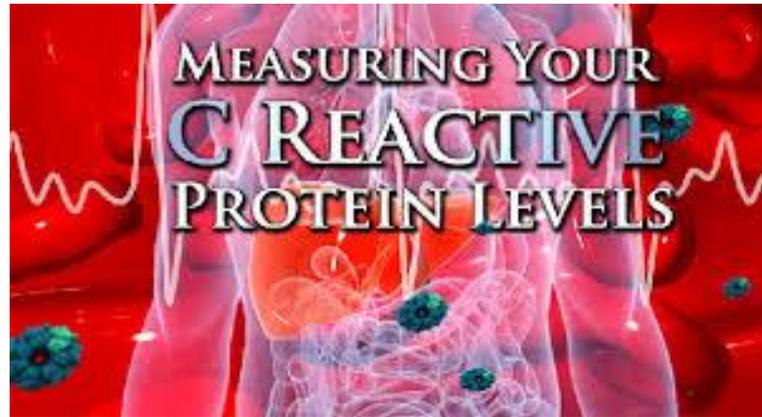


Mendelian Randomization: Practicum



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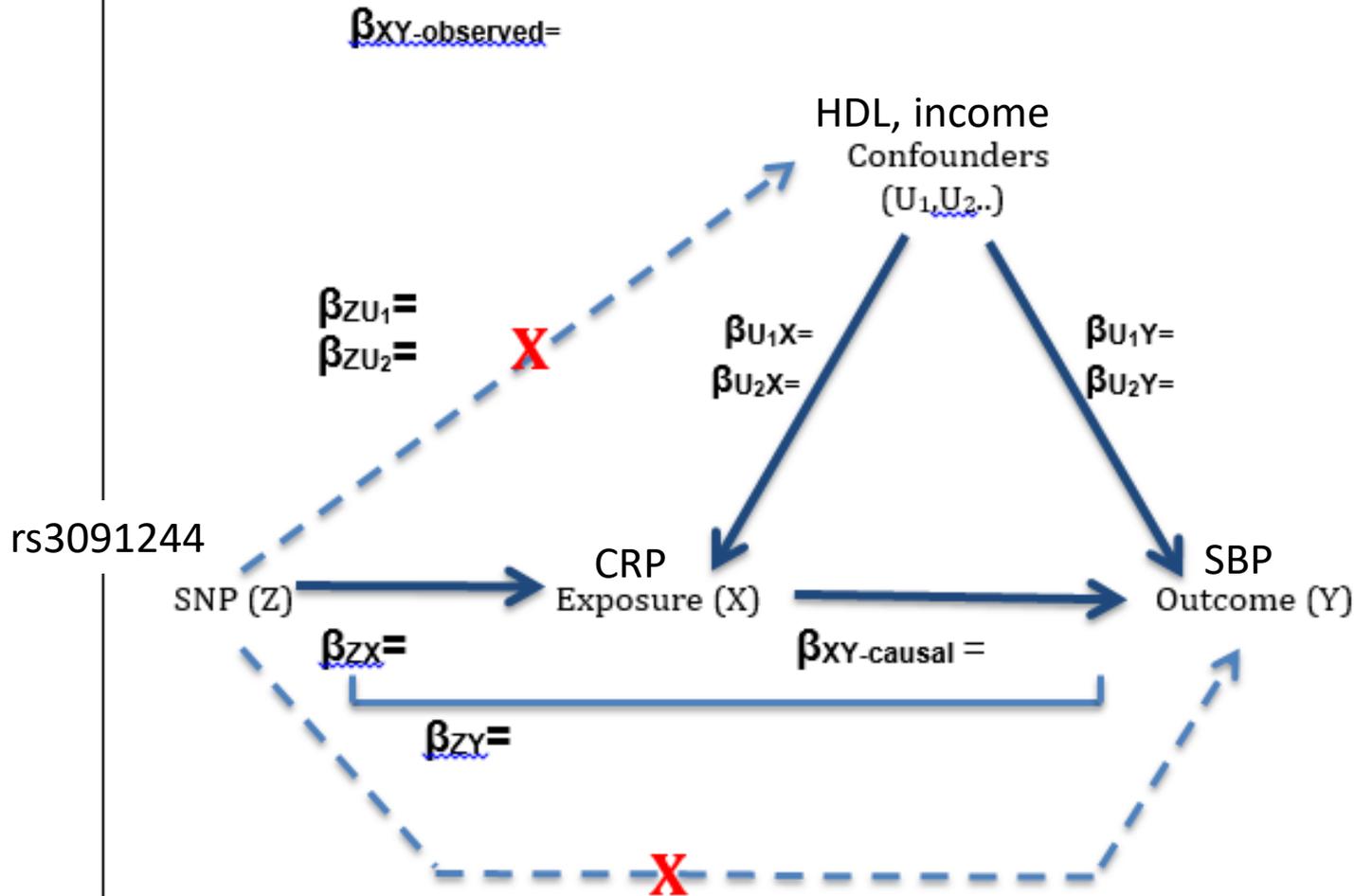
Applied research question:

