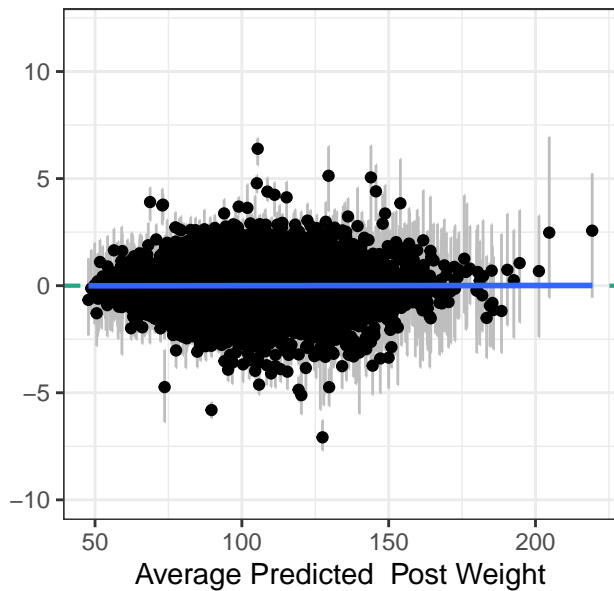
A



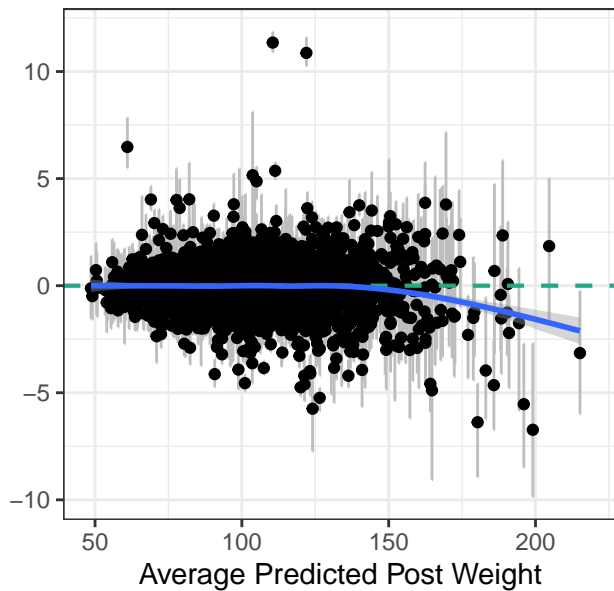
B

