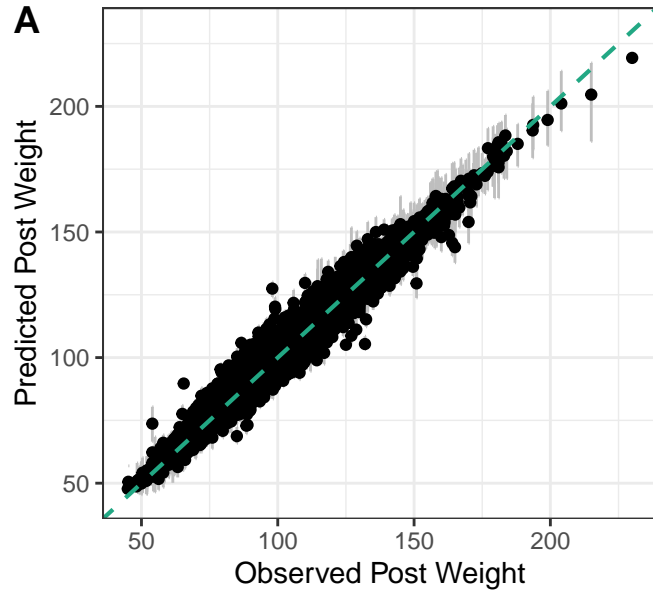
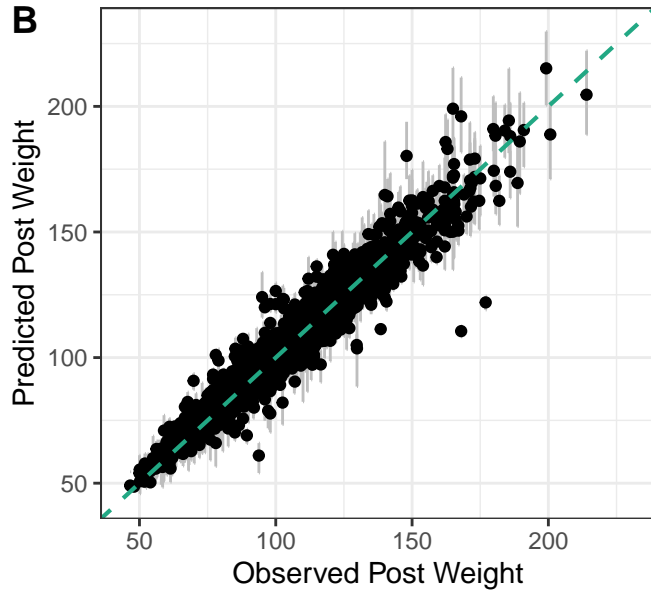
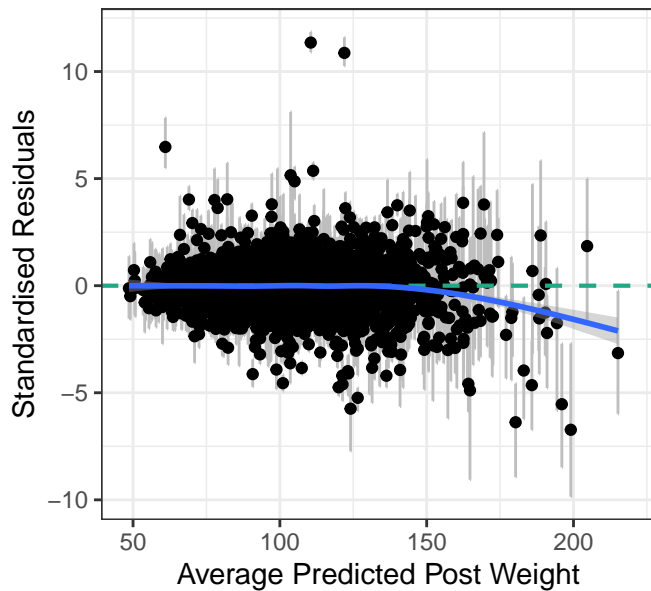
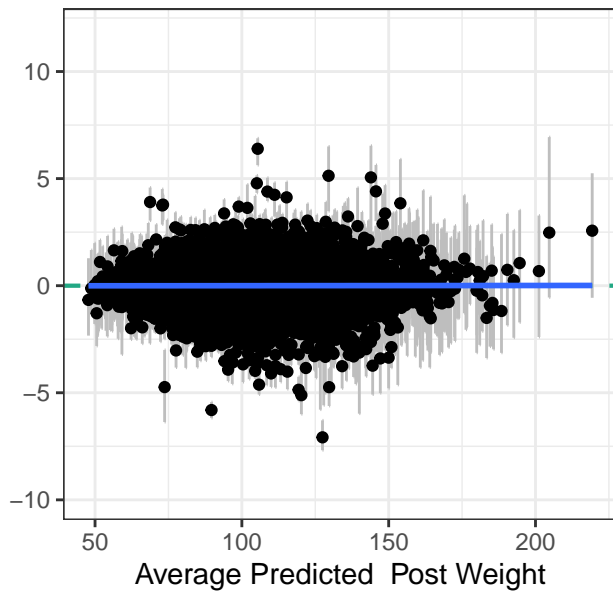
