

Addressing Confounding and Continuous Exposure Measurement Error Using Corrected Score Functions

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Acknowledgements

Bryan Blette, PhD



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Peter Gilbert, PhD



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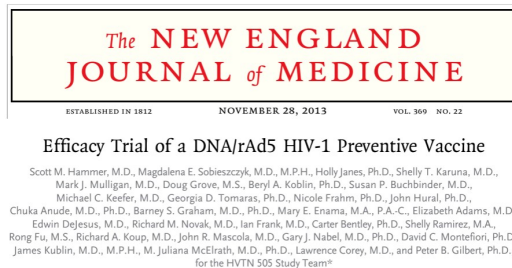
Michael Hudgens, PhD



UNC Chapel Hill

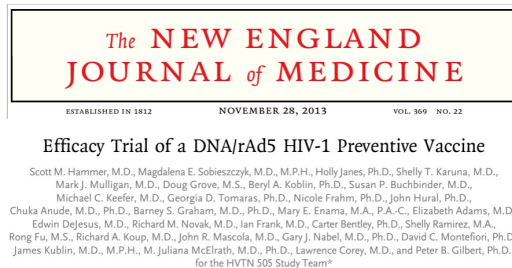
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Motivation: HVTN 505 trial



- **HVTN 505 trial:** trial of a preventive HIV vaccine

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- Stopped early after reaching predetermined cutoffs for efficacy futility [2]

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The Journal of Infectious Diseases

MAJOR ARTICLE

 IDSA
Infectious Diseases Society of America

 hivma
hiv medicine association

 OXFORD

Higher T-Cell Responses Induced by DNA/rAd5 HIV-1 Preventive Vaccine Are Associated With Lower HIV-1 Infection Risk in an Efficacy Trial

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“What would be the risk of HIV if, possibly counter to fact, somebody were to have biomarker level \mathbf{a} ?”

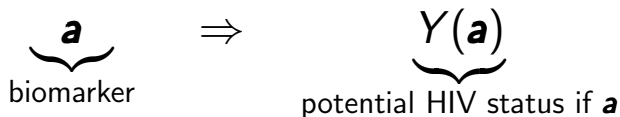
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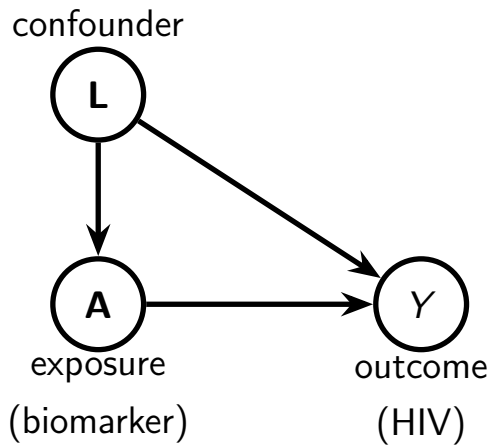


Estimand (**dose-response surface**): $\eta(\mathbf{a}) \equiv E\{Y(\mathbf{a})\}$ for $\mathbf{a} \in \mathcal{A}$

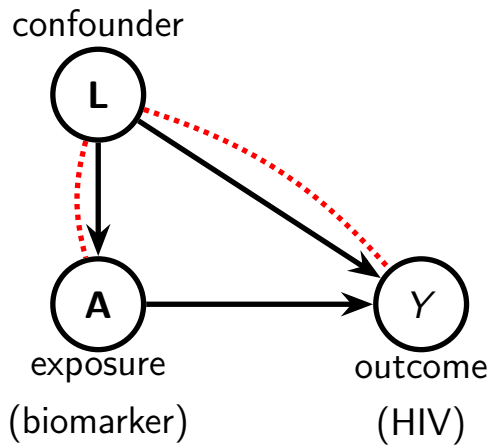
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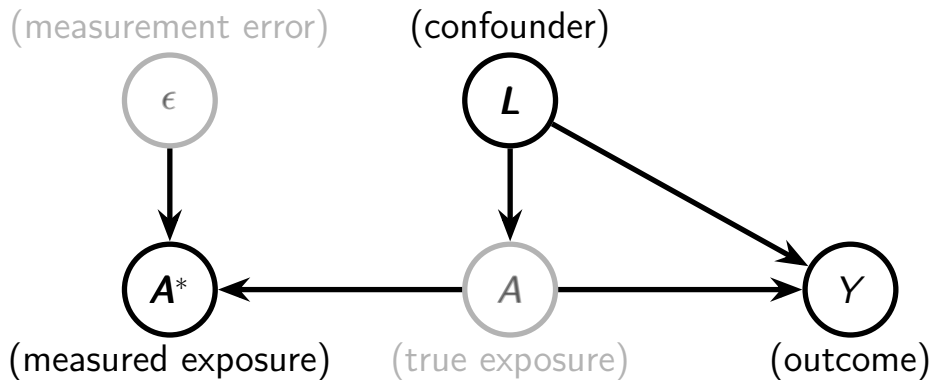
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Measurement Error