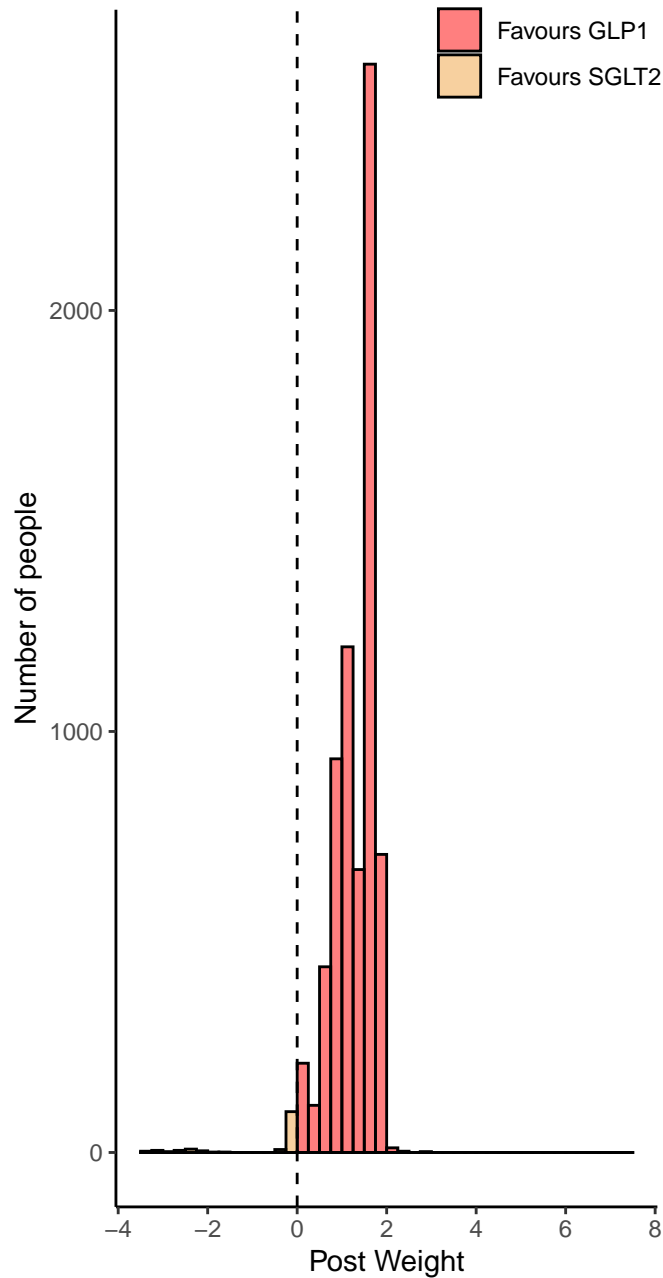
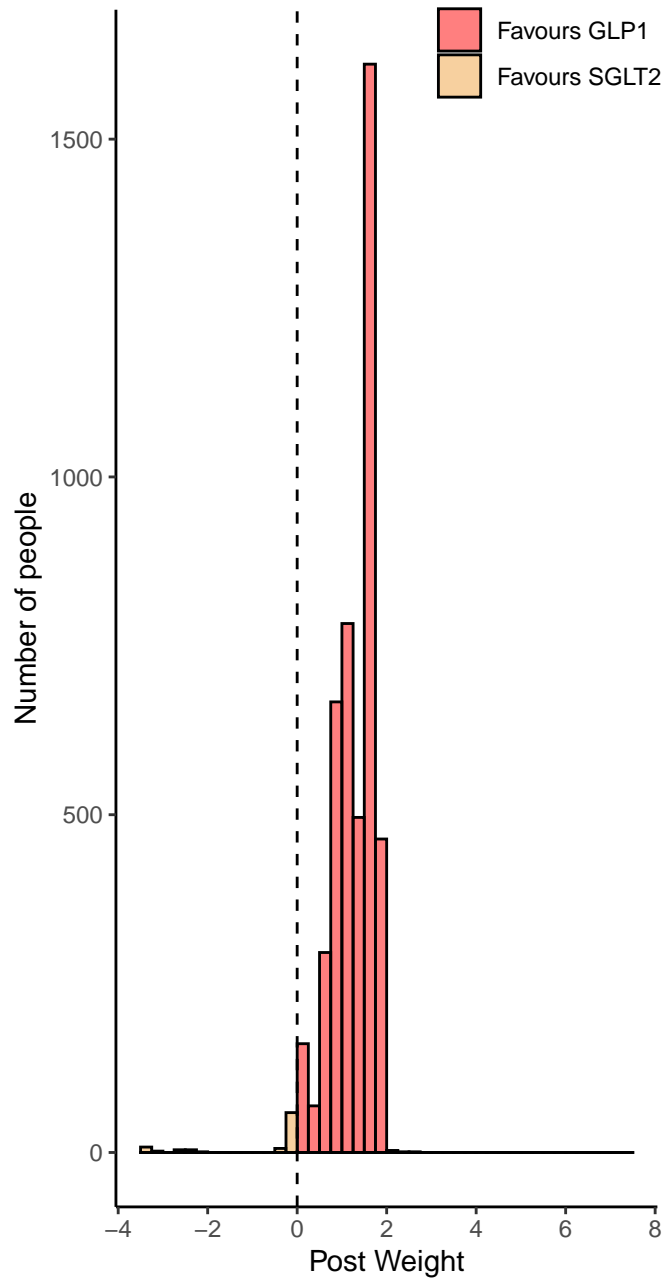


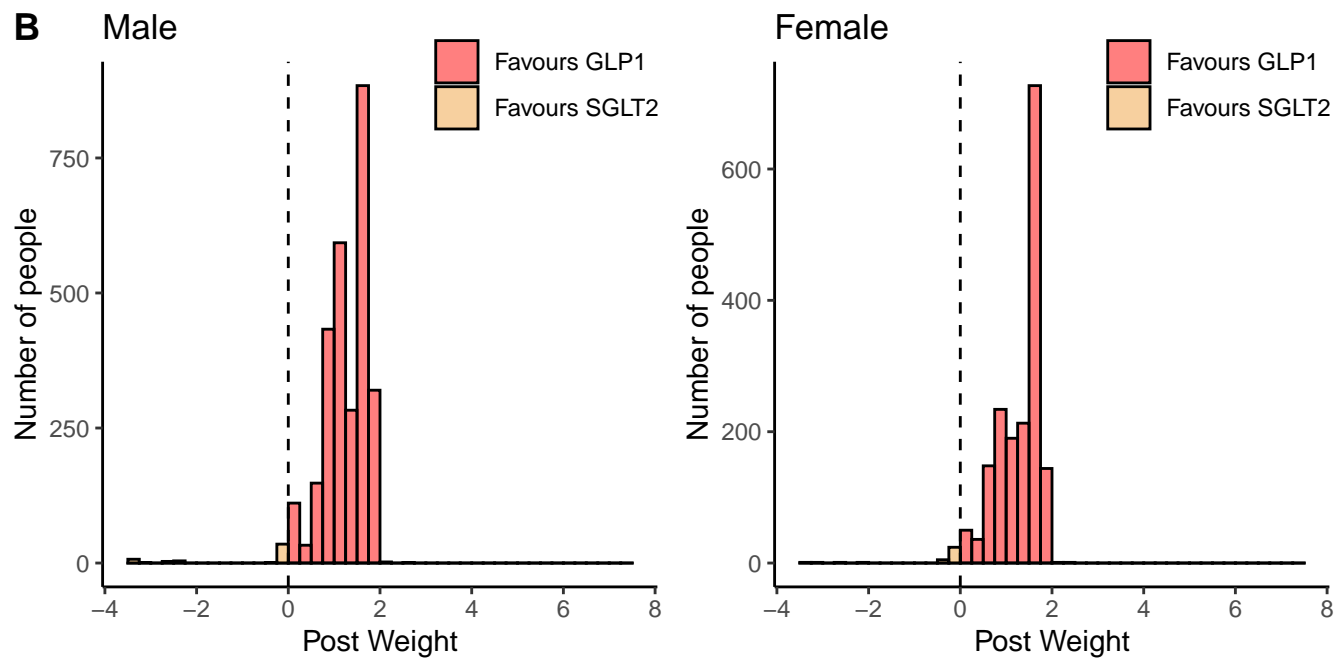