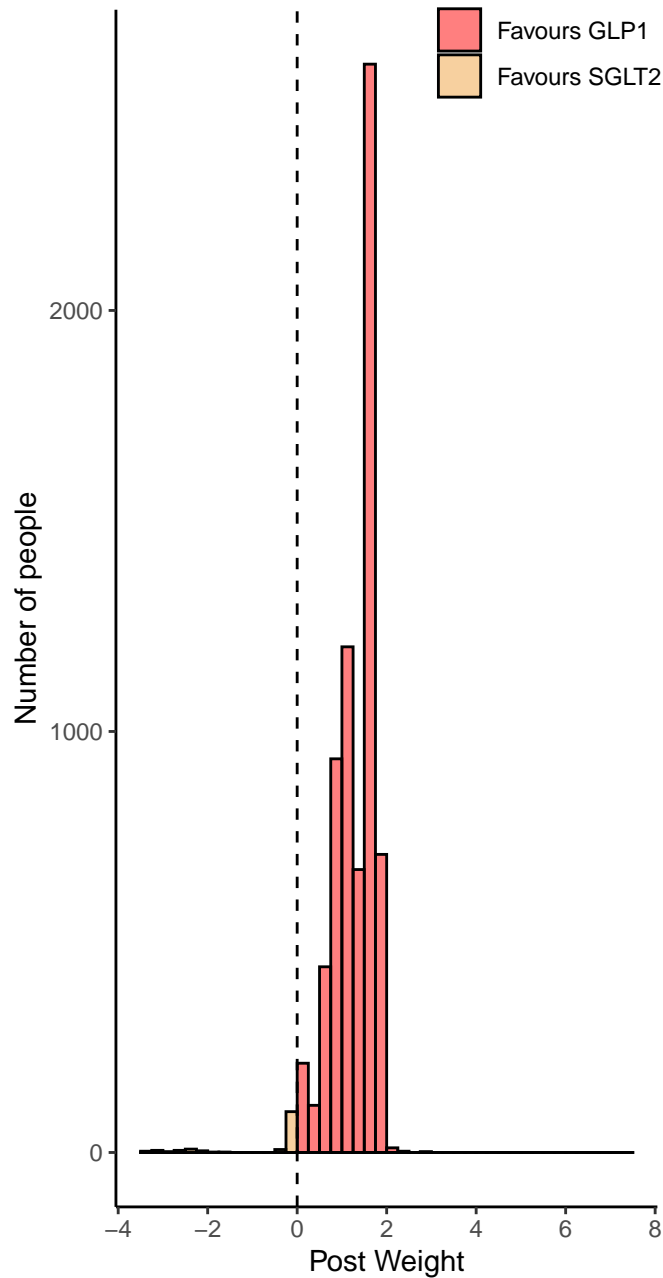
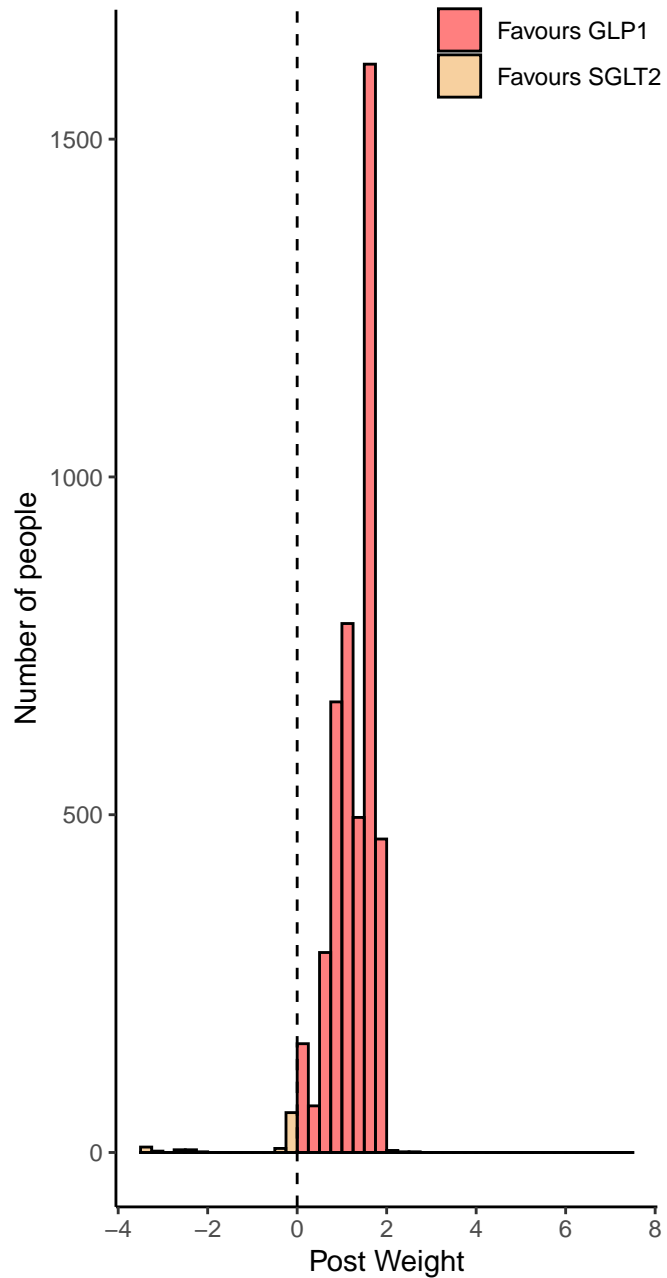
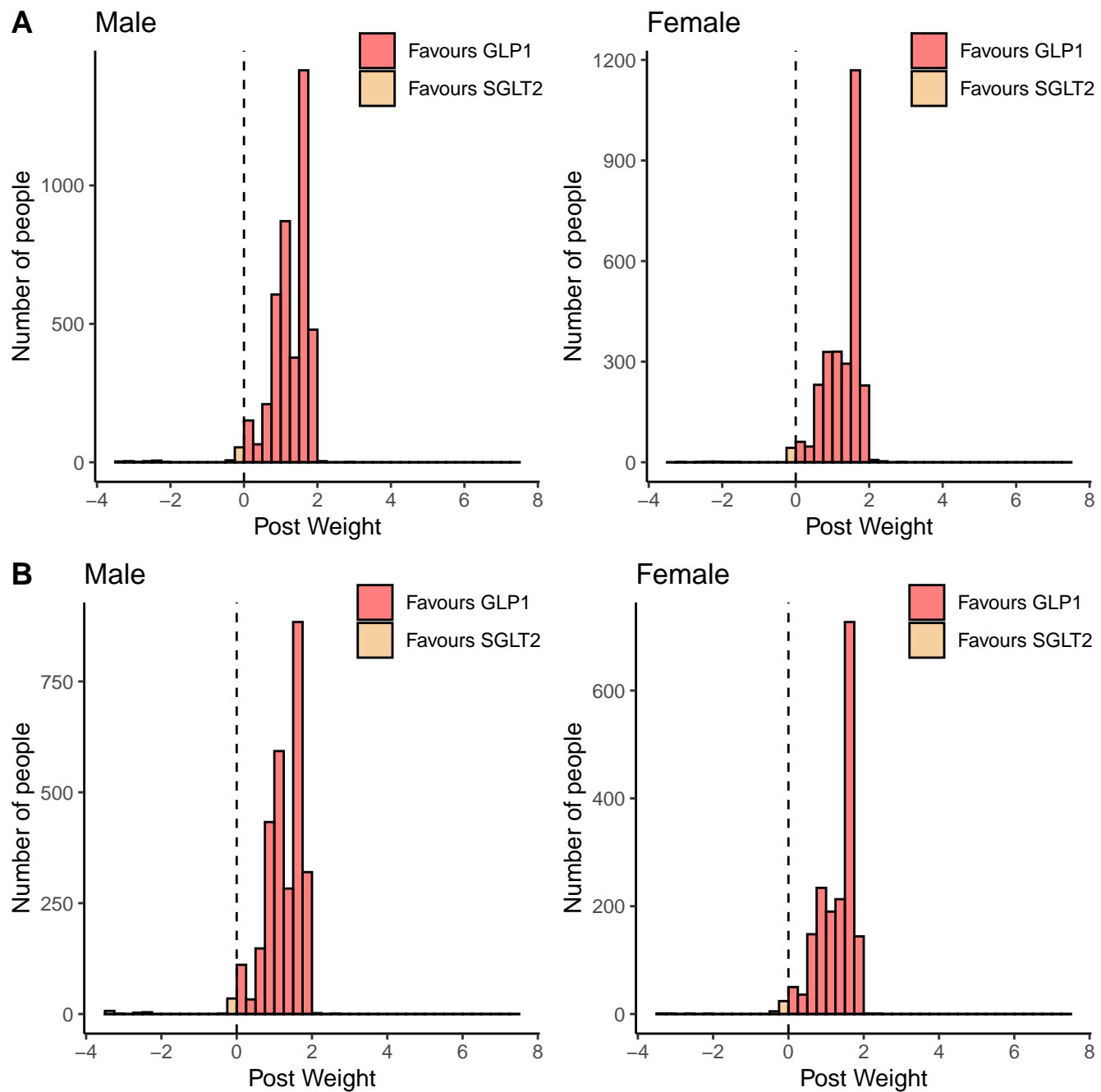


