

Toward Better Practice of Covariate Adjustment in Analyzing Randomized Clinical Trials

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HPTN
HIV Prevention
Trials Network

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- Covariate adjustment at the design and analysis stages
- Adjustment using linear working models
- Time-to-event outcomes
- Summary

Important: All the results in this talk hold **without assuming the models are correct!**

Why adjusting for covariates?

EMA (2015) guideline

*“Balance of treatment groups with respect to one or more specific prognostic covariates can **enhance the credibility** of the results of the trial”*

FDA (2021) guidance

*“Incorporating prognostic baseline factors in the primary statistical analysis of clinical trial data can result in a **more efficient** use of data to demonstrate and quantify the effects of treatment with minimal impact on bias or the Type I error rate”*

- At the design stage:
covariate-adaptive randomization
 - balance across baseline covariates to gain credibility and efficiency
- At the analysis stage:
model-assisted approach
 - more efficient use of data under the same assumption required by the unadjusted analysis

Design stage: covariate-adaptive randomization

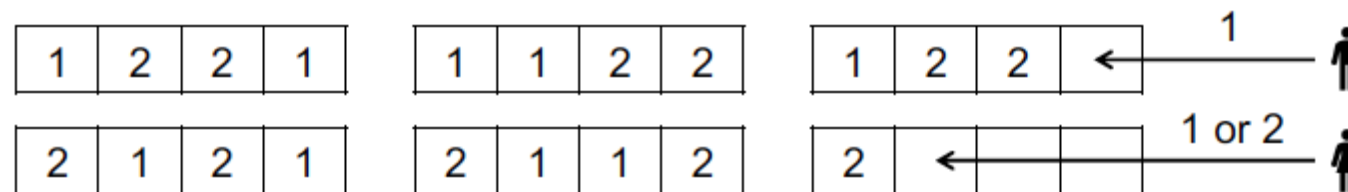
Simple Randomization (SR):

Treatment assignments are completely random

Covariate-Adaptative Randomization (CAR, also known as restricted randomization):

Balance treatment assignments across discrete baseline covariates (stratification variables)

- Example: Pocock-Simon's minimization, stratified urn design, stratified biased coin, stratified permuted-block randomization



Covariate adjustment in the analysis stage is a statistical method with high potential to **improve precision** for many trials

- **Pre-planned** adjustment for baseline variables when estimating the treatment effect
- **Target parameter** is the same as when using unadjusted method (e.g., difference in means)
- **Goal** is to avoid making any model assumption beyond what's assumed for the unadjusted method, i.e., **robustness to model misspecification** (FDA 2021)

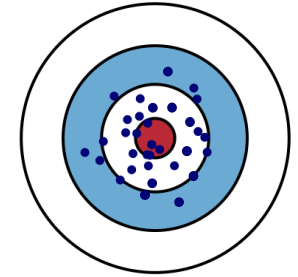
(e.g., Koch et al. 1998; Yang and Tsiatis, 2001; Rubin and van der Laan, 2008; Tsiatis, 2008; Lin, 2013; Bugni, 2018; Ye and Shao, 2020; Ye et al. 2022)

Example: analysis of covariance (ANCOVA)

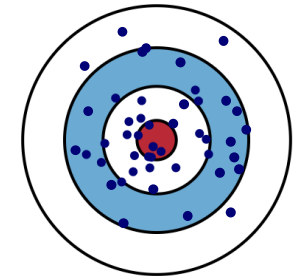
- ❑ Primary endpoint Y : continuous or binary
- ❑ Target parameter: $E(Y|A = 1) - E(Y|A = 0)$
- ❑ Estimator: $\hat{\theta}$ from fitting a linear model $E(Y|A, X) = \alpha + \theta A + \beta X$

- When the linear model is **incorrect**:
 - $\hat{\theta}$ still **correctly estimates** $E(Y|A = 1) - E(Y|A = 0)$
 - However, $\hat{\theta}$ can be **less precise** than simple mean difference $\bar{Y}_1 - \bar{Y}_0$
- Variance estimation should be **robust to model misspecification**
- Variance estimation should **account for CAR** (FDA, 2021)