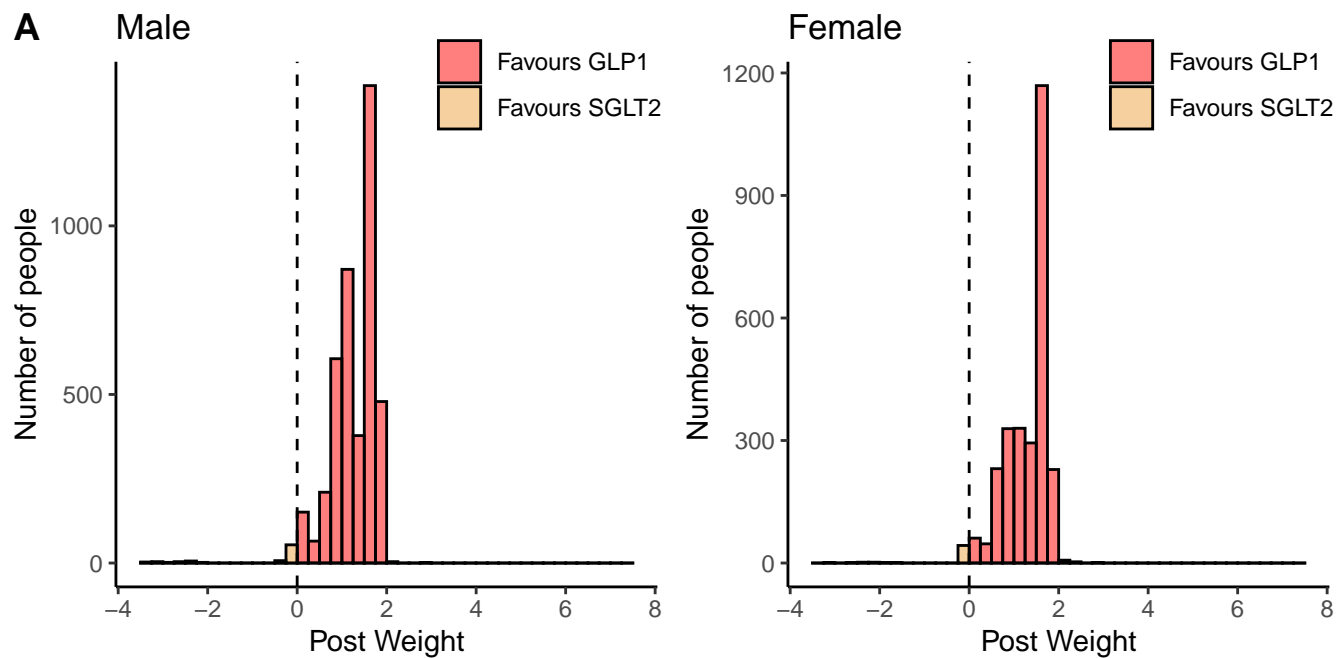
Standardised Residuals



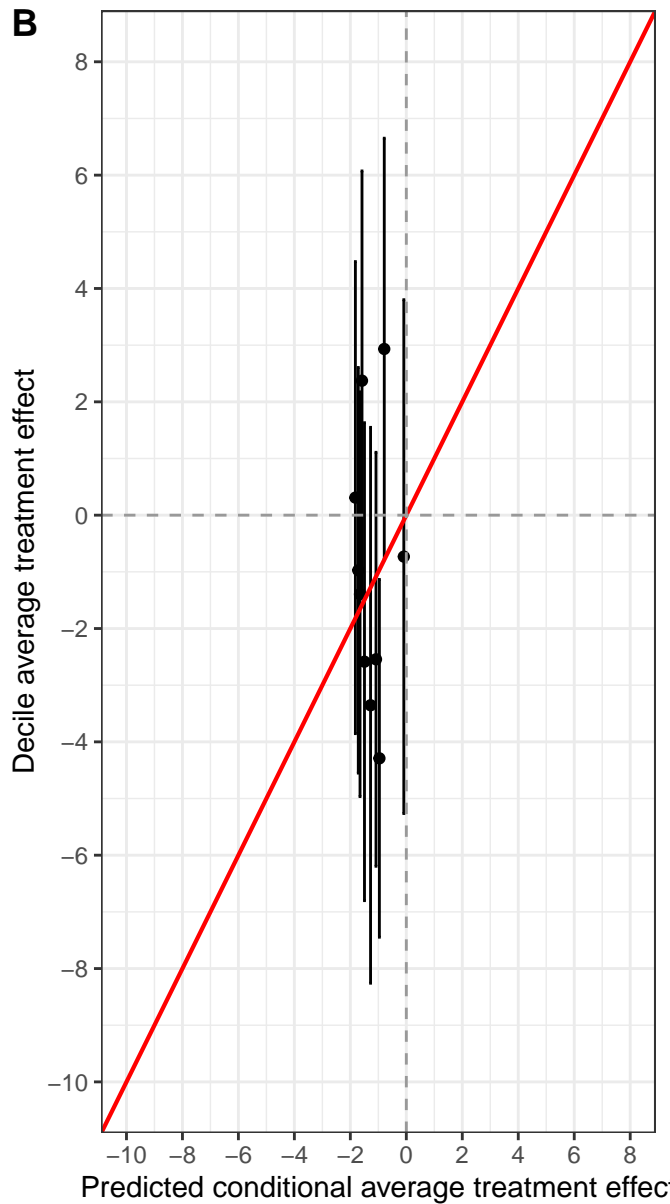