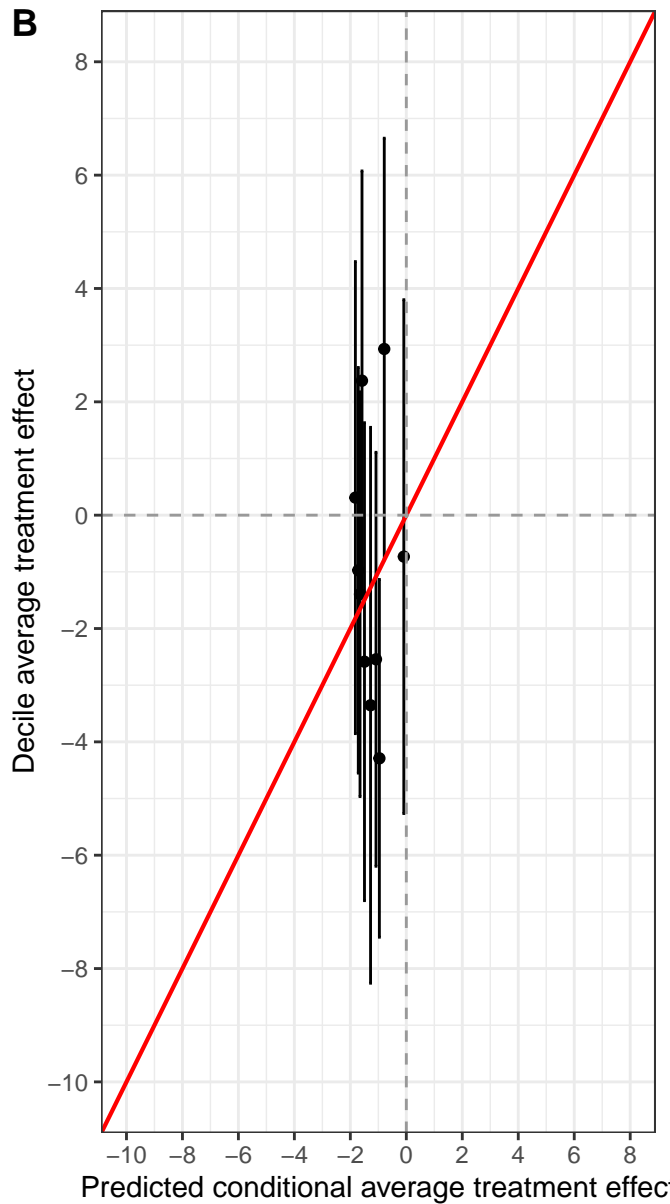
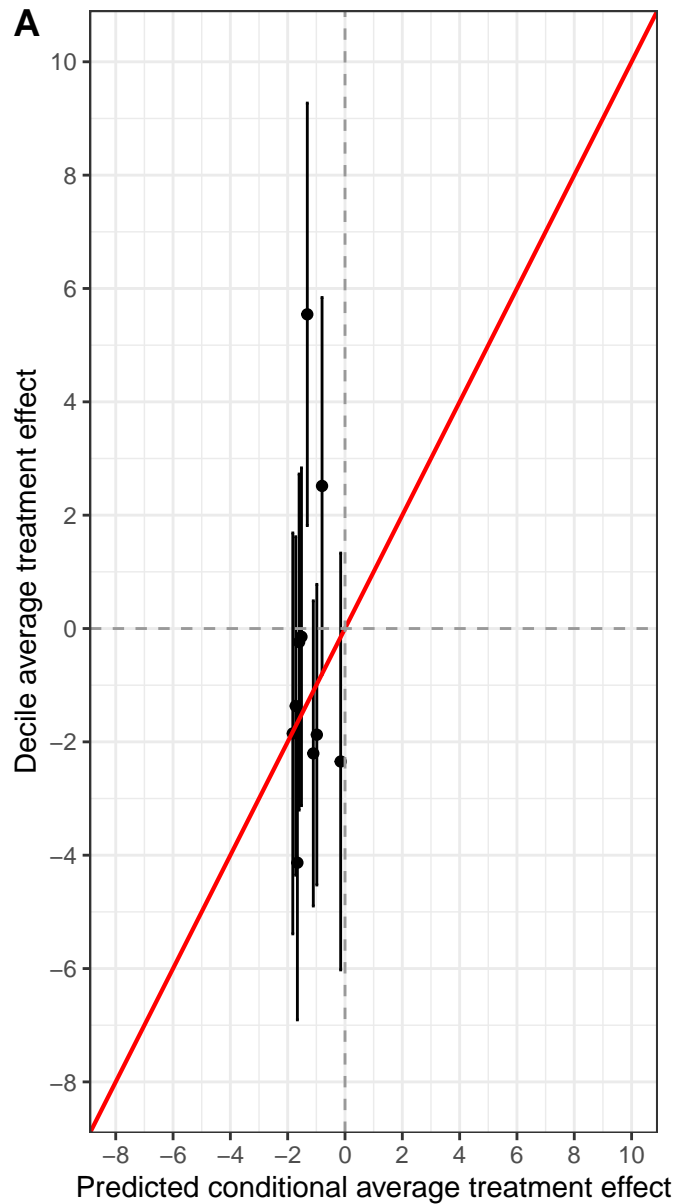
Residuals of Model 4