Mean Difference



ANCOVA



Proposal: analysis of heterogeneous covariance (ANHECOVA)

- ❑ Primary endpoint Y : continuous or binary
- ❑ Target parameter: $E(Y|A = 1) - E(Y|A = 0)$
- ❑ Estimator: $\hat{\theta}$ from fitting a linear model $E(Y|A, X) = \alpha + \theta A + \beta X + \gamma A(X - \bar{X})$, with X including indicators of all strata used in CAR

➤ When the linear model is **incorrect**:

- $\hat{\theta}$ still **correctly estimates** $E(Y|A = 1) - E(Y|A = 0)$
- $\hat{\theta}$ is **never less precise and often more precise** than $\bar{Y}_1 - \bar{Y}_0$

Guaranteed Precision Gain

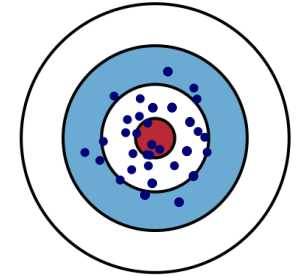
➤ Variance estimation is (not only) **robust to model misspecification**

Robust Variance Estimation

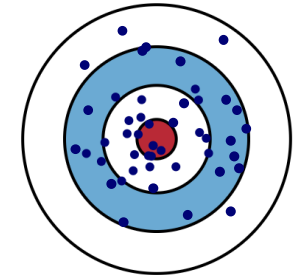
➤ Variance estimation is (but also) **robust to CAR**