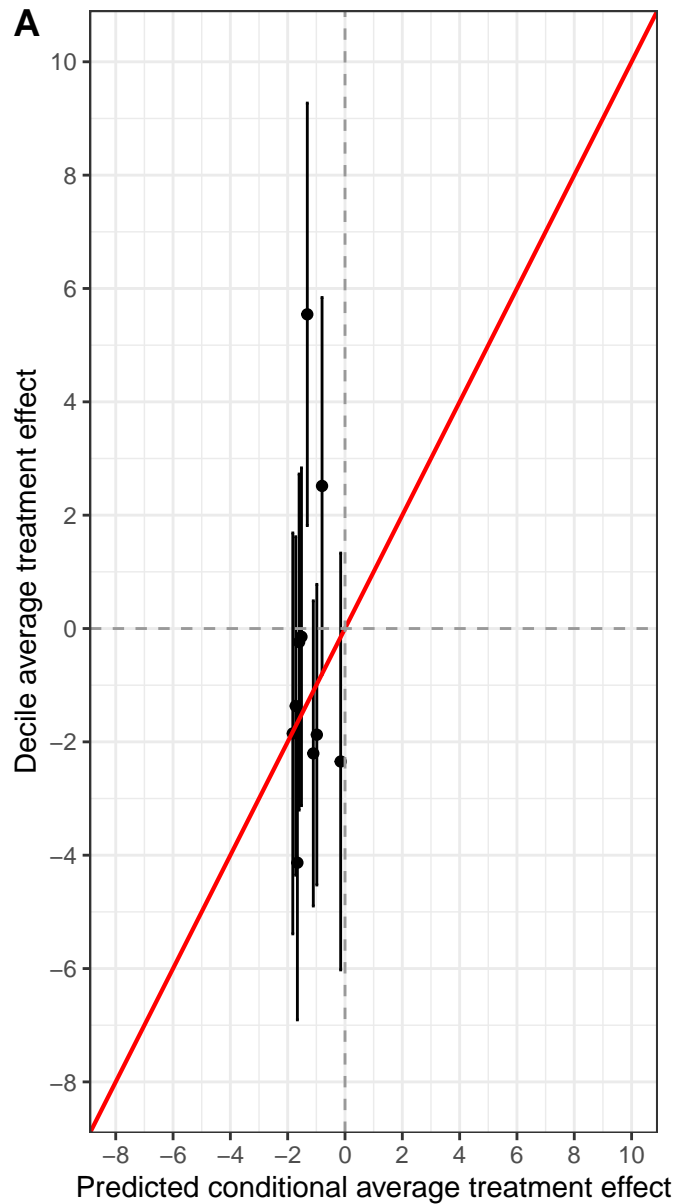
Standardised Residuals