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- ② Inverse Probability Weighting (IPW)

Addressing Confounding Alone

How can we address confounding in the absence of measurement error?

- 3 classical methods:
 - 1 G-Formula
 - 2 Inverse Probability Weighting (IPW)
 - 3 Doubly Robust Method (DR)
- These can all be framed as **M-estimators**

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Score Function: a function of the observed data and the parameter of interest

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$$E\left\{ \Psi_0(Y, L, A; \underbrace{\theta_0}_{\text{true}}) \right\} = \mathbf{0}.$$

Crash Course on M-Estimation (cont'd)

- Given a score function $\boldsymbol{\psi}_0$ and observed data $\{(Y_i, \mathbf{L}_i, \mathbf{A}_i) : i = 1, \dots, n\}$, we can find an estimator $\hat{\boldsymbol{\theta}}$ as the solution to

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- $\hat{\boldsymbol{\theta}}$ is consistent and asymptotically normal and has a simple variance estimator [6].

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This can be expressed as an M-estimator with score function

$$\Psi_{0-GF}(Y, \mathbf{L}, \mathbf{A}; \theta_{GF}) = \begin{bmatrix} \{Y - \mu(\mathbf{L}, \mathbf{A}; \beta)\} \partial_{\beta} \mu(\mathbf{L}, \mathbf{A}; \beta) \\ \eta(\mathbf{a}) - \mu(\mathbf{L}, \mathbf{a}; \beta) \end{bmatrix}$$

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Doubly robust* to models for $\mu(\mathbf{L}, \mathbf{A}; \beta)$ and $f_{\mathbf{A}|\mathbf{L}}(\mathbf{A}|\mathbf{L})$.

Addressing Confounding and Measurement Error

Can we just substitute \mathbf{A}^* for \mathbf{A} and find the solution to, e.g.,

$$\sum_{i=1}^n \psi_0(Y_i, L_i, \underbrace{A_i^*}_{\text{mismeasured}}; \theta) = \mathbf{0}?$$

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No! This leads to bias in $\hat{\boldsymbol{\theta}}$ because

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No! This leads to bias in $\hat{\theta}$ because

$$E\{\psi_0(Y, L, \mathbf{A}^*; \theta_0)\} \neq \mathbf{0}.$$

We need a new score function ψ_{CS} such that

$$E\left\{ \underbrace{\psi_{CS}}_{\text{new score fun.}} (Y, L, \underbrace{\mathbf{A}^*}_{\text{mismeasured}}; \theta_0) \right\} = \mathbf{0}.$$

Corrected Score Functions

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- 1 Add additional *imaginary* measurement error $i\tilde{\epsilon}$ to the mismeasured exposure
- 2 Keep only the real part of the resulting complex-valued function
- 3 Take the expectation over the additional measurement error $\tilde{\epsilon}$.

