# Fighting Noise with Noise Causal Inference with Many Candidate Instruments

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## Acknowledgements



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# Overweight



#### Overweight is associated with Quality of Life



#### Is this association causal?



#### Unmeasured confounding!

# Mendelian randomization (MR)



Nature rolls the dice!

#### The math behind



- 1. Most genetic variants are irrelevant!
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#### Modern approach: GWAS



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**Existing solution:** Valid causal inference with some invalid instruments (Bowden et al. [2015], Kolesár et al. [2015], Kang et al. [2016], Hartwig et al. [2017], Guo et al. [2018], Hartford et al. [2020])

## Valid causal inference with some invalid instruments

Solution:

#### All happy instruments are alike; each unhappy instrument is unhappy in its own way



Plurality rule: Guo et al., 2018

# Summary of existing solutions

Step 1 GWAS based on strength of the links  $Z_j \rightarrow X, j = 1, ..., p$ 

Step 2 Apply a mode-finding algorithm to identify the valid IVs

Step 3 Use the valid IVs to estimate the causal effect

# A toy simulation



- Sample size = 500
- $Z_j, j = 1, \dots, 50, 000, U \sim N(0, 1)$
- Linear models
- True causal effect  $X \rightarrow Y$  is 2

## Step 1: GWAS

**Step 1.1 (Marginal screening)**: Select the top 500 candidate IVs  $Z_r$  based on the marginal correlation

$$|Cor(Z_j, X)|, j = 1, \dots, 50, 000$$

Step 1.2 (Joint thresholding): Fit a debiased Lasso model

 $X \sim \mathbf{Z}_r$ 

Result: On average left with 21 candidate IVs

Include the 7 relevant IVs (5 valid, 2 invalid) every single time!

# Step 2: Mode-finding

Solution:

#### All happy instruments are alike; each unhappy instrument is unhappy in its own way



# Step 3: Causal effect estimation

#### C The causal effect estimates are biased!

• Coverage probability = 0.07 (nominal = 0.95)

Recall in Step 1: Select 21 candidate IVs

- 5 valid
- 2 invalid
- 14 irrelevant with spurious correlation

# What happened in Step 2



Distribution of causal effect estimates for candidate IVs passing joint thresholding across 1000 Monte Carlo runs

All irrelevant "instruments" are alike. Why?



- Single unmeasured confounder U
- X is only determined by U



Recall that under the IV framework,

$$X \rightarrow Y = rac{Z \rightarrow X \rightarrow Y}{Z \rightarrow X} = rac{Cov(Z, Y)}{Cov(Z, X)}$$



In the case of irrelevant "instrument,"

• 
$$Cov(Z, Y) = Z \rightarrow U \rightarrow X \rightarrow Y + Z \rightarrow U \rightarrow Y$$

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The bias is the same for all irrelevant "instruments"!

In fact, also same as the bias from Y ~ X

# What happened in Step 2



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What we know so far:

- All valid instruments are alike
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How to distinguish valid instruments from irrelevant ones?

**Key idea:** Generate pseudo variables by permuting rows of Z, denoted by  $Z^*$ 

- Group 1: valid instruments in Z
- Group 2: invalid instruments in Z
- Group 3: irrelevant instruments in Z
- Group 4: irrelevant instruments in Z\*

Groups 3 and 4 are alike! We can track variables in Group 4.



Distribution of causal effect estimates for candidate IVs passing joint thresholding across 1000 Monte Carlo runs

**Step 2:** Remove causal effect estimates inside the range of pseudo effect estimates



Distribution of causal estimates after removing spurious IVs

# **Step 3:** Develop a mode-finding algorithms to identify the valid instruments



Distribution of causal estimates after mode-finding

**Step 4:** Use the selected valid instruments to estimate the causal effects



## Theoretical guarantees

Theorem 1 (Zhang et al., 2022)

(Spurious IV) Under some regularity conditions, with probability tending to 1, causal estimates for spurious IVs are concentrated within

$$[\beta^* + C_* - d, \ \beta^* + C_* + d]$$

for  $d > L(\omega, \sigma_X^2, \sigma_u^2, \sigma_Y^2, \alpha_X^*, \alpha_Y^*)$ , where  $C_* = (\alpha_Y^* \alpha_X^* \sigma_u^2)/(\alpha_X^{*2} \sigma_u^2 + \sigma_X^2)$ .

Theorem 2 (Zhang et al., 2022)

(Valid IV) Under some regularity conditions, with probability tending to 1, causal estimates for valid IVs are concentrated around  $\beta^*$ .

**Separation**: If  $d < |C_*|$ , then spurious IVs and valid IVs are separable.

Easier separation if stronger U OR sufficiently large  $\omega$ 

# Wisconsin Longitudinal Study



- Participants were graduates from Wisconsin high schools in 1957 and their siblings
- Use 3023 unrelated individuals re-interviewed in 2011
- Exposure: BMI (only those ≥ 25)
  mean (SD): 30.6 (4.9)
- Outcome: Health Utility Index Mark 3 (HUI-3)
  - between -0.22 and 1
  - mean (SD): 0.79 (0.23)

#### Crude analysis



After adjusting for age, gender and education, one unit BMI increase associated with 0.011 unit decrease in HUI-3 (95% CI = [0.009, 0.013])

## Conceptual Mendelian randomization model



 Consider baseline covariates including age, gender, education and population stratification (top 6 principal components)

## Data analysis: Prepossessing + Step 1

- Quality control 🖙 3,683,868 SNPs left
- Generate the same number of pseudo SNPs by permuting the rows of the Z matrix
- After GWAS: 44 SNPs + 42 pseudo

#### Data analysis: Steps 2-4



Causal estimate: one unit increase in BMI will result in 0.039 unit decrease of HUI-3 (95% CI = [-0.052, -0.025])

## Data analysis: Ignoring spurious instruments



Two-stage hard thresholding with voting (Guo et al., 2018): true SNPs only

Causal estimate: one unit increase in BMI will lead to 0.008 unit decrease in HUI-3 (95% CI = [-0.013, -0.003])

## Data analysis: Ignoring spurious instruments

Two-stage hard thresholding with voting (Guo et al., 2018): same data set as the proposed (true + pseudo)



Causal estimate: one unit increase in BMI will lead to 0.011 unit decrease in HUI-3 (95% CI = [-0.015, -0.008])

## Comparison of results

Proposed:

1 SD increase in BMI leads to 1 SD (roughly) decrease in HUI-3

Guo et al. (2018)'s method (ignore spurious instruments; true only): 1 SD increase in BMI leads to 0.2 SD (roughly) decrease in HUI-3

Guo et al. (2018)'s method (ignore spurious instruments; true + pseudo): 1 SD increase in BMI leads to 0.2 SD (roughly) decrease in HUI-3

Crude analysis (OLS):

1 SD increase in BMI is associated with 0.2 SD (roughly) decrease in HUI-3

## Summary

- Mendelian randomization is a powerful tool for causal effect estimation
- Challenges for MR studies
  - Find relevant IVs 🖙 GWAS
  - Deal with pleiotropy is find the mode of effect estimates
  - Bias due to spurious variables in GWAS
- Spurious IV bias is potentially a serious problem
  - Ignoring Spurious IV bias has a similar effect as ignoring unmeasured confounding bias

#### Fight noise with noise

Use pseudo variables to correct for Spurious IV bias

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