A**B****Standardised Residuals**

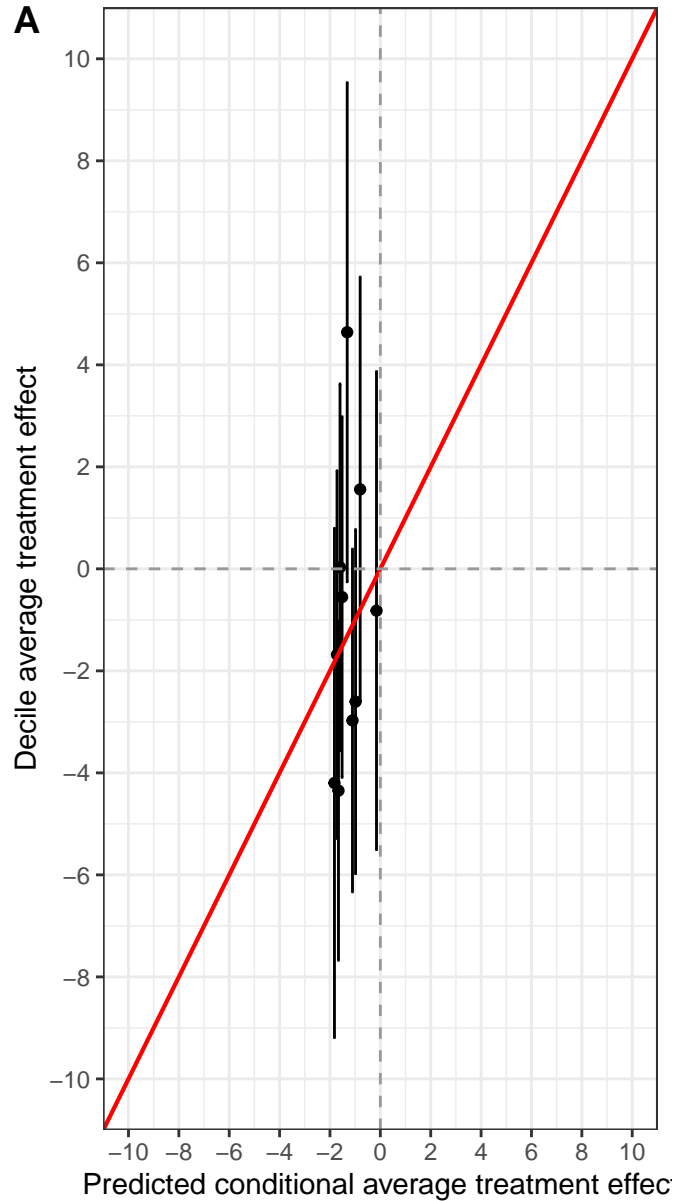
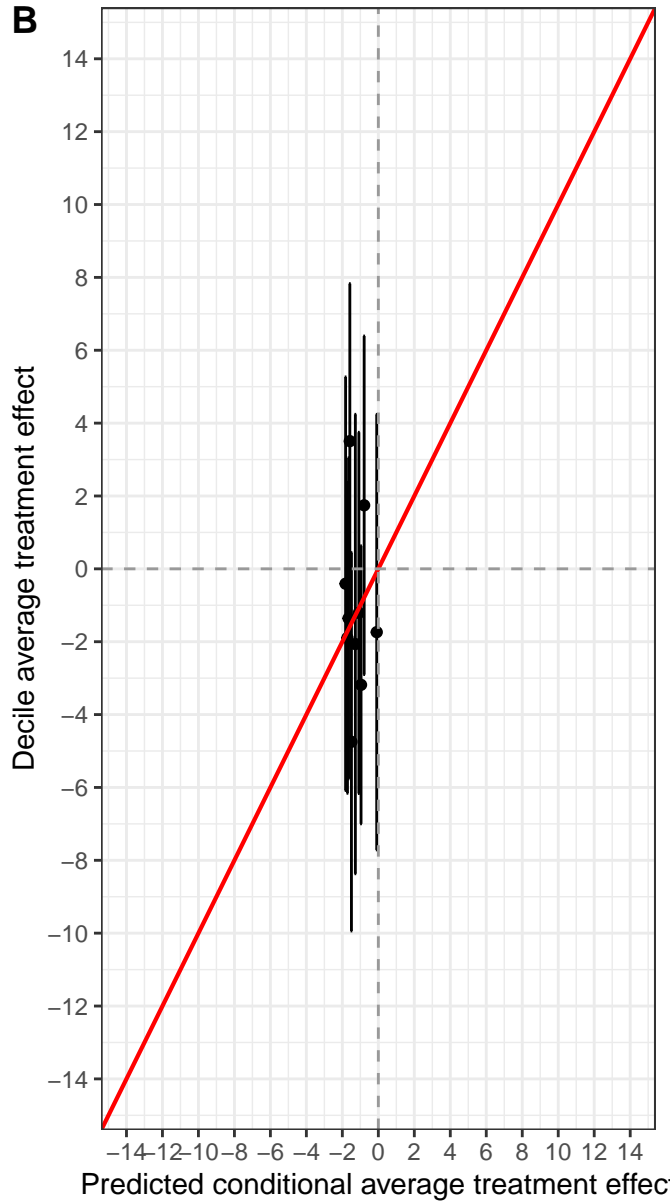
A**B**