Corrected Score Functions (contd)

Under certain conditions, the corrected score function Ψ_{CS} is then unbiased, meaning

$$E\{\Psi_{CS}(Y, L, \underbrace{A^*}_{\text{mismeasured}}; \theta_0)\} = \mathbf{0}.$$

Corrected Score Functions (contd)

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The G-Formula, IPW, and DR score functions all satisfy these conditions, and so can be “corrected.”

$$\Psi_{GF} \longrightarrow \Psi_{CS-GF}$$

$$\Psi_{IPW} \longrightarrow \Psi_{CS-IPW}$$

$$\Psi_{DR} \longrightarrow \Psi_{CS-DR}$$

Monte Carlo Corrected Score Functions

- Sometimes we can find a closed-form algebraic expression for

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$$\mathbb{E}\{f(\tilde{\boldsymbol{\epsilon}})\} \approx B^{-1} \sum_{b=1}^B f(\tilde{\boldsymbol{\epsilon}}_b)$$

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$$\boldsymbol{\Psi}_{CS}(Y, \mathbf{L}, \mathbf{A}^*; \boldsymbol{\theta}) = \underbrace{\mathbb{E}[\operatorname{Re}\{\boldsymbol{\Psi}_0(Y, \mathbf{L}, \mathbf{A}^* + i\tilde{\boldsymbol{\epsilon}}; \boldsymbol{\theta})\} \mid Y, \mathbf{L}, \mathbf{A}^*]}_{\mathbb{E}\{f(\tilde{\boldsymbol{\epsilon}})\}}.$$

- Alternatively, we can approximate this expectation with Monte Carlo replicates

$$\begin{aligned}\mathbb{E}\{f(\tilde{\boldsymbol{\epsilon}})\} &\approx B^{-1} \sum_{b=1}^B f(\tilde{\boldsymbol{\epsilon}}_b) \\ \implies \boldsymbol{\Psi}_{MCCS}^B(Y, \mathbf{L}, \mathbf{A}^*; \boldsymbol{\theta}) &= B^{-1} \sum_{b=1}^B \operatorname{Re}\{\boldsymbol{\Psi}_0(Y, \mathbf{L}, \mathbf{A}^* + i\tilde{\boldsymbol{\epsilon}}_b; \boldsymbol{\theta})\}\end{aligned}$$

Simulation Setting

$$L \sim \mathcal{N}(0, 0.36)$$



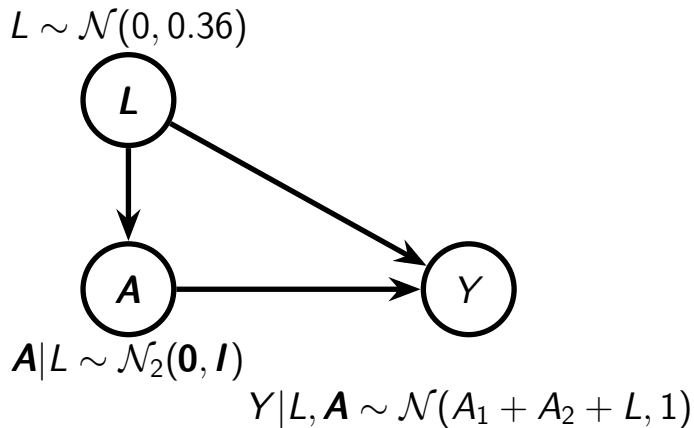
Simulation Setting

$$L \sim \mathcal{N}(0, 0.36)$$



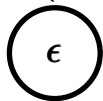
$$\mathbf{A}|L \sim \mathcal{N}_2(\mathbf{0}, I)$$