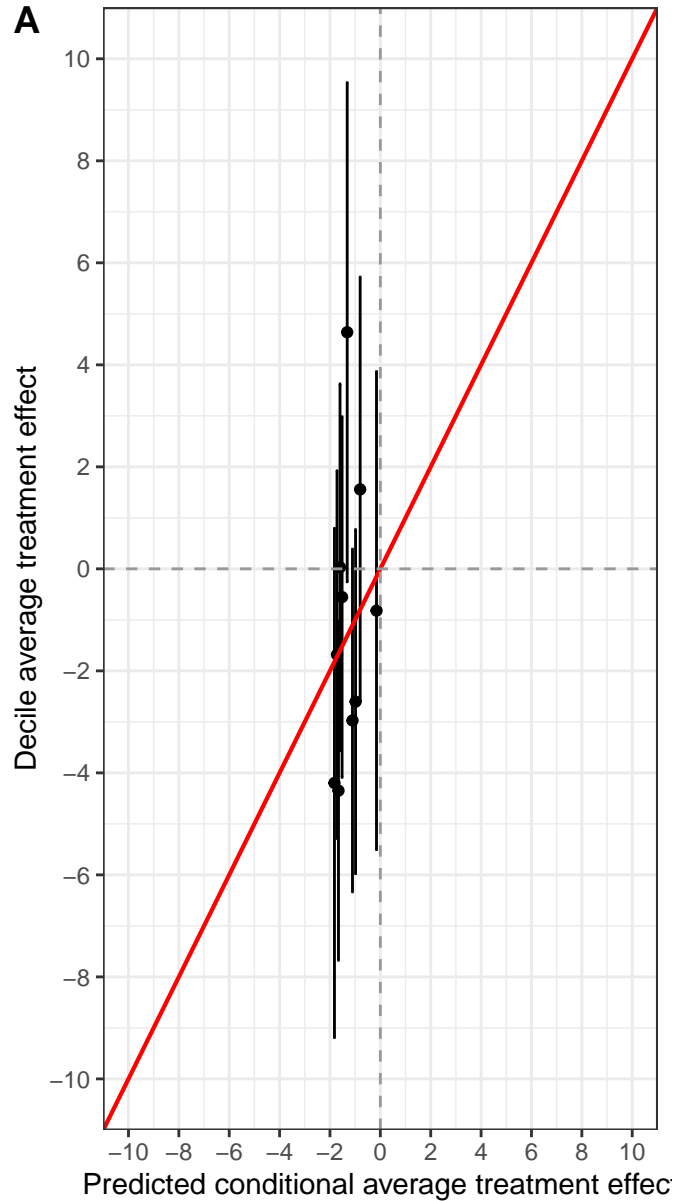
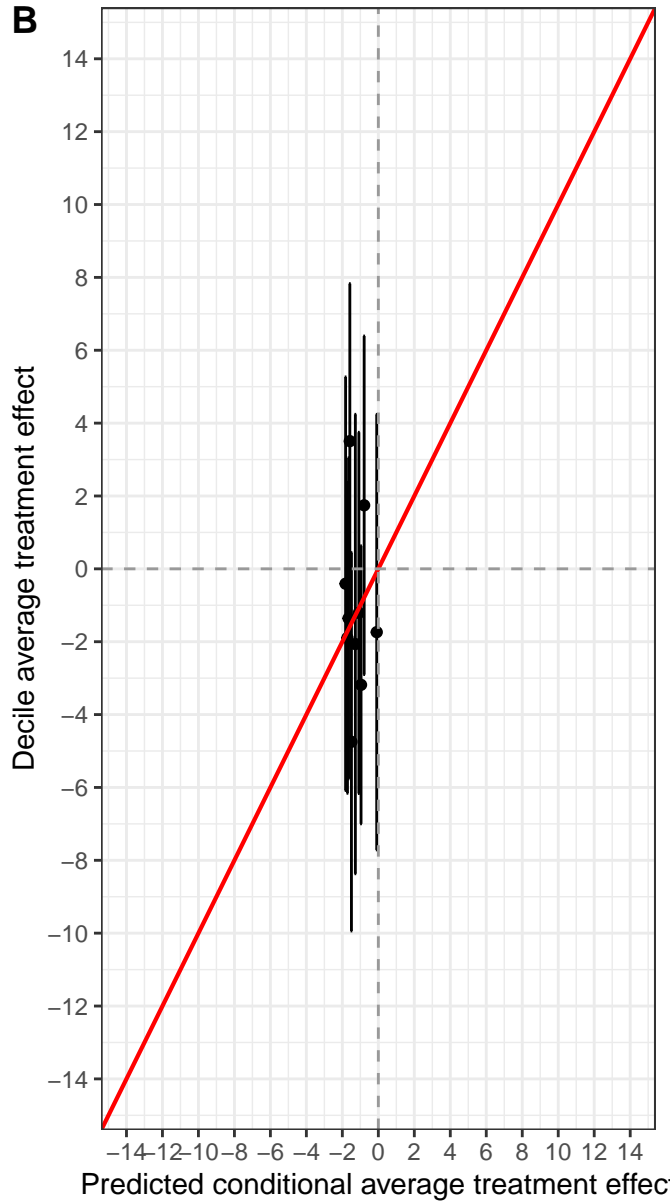
A**B**



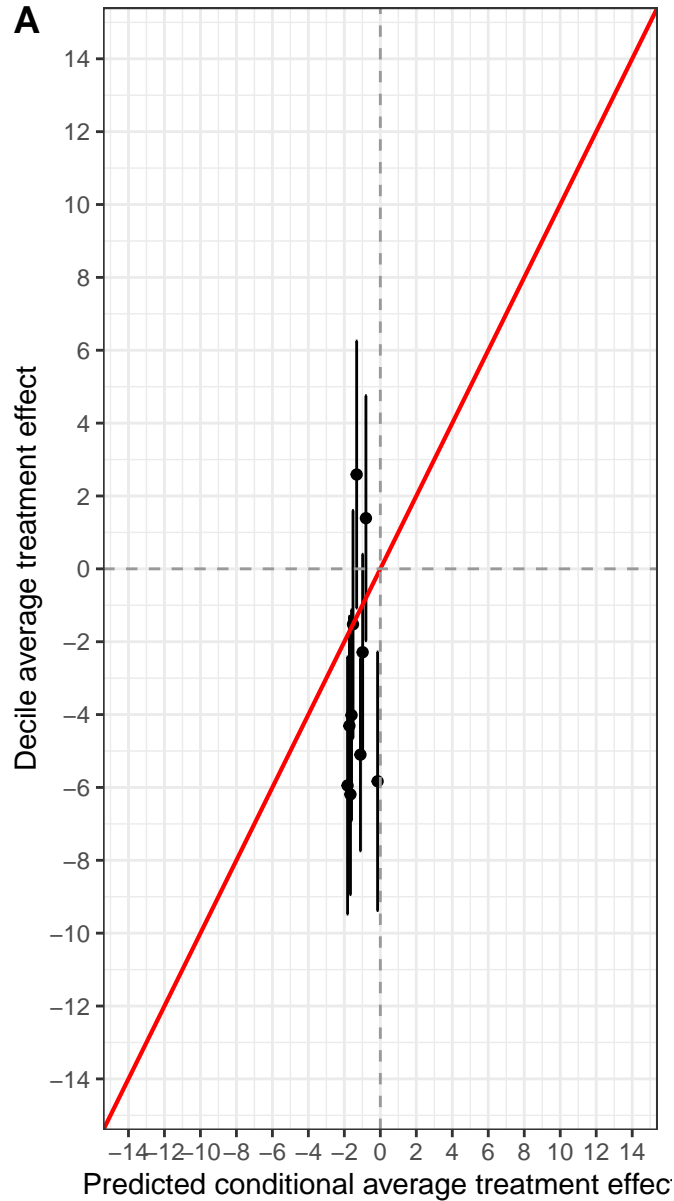
Effects validation $\text{lm}(\text{weight} \sim \text{drugclass} + \text{prop_score})$



Effects validation prop score matching

A**B**

Effects validation prop score inverse weighting

A**B**