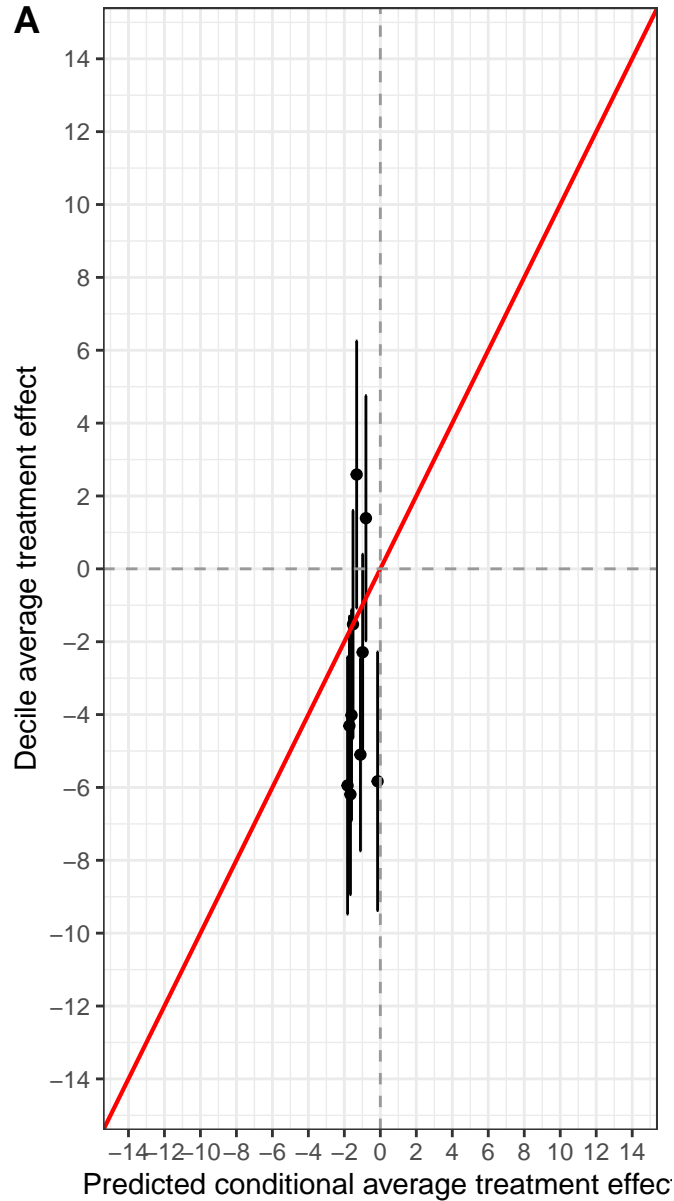
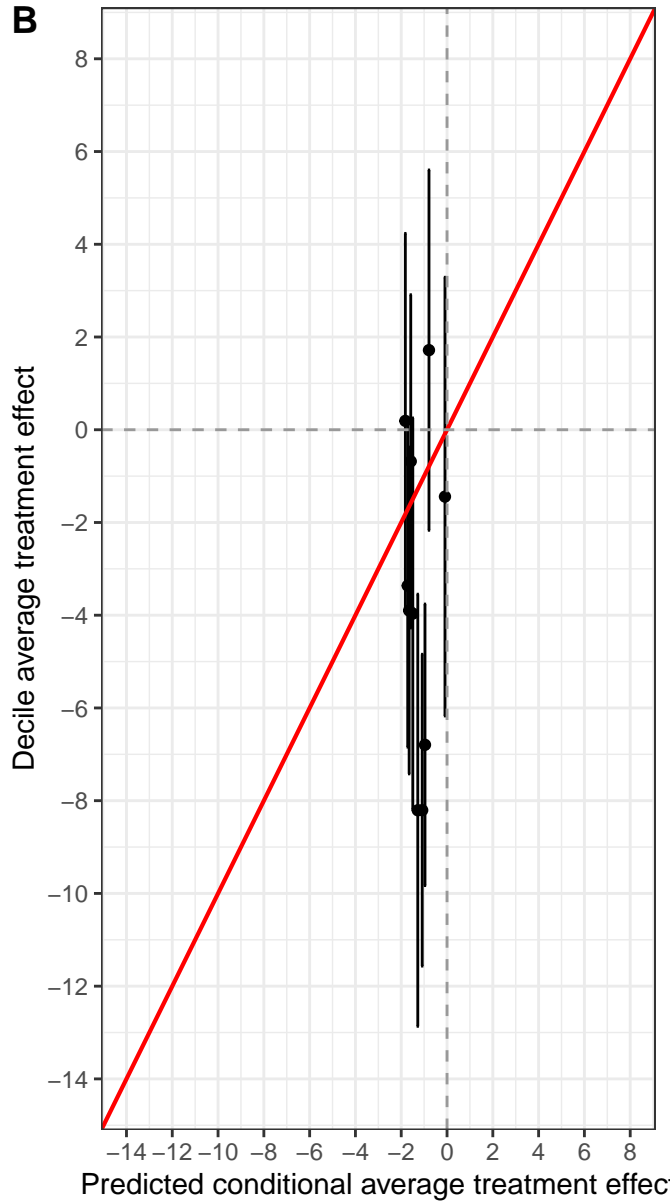
Effects validation $\text{lm}(\text{weight} \sim \text{drugclass} + \text{prop_score})$