Simulation Setting

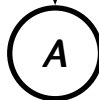


Simulation Setting

$$\epsilon \sim \mathcal{N}_2(\mathbf{0}, \sigma_{me}^2 \mathbf{I})$$



$$L \sim \mathcal{N}(0, 0.36)$$

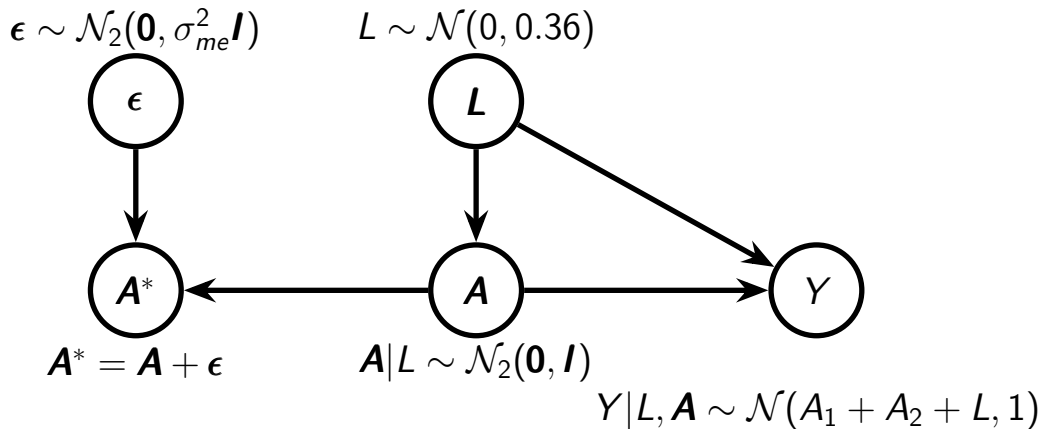


$$\mathbf{A}|L \sim \mathcal{N}_2(\mathbf{0}, \mathbf{I})$$

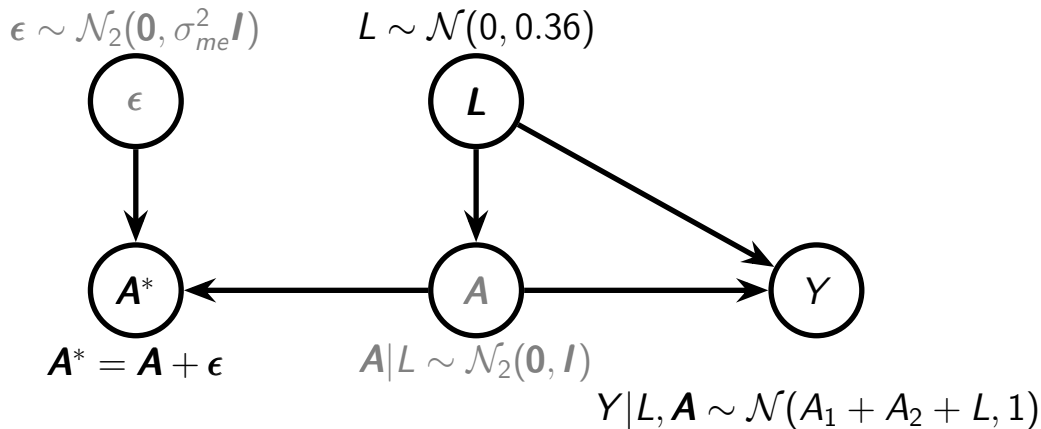


$$Y|L, \mathbf{A} \sim \mathcal{N}(A_1 + A_2 + L, 1)$$

Simulation Setting



Simulation Setting



Simulation Setting

- implies a dose-response curve of $\eta(\mathbf{a}; \gamma) = \gamma_0 + \underbrace{\gamma_1}_{\text{estimand}} a_1 + \gamma_2 a_2$

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- sample size $n = 800$

Simulation Setting

- implies a dose-response curve of $\eta(\mathbf{a}; \gamma) = \gamma_0 + \underbrace{\gamma_1}_{\text{estimand}} a_1 + \gamma_2 a_2$
- sample size $n = 800$
- 2 estimators compared:

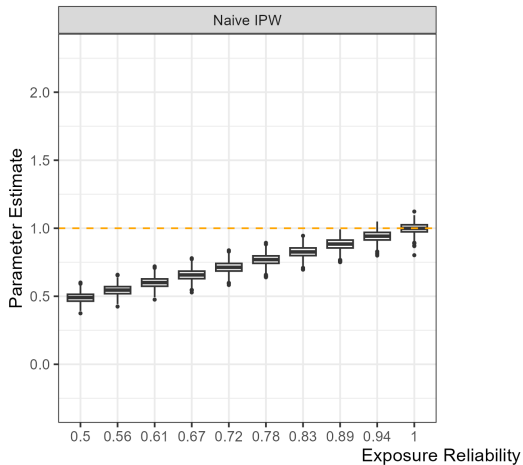
Simulation Setting

- implies a dose-response curve of $\eta(\mathbf{a}; \gamma) = \gamma_0 + \underbrace{\gamma_1}_{\text{estimand}} a_1 + \gamma_2 a_2$
- sample size $n = 800$
- 2 estimators compared:
 - ① naive IPW (ignores measurement error)

