# Improving Inferences from Randomized Trials: Using per-protocol analyses obtain better estimates of HIV treatment effects.

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#### Outline

**RCTs** 

Per-protocol effects

Example using ACTG 5202 Trial

Population

Analysis Plan

Results

Limitations and Future

Closing Remarks

# Randomized Trials are a gold standard

- Require clear enrollment criteria.
- Unabmiguous intervention protocol.
- Exchangeability:  $Y^a \perp \!\!\!\perp A$  for  $A \in \{0,1\}$
- Consistency:  $Y = Y^{a=1}A + Y^{a=0}(1 A)$
- Positivity: Pr(A = a) > 0
- This allows for unbiased estimation of treatment effects.

 $Y^a=\mbox{counterfactual}$  outcome under treatment  $A=a,\ Y=\mbox{observed}$  outcome,  $A=\mbox{treatment}$ 

#### RCT estimands

- Intention-to-treat (ITT) effect:  $E[Y^{r=1}] E[Y^{r=0}]$
- This is the effect of treatment assignment on outcomes.
- Public health focused.

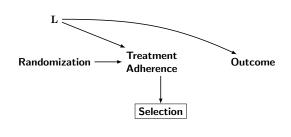
- Typical Per-protocol (PP) effect:  $E[Y^{r=1,\bar{a}=1}] E[Y^{r=0,\bar{a}=0}]$
- This is the effect of treatment assignment and adherence on outcomes.
- Patient focused.<sup>1</sup>

 $R = \text{randomization}; \bar{A} = \text{history of treatment}$ 

<sup>&</sup>lt;sup>1</sup>Hernan and Robins, NEJM 2016

# Per-protocol effects can be biased

- Frequently done by excluding those not adhering.<sup>1</sup>
- Susceptible to selection bias.
- Can be addresed: e.g with inverse probability weighting (see next talk).



<sup>&</sup>lt;sup>1</sup>Cole *et al.* JAMA Net. Open 2023, Dodd *et al.* Trials 2012

### There is no *one* per-protocol effect

- Accounts for adherence.
- "Doc, what if I take all my doses like you tell me to?"
- There are at least 6 per-protocol parameters that can be estimated.<sup>1</sup>
- There are also  $k \ge 1$  protocols depending on how the investigator(s) define adherence.

<sup>&</sup>lt;sup>1</sup>Rudolph et al. Epidemiology 2020

#### Per Protocol Causal Identification

Let g be a deterministic treatment strategy and overbar denote a history of values

• Conditional Exchangeability:

$$Y^g \perp (A_t, C_{t+1}) \mid (\bar{A}_{t-1} = \bar{a}_{t-1}^g, \bar{L}_t = \bar{\ell}_t, C_t = Y_t = 0) \quad \forall t$$

- Consistency: if  $\bar{A}_t$  =  $\bar{A}_t^g$  then  $\bar{Y}_t$  =  $\bar{Y}_t^g$
- Positivity:  $f(a_t^g, C_t = 0 \mid \bar{a}_{t-1}^g, \bar{\ell}_t, C_t = Y_t = 0) > 0 \ \forall t$
- No missclassification and correct model specification.

Y= outcome, C= censoring, A= treatment , L= covariates, t= time point. Wen et al. Biometrics 2019

### Example: adherence in HIV treatment

- Adherence needed for HIV viral suppression varies by treatment regimen.
- Blanket recommendations fail.
- Understanding of how is critical<sup>a</sup> for:

0.8

0.7

0.6

All regimen types

INSTI-based regimens<sup>‡,¶</sup>

PDC\*= 0.82

0.8

0.6

<sup>0.8</sup> 0.7 adherence impacts efficacy 0.6 NNRTI-based regimens<sup>§,¶</sup> Developing new treatments. suppression r 80 Maximizing current 0.7 treatments. 0.6 Adimora. Cole and Eron CID 2017 0.2 Antiretroviral PDC\* level

# An example using an HIV Trial: ACTG 5202 Study Population

Phase 3b RCT at 59 sites, US and Puerto Rico

	ABC/3TC	TDF/FTC		
N	928	929		
Male at birth %	81.4	84.0		
Age Group %				
$\leq 25$	10.1	10.5		
26-49	77.0	74.8		
≥ 50	12.8	14.6		
Baseline log <sub>10</sub> RNA copies/mL(med [IQR])	4.66 [4.31, 5.06]	4.65 [4.34, 4.96]		
Baseline CD4 count/mL (med [IQR])	229 [84, 338]	230 [97, 330]		

# Example: ACTG 5202 Reanalysis

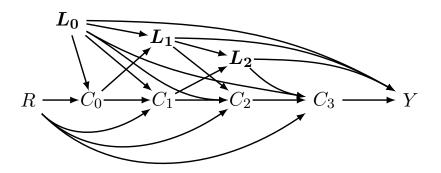
#### Objective:

- Estimand:  $E\left[Y^{r=1,\bar{a}=1}\right] E\left[Y^{r=0,\bar{a}=0}\right]$  at 48 weeks.
- Will vary protocol definition depending on the number of doses missed.