Universal Applicability

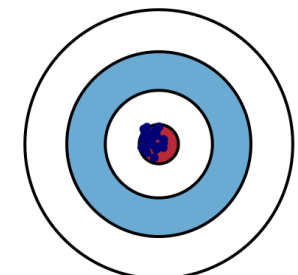
Mean Difference



ANCOVA

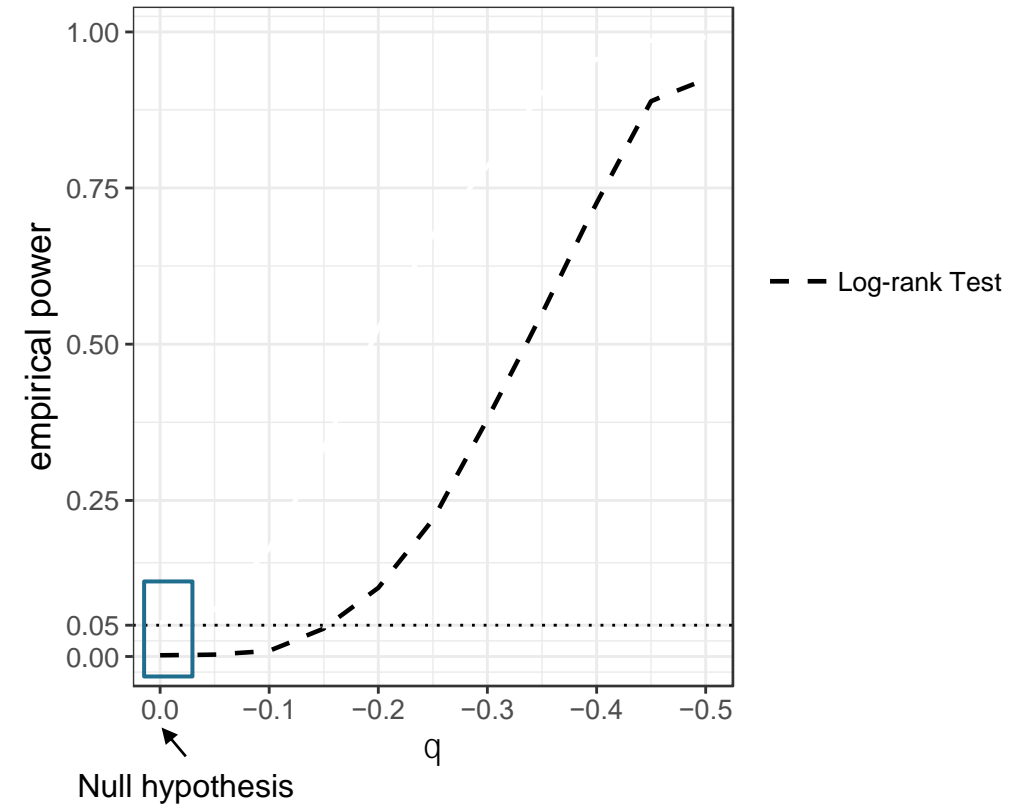


ANHECOVA



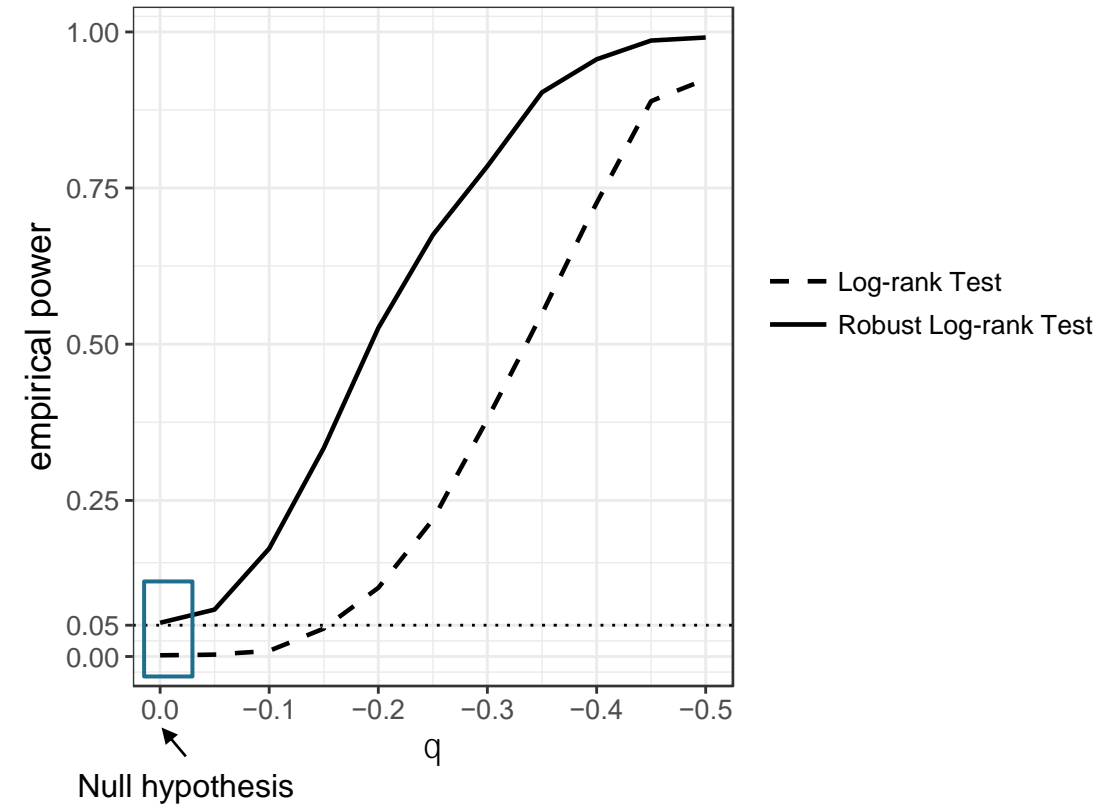
Time-to-event outcomes (Log-rank Test)

- ❑ Primary endpoint: time-to-event
 - ❑ Null Hypothesis $H_0: \lambda_0(t) = \lambda_1(t)$
 - ❑ Test statistics: **Log-rank test**
-
- Does NOT adjust for any covariate
 - **Conservative** under CAR