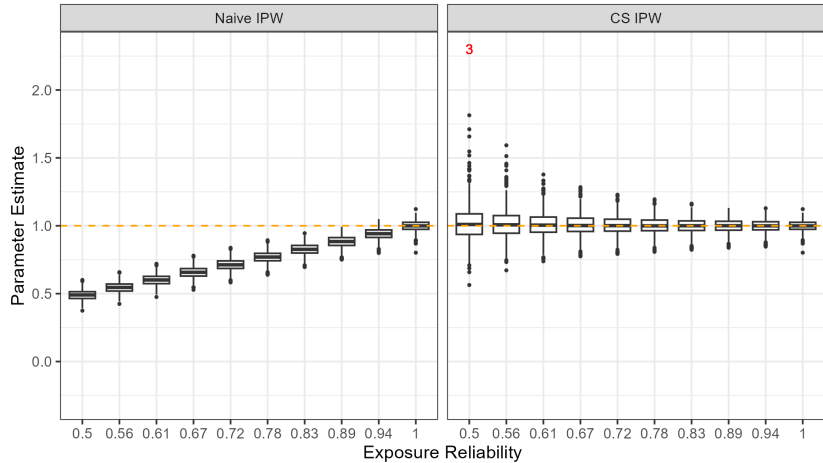
Simulation Setting

- implies a dose-response curve of $\eta(\mathbf{a}; \gamma) = \gamma_0 + \underbrace{\gamma_1}_{\text{estimand}} a_1 + \gamma_2 a_2$
- sample size $n = 800$
- 2 estimators compared:
 - 1 naive IPW (ignores measurement error)
 - 2 Corrected Score IPW

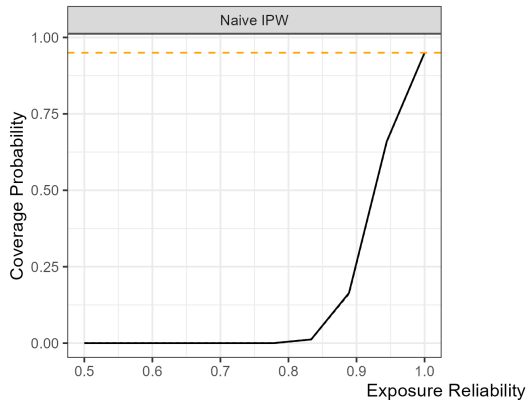
Simulation Results: Estimator



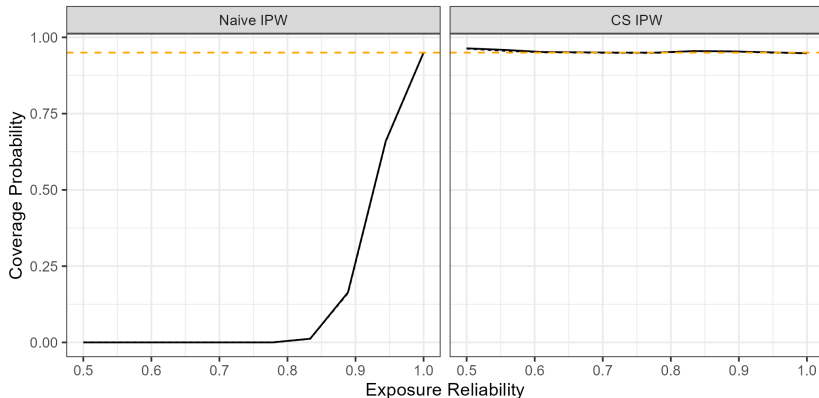
Simulation Results: Estimator



Simulation Results: Confidence Interval



Simulation Results: Confidence Interval



Application: HVTN 505 Trial

- **two exposures** (both log-transformed):