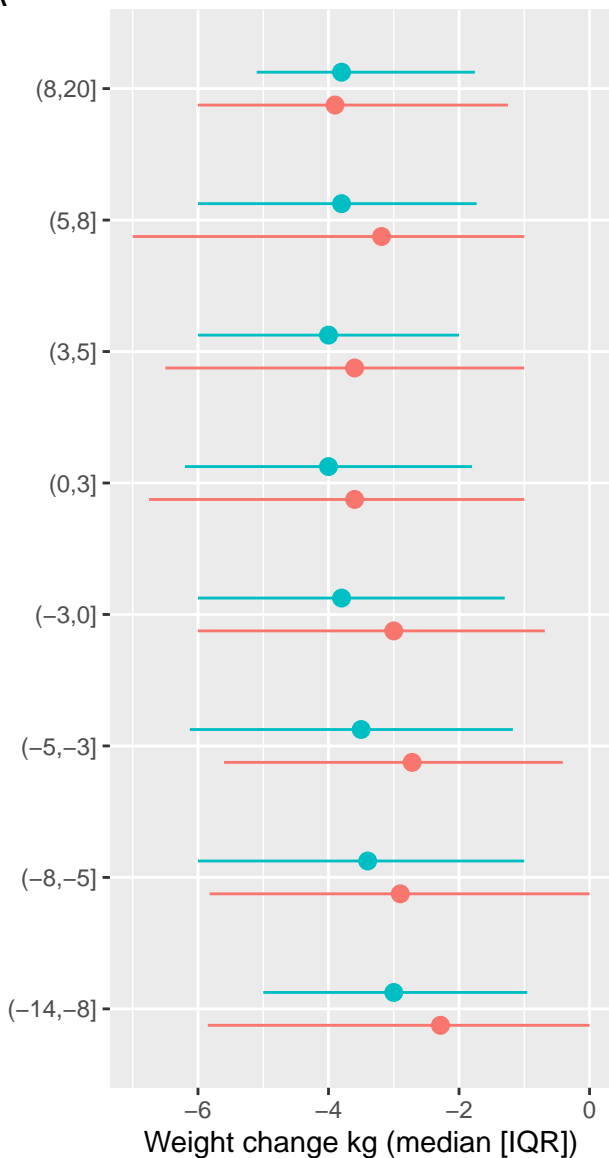
Effects validation prop score matching

A**B**

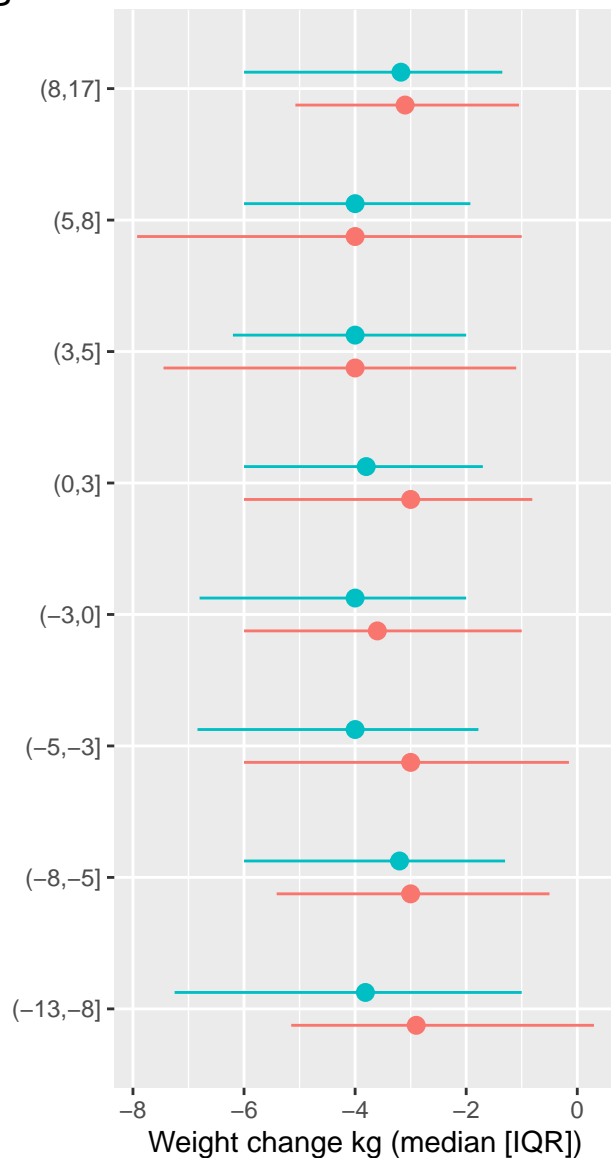