Time-to-event outcomes (Ye-Shao's Robust Log-rank Test)

- ❑ Primary endpoint: time-to-event
 - ❑ Null Hypothesis $H_0: \lambda_0(t) = \lambda_1(t)$
 - ❑ Test statistics: Ye-Shao's **Robust log-rank test**
-
- Does NOT adjust for any covariate
 - **Correct** under CAR
 - but NOT universally applicable

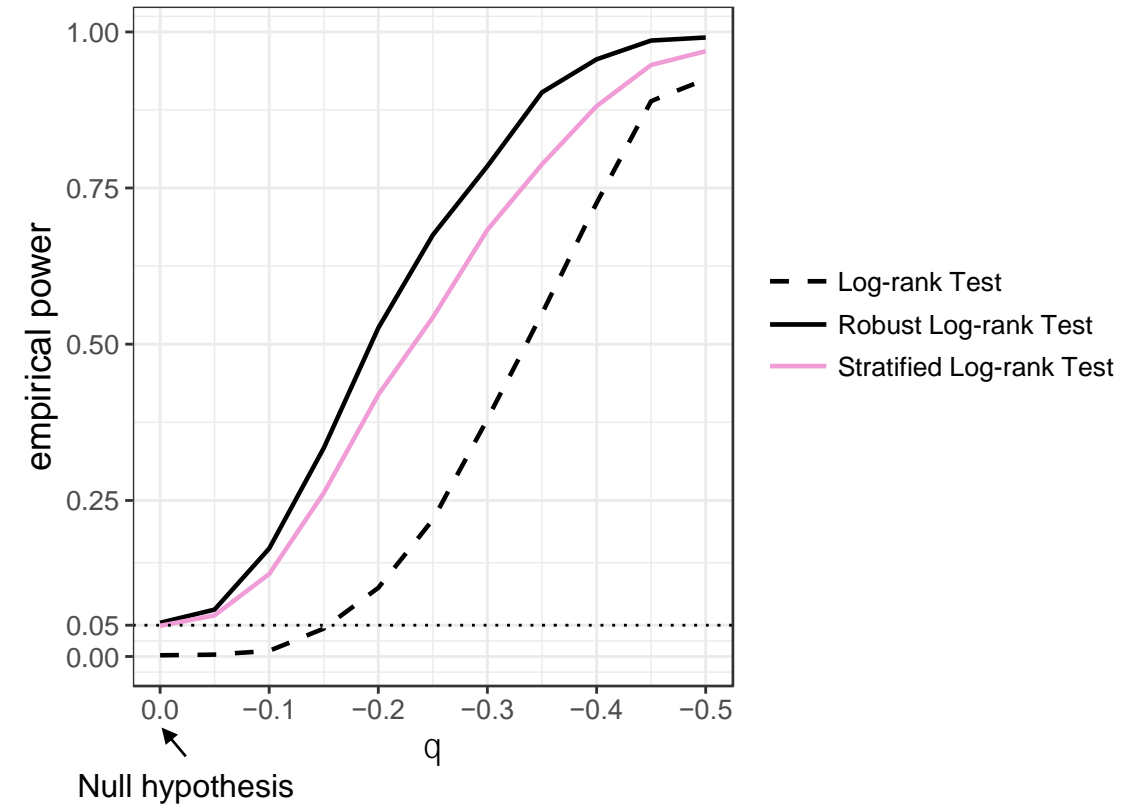


Time-to-event outcomes (Stratified Log-rank test)

- ❑ Primary endpoint: time-to-event
- ❑ Null Hypothesis $H_0: \lambda_{0z}(t) = \lambda_{1z}(t)$
- ❑ Test statistics: **Stratified Log-rank test**

- Adjust for discrete covariate
- **Correct** under CAR
 - **universally applicable**

- Can be **less powerful** than Ye-Shao's Robust Log-rank Test



Time-to-event outcomes (Covariate-adjusted Log-rank test)