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 - (ii) recruitment of $\text{Fc}\gamma\text{RIIa}$ of the H131-Con S gp140 protein (**RII**)

Application: HVTN 505 Trial

- **two exposures** (both log-transformed):
 - (i) antibody-dependent cellular phagocytosis (**ADCP**)
 - (ii) recruitment of $\text{Fc}\gamma\text{RIIa}$ of the H131-Con S gp140 protein (**RII**)
- **case-cohort sampling**: biomarkers only measured in stratified random sample of controls

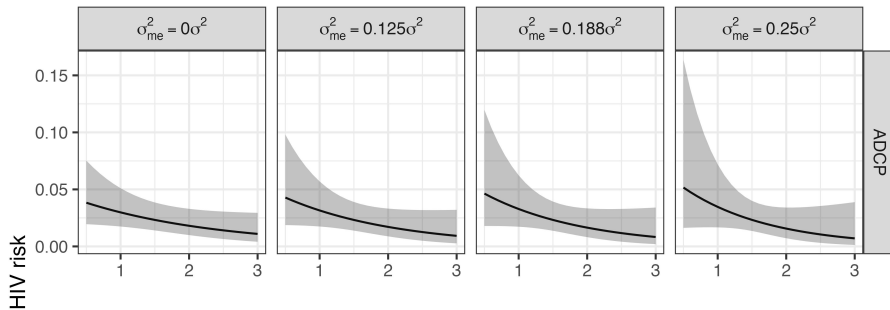
Application: HVTN 505 Trial

- **two exposures** (both log-transformed):
 - (i) antibody-dependent cellular phagocytosis (**ADCP**)
 - (ii) recruitment of $\text{Fc}\gamma\text{RIIa}$ of the H131-Con S gp140 protein (**RII**)
- **case-cohort sampling**: biomarkers only measured in stratified random sample of controls
- **covariates**: age, race, BMI, behavior risk, CD4-P, and CD8-P

Application: HVTN 505 Trial

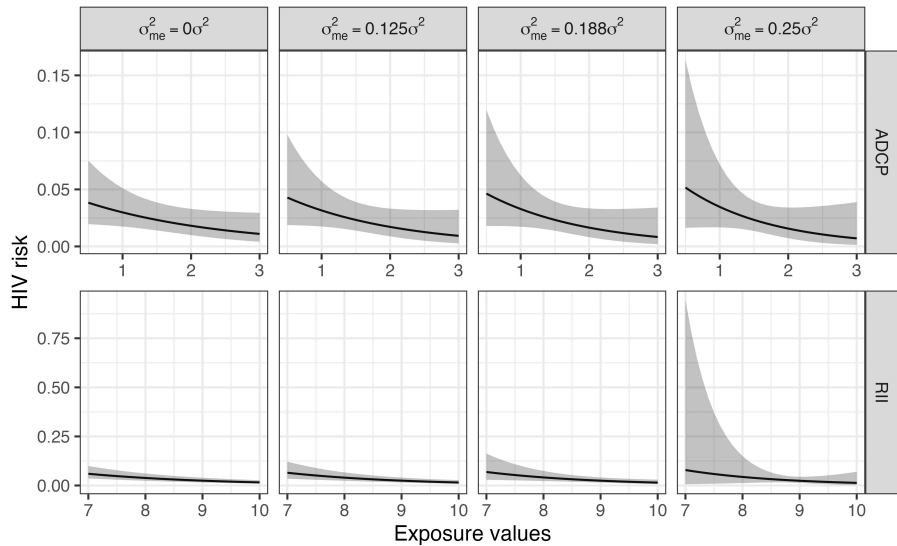
- **two exposures** (both log-transformed):
 - (i) antibody-dependent cellular phagocytosis (**ADCP**)
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- **case-cohort sampling**: biomarkers only measured in stratified random sample of controls
- **covariates**: age, race, BMI, behavior risk, CD4-P, and CD8-P
- **analysis**: DR estimator with a log-linear outcome model