Outcomes: Composite virologic failure and all-cause mortality.

Methods: IPW analysis accounting for

- Age, Sex, Ethnicity
- Baseline CD4
- Baseline HIV RNA and screening RNA
- Time varying CD4
- Time varying HIV RNA



t: follow up time,  $\mathbf{L_t}$ : vector of covariates at follow up time t  $C_t$ : protocol deviation at follow up time t, R: randomization, Y: viral failure or death

#### Adherence and Protocol

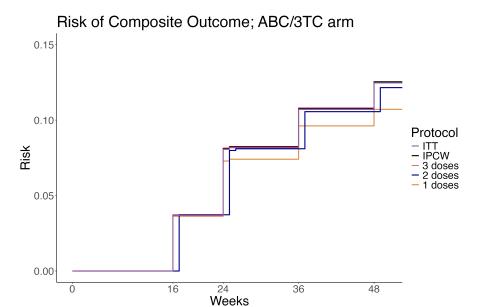
Adherence evaluated in-person at 8, 24, 48, 72, 96, then every 24 weeks and either at the final study evaluation or after virologic failure.

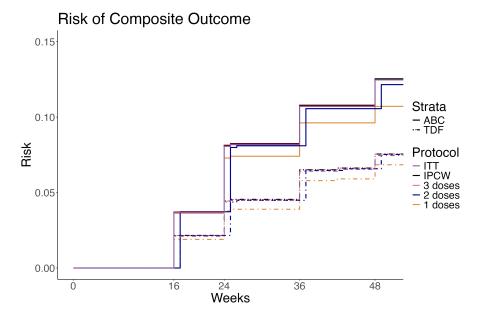
Last Time Missed Medication	How Close Was Dose Schedule Followed
Never	Never
>3 months ago	Some of the time
1-3 months ago	About half the time
2-4 weeks ago	Most of the time
1-2 weeks ago	All the time
Within the past week	

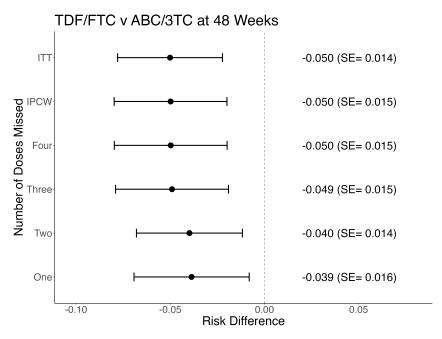
Protocol Definition	Description of Protocol
0 dose missed OK	No report of missed medication doses
1 dose missed OK	Participant with only one report of missed medication doses
:	
4 doses missed OK	Participant with $\geq 4$ reported missed medication doses without overlap in reported timing

### Deviation from Defined Protocols

	Doses Missed						
Treatment Group	LTFU	1	2	3	4	5	Total
ABC/3TC TDF/FTC	234 211		110 79		_	-	







#### Limitations and Future Plans

- 1. Completed with public access data.<sup>1</sup>
- 2. Reliance on coarse, self-reported medication adherence.
- 3. Assume identification conditions met.
  - Not guaranteed even though in a trial. Just like observational studies.
- 4. Future directions include repeating analysis with g-formula, considering additional protocols, and extending to different diseases.

<sup>&</sup>lt;sup>1</sup>Approved for more granular data from ACTG, awaiting dataset

## **Takeaways**

- Per protocol analysis should be treated like an observational analysis.
- Time varying covariates need to be taken into account.
- There are many ways protocols can be defined.
- The way protocol is defined can have meaningful impacts on estimates.

# Thank you!



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- Cole Lab Members



My website where you can find a link to my github.

#### Outcome Definition

- plasma HIV-1 RNA level ≥1000 copies /mL between 16 weeks and 24 weeks
- or ≥200 copies/mL at or after 24 weeks
- all cause mortality

# Censoring Risk

