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A COMPARISON OF ANALYTICAL APPROACHES TO OBTAIN MENDELIAN RANDOMIZATION ESTIMATES WITH LONGITUDINAL EXPOSURES

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MOTIVATIONS.



- Genetic variants are proposed as an instrumental variable (IV)
- Increasingly popular approach to causal inference



- Exposures of interest are often time-varying,
 - e.g. alcohol, cholesterol levels
- Conventional IV methods designed to only handle time-fixed
 exposures
- MR estimates based on a single measurement of the exposure



Describe the **instrumental conditions** in the context of a **time-varying exposure**

OBJECTIVES.



Review the possible causal interpretations of MR estimates and the additional assumptions required to do



Propose methods to incorporate timevarying exposures in MR analyses of continuous outcomes based on gestimation of structural mean models

1. IV CONDITIONS

IV CONDITIONS.



(1) Genetic variant is associated with the exposure.

- Z = Genetic variant
- A = Exposure
- **Y** = Outcome
- **U** = (Unmeasured) confounders



(2) The genetic variant does not affect the outcome except through its potential effect on the exposure.



(3) The genetic variant and the outcome do not share common causes.

IV CONDITIONS WITH A TIME-VARYING EXPOSURE.





TIME-FIXED EXPOSURE

TIME-VARYING EXPOSURE

- **Z** = Genetic variant
- A = Exposure
- **Y** = Outcome
- U = (Unmeasured) confounders

IV CONDITIONS WITH A TIME-VARYING EXPOSURE.



TIME-FIXED EXPOSURE

- Z = Genetic variant
- A = Exposure
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TIME-VARYING EXPOSURE



Second IV condition: the genetic variant does not affect the outcome except through its possible effect on the exposure

IV CONDITIONS WITH A TIME-VARYING EXPOSURE.



TIME-FIXED EXPOSURE

- Z = Genetic variant
- A = Exposure
- Y = Outcome
- **U** = (Unmeasured) confounders



TIME-VARYING EXPOSURE



Recall: the second IV condition states that the instrument does not affect the outcome except through its possible effect on the exposure

Suppose for each individual, we have measurements on:



Outcome that an individual would have had had they received exposure level a at age m (Y^{a_m}) exposure level a' at age m ($Y^{a'_m}$)

Requires genetic variant Z to be a valid instrument for A_m (and only A_m)



Point effect:
$$E[Y^{a_m}] - E[Y^{a'_m}]$$

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• No arrows from Z to other exposure time points; or



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- No arrows from Z to other exposure time points; or
- No arrows from other exposure time points to the outcome

The assumptions to estimate the point effect in MR studies generally do not hold.





SECOND TYPE OF CAUSAL EFFECT: THE PERIOD EFFECT.

Suppose we are now interested in the effect of an exposure during a specific period,

e.g. [m - p, ..., m - 1, m]



Interested counterfactual outcomes under:

1. Some exposure trajectory

 $(a_{m-p}, \ldots, a_{m-1}, a_m)$

2. Same exposure trajectory as (1), but shifted upwards by 1 unit



SECOND TYPE OF CAUSAL EFFECT: THE PERIOD EFFECT.

If we measure the period effect with a single measurement time point:

- 1. The genetic variant Z is a valid instrument for the exposure during the period [m p, ..., m 1, m]
- 2. The association between the genetic variant and the exposure stays constant over time

This is a weaker form of the assumption required for the point effect

With multiple measurements of an exposure, we can relax this assumption by allowing the association to change over time.

For the period effect:

$$E[Y^{a_{m-p}=a,...,a_{m-1}=a,a_{m}=a}-Y^{\overline{0}}|A_{m-p},...,A_{m-1},A_{m},Z]=\gamma(a;\psi)$$

STRUCTURAL MEAN MODELS.

Model for the expected difference between two counterfactual outcomes

For the period effect:

 $E[Y^{a_{m-p}=a,...,a_{m-1}=a,a_{m}=a}-Y^{\overline{0}}|A_{m-p},...,A_{m-1},A_{m},Z]=\gamma(a;\psi)$

STRUCTURAL MEAN MODELS.

Model for the expected difference between two counterfactual outcomes Function of the exposure, indexed by the unknown parameter ψ

E.g., with one measurement of the exposure, $\gamma(a;\psi)=\psi a$

with p + 1 measurements of the exposure: $\gamma(a; \psi) = \sum_{j=m-p}^{p} \psi_j a$

Homogeneity: Z does not modify the relationship between the exposure and the outcome

G-ESTIMATION.

Define an estimating equation

 $U(\psi; Z) = H(\psi)(Z - E(Z))$ where $H(\psi) = Y - \gamma(a; \psi)$

Set $E[U(\psi; Z)] = 0$ and solve for ψ ; for continuous outcomes,

the solution is:

$$\hat{\psi} = \sum_{i=1}^{n} Y_i(Z_i - E[Z]) / \sum_{i=1}^{n} A_{m,i}(Z_i - E[Z])$$

Set the estimating equation to 0 and solve for ψ

Goal: use data from the Framingham Heart Study (FHS) to estimate the **period effect*** of shifting one's **alcohol intake trajectory** upwards by 1 drink/day on **gamma-glutamyl transferase (GGT) levels**

*The period was defined to be over the duration of follow-up



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INSTRUMENT: 26 genetic variants identified from prior studies that were combined in different ways to create multiple weighted allele scores;

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EXPOSURES: alcohol intake (via FFQ) at examination 1 and examination 2

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OUTCOME: gamma-glutamyl transferase levels (measured in blood) at examination 2; marker for liver function

APPLICATION: ANALYSIS.

Proposed two structural mean models

Only consider a single exposure measurement

- 1. Exposure measured at examination 1
- 2. Exposure measured at examination 2
- 3. Average exposure across the two examination time points

Consider multiple exposure measurements Estimated the parameters using different weighted allele scores

i
$$E[Y - Y^{\overline{0}} | A_{m-p}, ..., A_{m-1}, A_m, Z] = \psi_1 A_m$$

i
$$E[Y - Y^{\overline{0}}|A_{m-p}, ..., A_{m-1}, A_m, Z] = \sum_{j=1}^{2} \psi_j A$$

Examination 1 only Examination 2 only Average Joint, algorithm 1 Joint, algorithm 2 Joint, algorithm 3

Highlights from our data application:

- Similarity of results driven by limited number of exposure measurements
- Precision of estimates

Takeaway: estimates were similar regardless of choice of model, although confidence intervals vary

8.0

2.0

Period effect of alcohol

intake on GGT levels

4.0

APPLICATION: RESULTS.

3. STRUCTURAL MEAN MODELS



Possible **causal interpretations** of MR estimates and their assumptions

- Point effect
- Period effect

SUMMARY.



G-estimation of **structural mean models** to accommodate multiple measures of a time-varying exposure in an MR analysis.



Advantages of our proposed method:

- Include >2 time points
- Model non-linear outcomes
- Include other covariates into the model

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SCHOOL OF PUBLIC HEALTH

QUESTIONS.