Effects validation prop score inverse weighting

A**B**

A



B



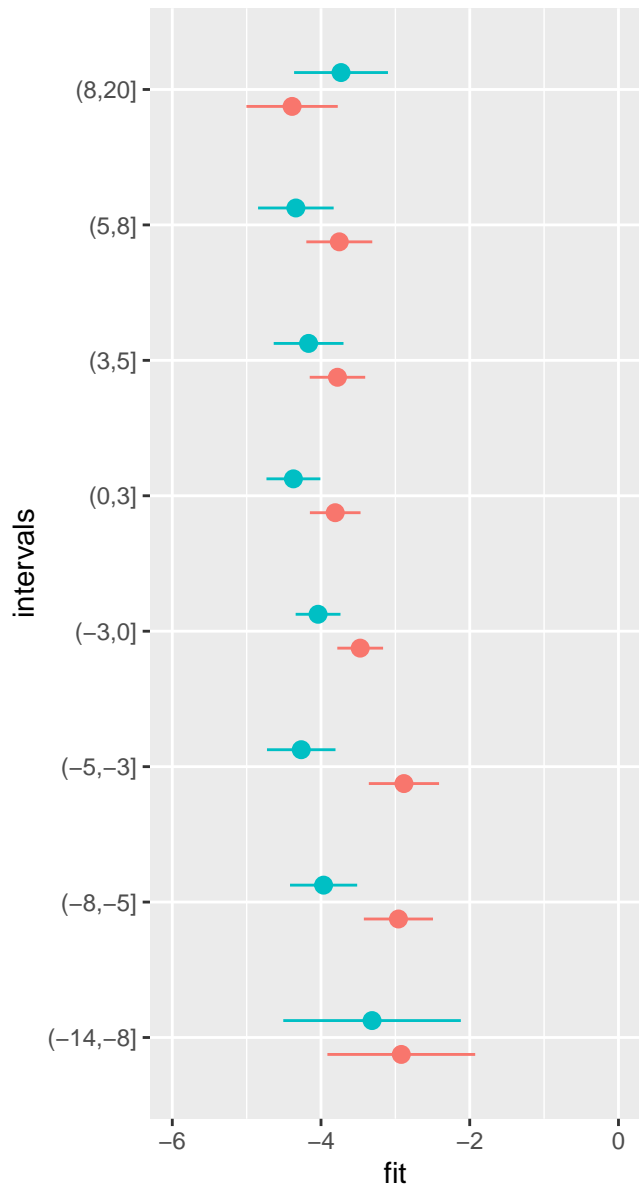
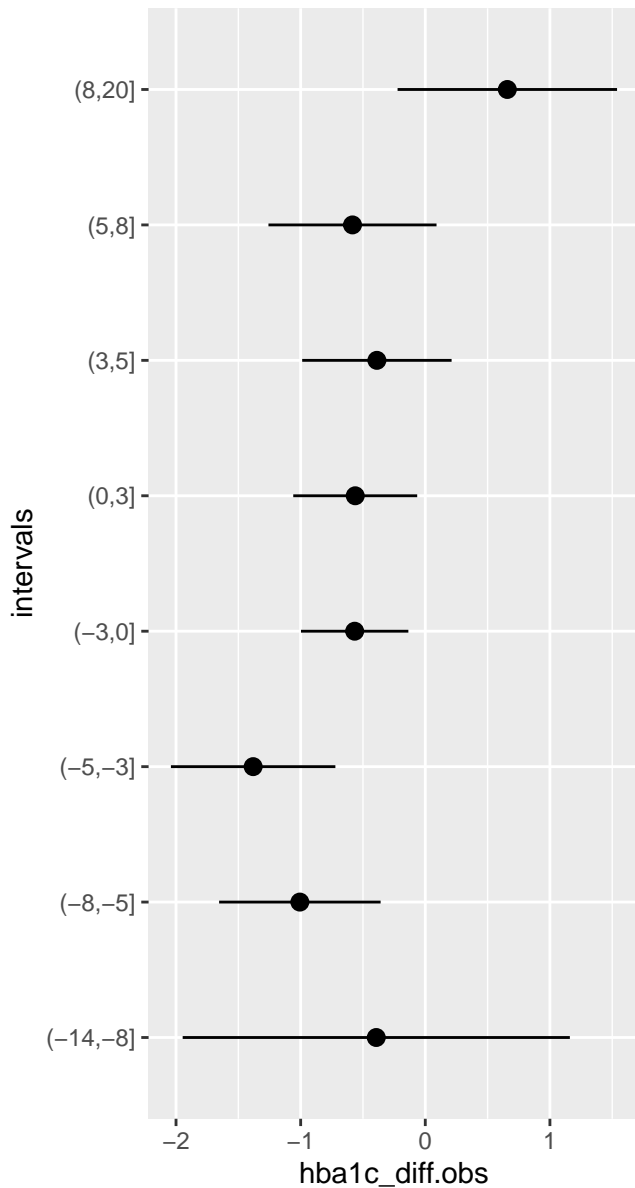
drugclass