- ❑ Primary endpoint: time-to-event
- ❑ Null Hypothesis $H_0: \lambda_0(t) = \lambda_1(t)$
- ❑ Test statistics: **Covariate-Adjusted Log-rank test**
(apply ANHECOVA to Log-rank test)

➤ Adjust for any covariate

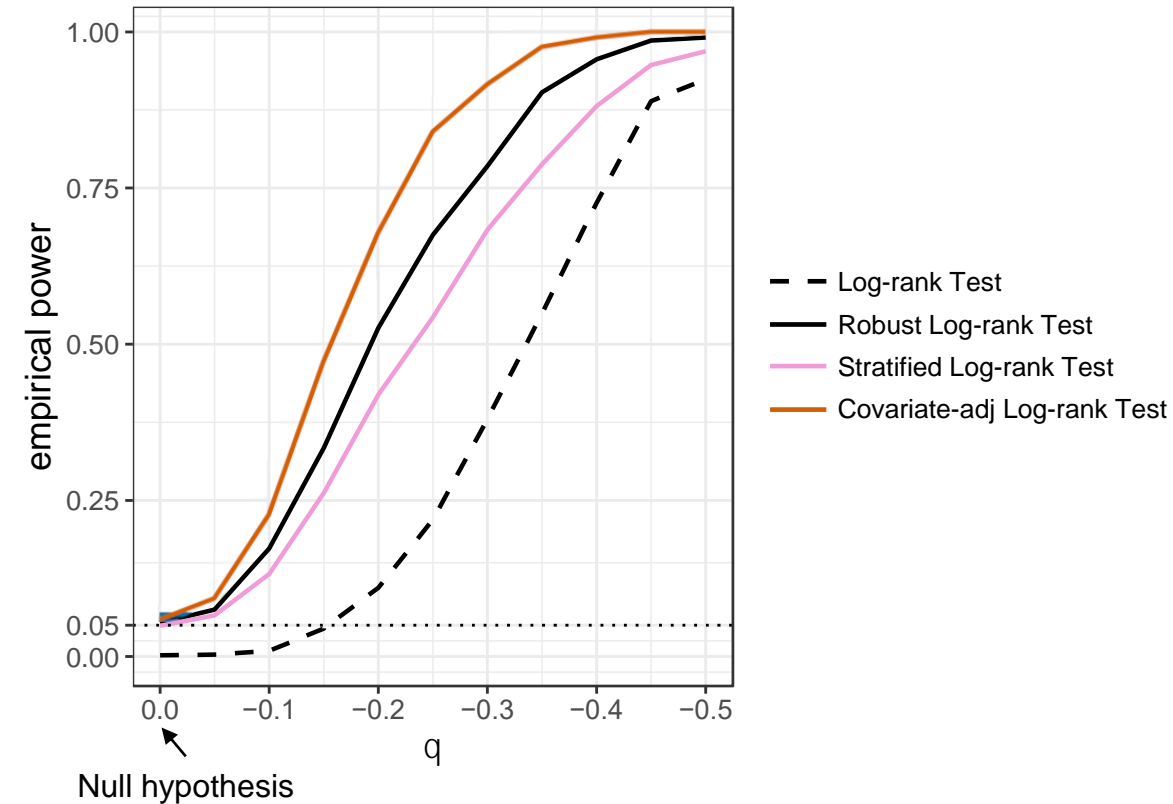
➤ **Correct** under CAR

- **universally applicable**

Universal Applicability

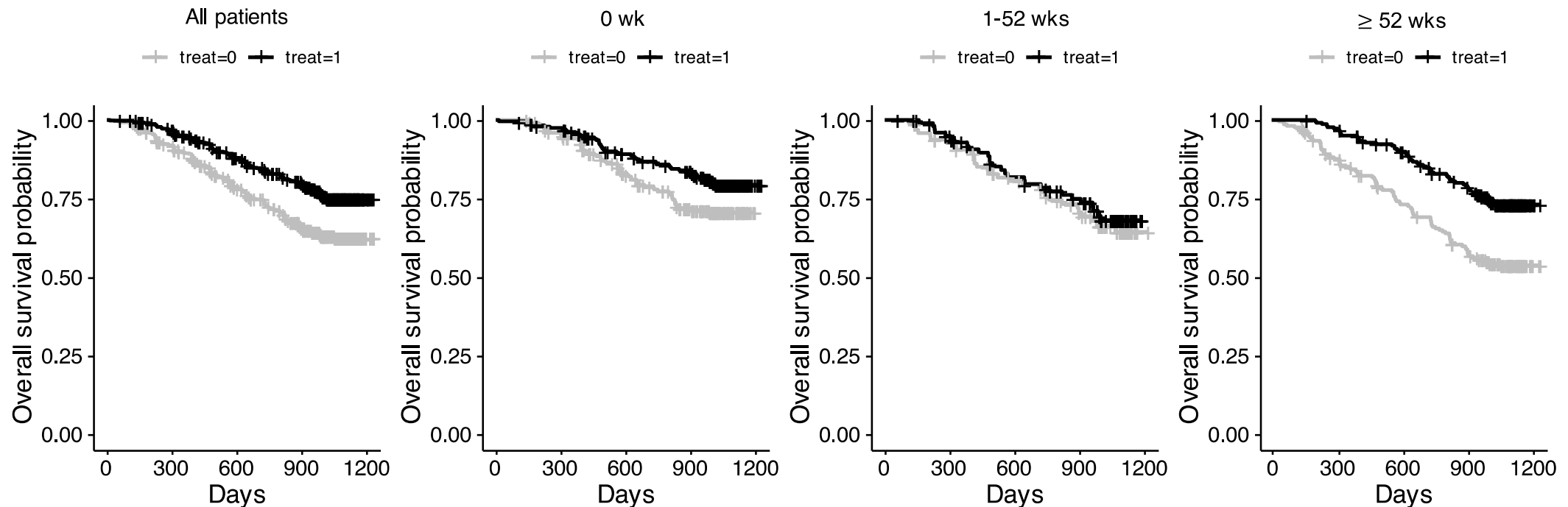
➤ **Never less powerful and often more powerful** than Ye-Shao's Robust Log-rank Test

Guaranteed Power Gain



Application to ACTG175 (Hammer et al. 1996)

- **Population:** Adults infected with HIV type 1 whose CD4 cell counts were 200-500 per cubic millimeter
- **Primary endpoint:** composite event ($\geq 50\%$ decline in CD4 cell count, an AIDS-defining event, or death)
- **Stratified permuted block randomization:** equal allocation and three strata: 0 week, 1-52 weeks, and ≥ 52 weeks of prior antiretroviral therapy.
- **Treatments:** Zidovudine (control) and Didanosine (treated).
- **Covariates for adjustment:** baseline CD4 cell count and number of days receiving prior antiretroviral therapy.



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	All patients	Sub-group		
		0 wk	1-52 wks	≥ 52 wks
Number of patients	1,093	461	198	434
Log-rank test	4.62	2.31	0.53	4.46
Two-sided p-value (adjusted for sub-group analysis)	<0.001	0.064	1	<0.001
Covariate-adjusted log-rank test	4.95	2.40	0.49	4.90
Two-sided p-value (adjusted for sub-group analysis)	<0.001	0.049	1	<0.001

Summary – ANHECOVA and three considerations

- ❖ **ANHECOVA: a general covariate adjustment strategy**
 - Adjust for **indicators of all strata used in CAR** and **all treatment-by-covariate interactions**
- 1. Guaranteed Precision Gain**
 - ANHECOVA does no harm
- 2. Robust Variance Estimation**
 - We recommend using variance estimators that are robust to model misspecification
- 3. Universal Applicability**
 - Variance estimator for ANHECOVA can be universally used under SR and all CAR

R package [RobinCar] is available on GitHub.

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