Does having higher proinflammatory CRP causally increase your blood pressure?

1. Observational analyses (simple linear regressions in R)
2. MR/IV Analyses: Wald Estimator (simple linear regression)
3. MR/IV Analyses: TSLS (AER-package)
4. Manual TSLS (simple linear regression)
5. Weak instrument bias

Applied research question: Does having higher proinflammatory CRP causally increase your blood pressure?

Graphical representation of a Mendelian Randomization IV analysis



Applied research question: Does having higher proinflammatory CRP causally increase your blood pressure?

Why: to get a grip on the data, look at the MR assumptions

1. **Observational analyses** (simple linear regressions in R)
 - a. CRP-SBP OLS association
 - b. SNP rs3091244 – CRP association
 - c. confounders' (*HDL, Income*) effect on CRP & SBP
 - d. comparison unadjusted and covariate-adjusted OLS observational CRP-SBP regressions
 - e. conclude

Applied research question: Does having higher proinflammatory CRP causally increase your blood pressure?

Why: 1st MR/IV method to test causality

2. MR/IV Analyses: **Wald Estimator** (simple linear regressions)

Formulas for Wald Estimator

Where Z=SNP instrument, X=Exposure, Y=Outcome

$$\text{Causal } \beta_{IV} = \frac{\beta_{ZY}}{\beta_{ZX}}$$

$$SE_{IV} = \frac{SE_{ZY}}{\beta_{ZX}}$$

$$95\% \text{ CI} = \beta_{IV} \pm 1.96^* SE_{IV}$$

- compute the causal effect using the Wald estimator
- compare Wald with the observational OLS of CRP-SBP

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Why: 2nd method to test causality

3. MR/IV Analyses: **TSLS** (AER R-package)

General format for TSLS command:

`summary(ivreg(Outcome~Exposure | Instrument))`

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Why: 2nd method to test causality - TSLS how does it work?

4. **Manual TSLS** (two stage least squares)

a. Stage1: regress CRP on SNP

b. Stage2: use predicted CRP

`(predicted_CRP <- predict(Stage1))` to predict
SBP

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5. Weak instrument bias: an issue ?

Assessing instrument strength with the F-stat (looking for ≥ 10).
For Single SNP MR, the F-statistic is calculated as:

$$F_{\text{stat}} = \frac{R^2 * (N-1)}{(1-R^2)}$$

where R^2 is the variance explained in exposure by the SNP, and N is number of individuals in the study. This statistic is available in the output for OLS and TSLS

Applied research question: Does having higher proinflammatory CRP causally increase your blood pressure?

Steps:

1. Copy the data, the scripts and the ppt

```
mkdir MRpractical  
cp -r /faculty/camelia/2017/friday/* MRpractical
```

2. Run the analyses as described in the ppt & the R-script (easy/difficult)

3. Go to **mr.surge.sh** to provide your answers (10 MC questions)

4. As you're running the commands, fill in the graphical representation of the IV analysis with the appropriate variables and beta-coefficient

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