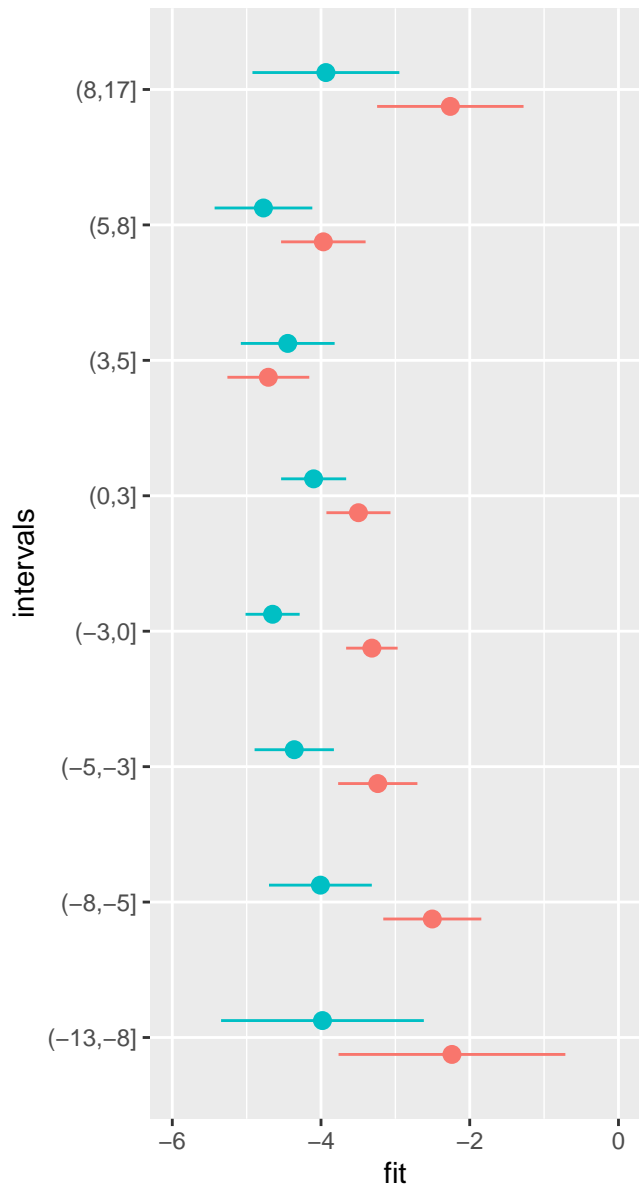
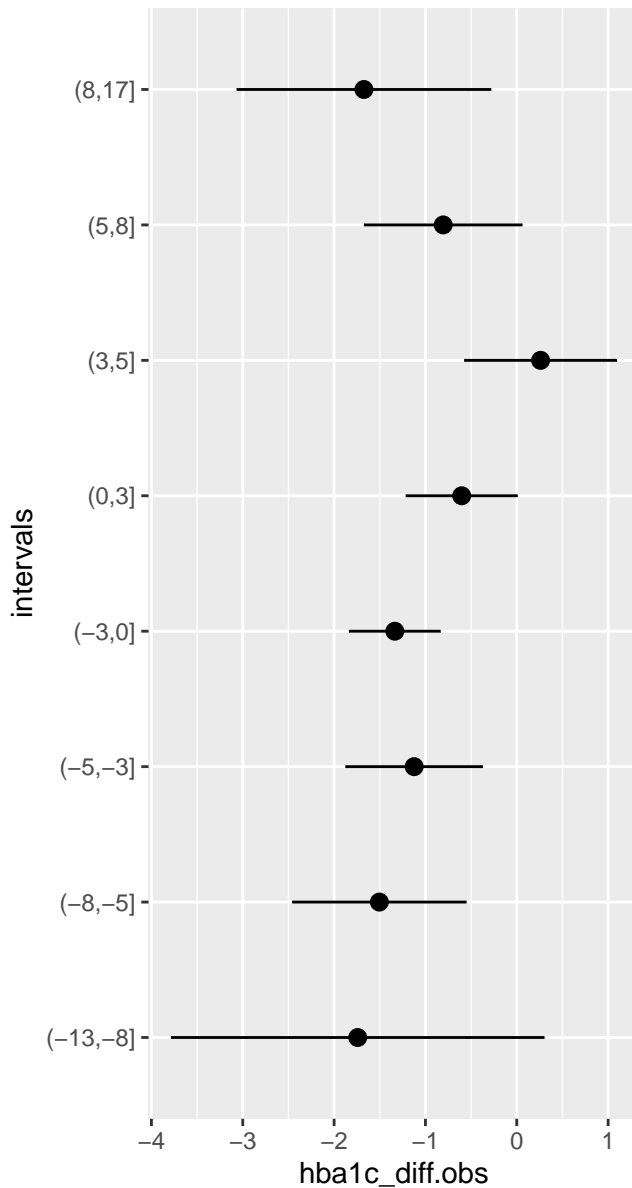
GLP1



SGLT2



drugclass GLP1 SGLT2



drugclass GLP1 SGLT2