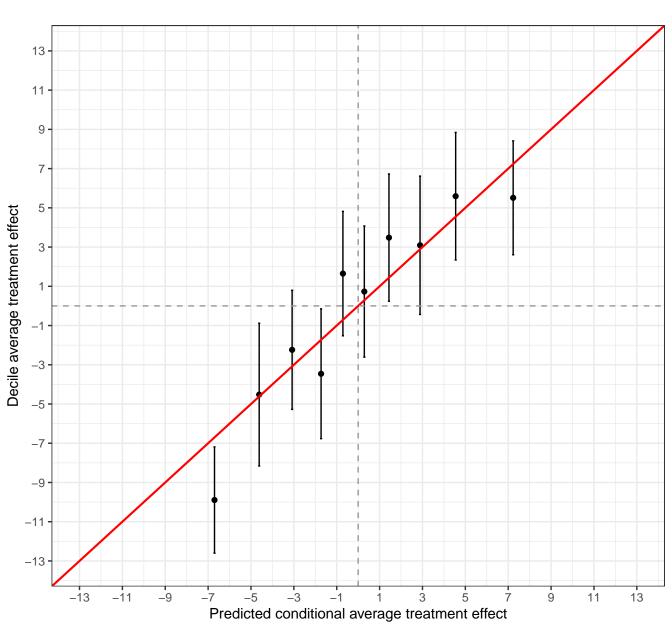
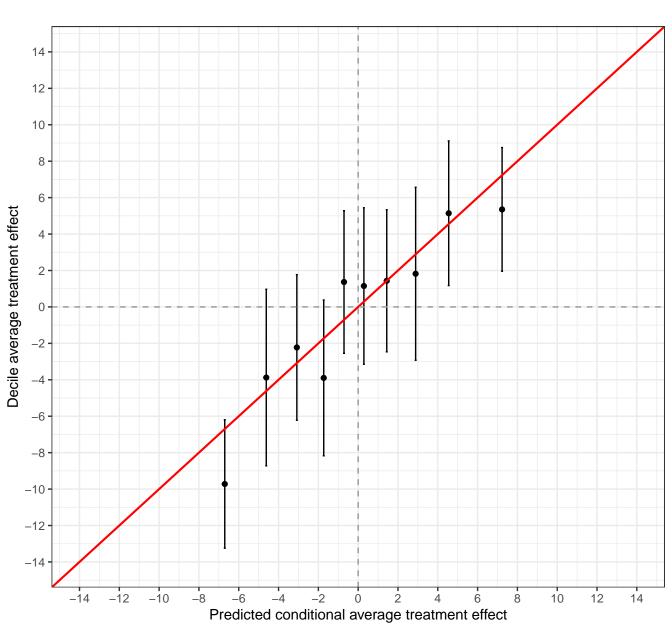


Effects validation: lm(hba1c~drugclass+prop_score)



Effects validation: propensity score matching



Effects validation: inverse propensity score weighting