Application: HVTN 505 Trial



Exposure values

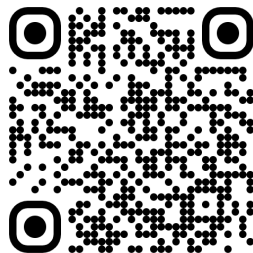
Application: HVTN 505 Trial



Mismex: Causal Inference for Mismeasured Exposures



Paper in *Biometrics*



GitHub R package

References

- [1] Youyi Fong, Xiaoying Shen, Vicki C Ashley, Aaron Deal, Kelly E Seaton, Chencheng Yu, Shannon P Grant, Guido Ferrari, Allan C deCamp, Robert T Bailer, et al. Modification of the association between T-cell immune responses and human immunodeficiency virus type 1 infection risk by vaccine-induced antibody responses in the HVTN 505 trial. *The Journal of Infectious Diseases*, 217(8):1280–1288, 2018.
- [2] Scott M Hammer, Magdalena E Sobieszczyk, Holly Janes, Shelly T Karuna, Mark J Mulligan, Doug Grove, Beryl A Koblin, Susan P Buchbinder, Michael C Keefer, Georgia D Tomaras, et al. Efficacy trial of a DNA/rAd5 HIV-1 preventive vaccine. *New England Journal of Medicine*, 369(22):2083–2092, 2013.
- [3] Holly E Janes, Kristen W Cohen, Nicole Frahm, Stephen C De Rosa, Brittany Sanchez, John Hural, Craig A Magaret, Shelly Karuna, Carter Bentley, Raphael Gottardo, et al. Higher T-cell responses induced by DNA/rAd5 HIV-1 preventive vaccine are associated with lower HIV-1 infection risk in an efficacy trial. *The Journal of Infectious Diseases*, 215(9):1376–1385, 2017.
- [4] Scott D Neidich, Youyi Fong, Shuying S Li, Daniel E Geraghty, Brian D Williamson, William Chad Young, Derrick Goodman, Kelly E Seaton, Xiaoying Shen, Sheetal Sawant, et al. Antibody Fc effector functions and IgG3 associate with decreased HIV-1 risk. *Journal of Clinical Investigation*, 129(11):4838–4849, 2019.
- [5] Steven J Novick and Leonard A Stefanski. Corrected score estimation via complex variable simulation extrapolation. *Journal of the American Statistical Association*, 97(458):472–481, June 2002. ISSN 0162-1459, 1537-274X. doi: 10.1198/016214502760047005. URL <http://www.tandfonline.com/doi/abs/10.1198/016214502760047005>.
- [6] Leonard A Stefanski and Dennis D Boos. The calculus of M-estimation. *The American Statistician*, 56(1):29–38, 2002.

Appendix

Appendix: Corrected Score Functions

- Suppose the oracle score function is **conditionally unbiased**, meaning

$$E\{\Psi_0(Y, L, \mathbf{A}; \theta) | \mathbf{A}\} = 0.$$

- Define the corrected score function as

$$\Psi_{CS}(Y, L, \mathbf{A}^*; \theta) = E \left[\text{Re} \left\{ \Psi_0(Y, L, \tilde{\mathbf{A}}; \theta) \right\} \mid Y, L, \mathbf{A}^* \right],$$

where $\tilde{\mathbf{A}} = \mathbf{A}^* + i\tilde{\epsilon}$, $i = \sqrt{-1}$, $\text{Re}(\cdot)$ denotes the real component of a complex number, and $\tilde{\epsilon} \sim \mathcal{N}(\mathbf{0}, \Sigma_{me})$.

- Then

$$\begin{aligned} E\{\Psi_{CS}(Y, L, \mathbf{A}^*; \theta) \mid Y, L, \mathbf{A}\} &= \Psi_0(Y, L, \mathbf{A}; \theta) \\ \implies E[E\{\Psi_{CS}(Y, L, \mathbf{A}^*; \theta) \mid Y, L, \mathbf{A}\}] &= E\{\Psi_0(Y, L, \mathbf{A}; \theta)\} \\ \implies E\{\Psi_{CS}(Y, L, \mathbf{A}^*; \theta)\} &= 0 \end{aligned}$$