A Unified Framework for Rerandomization using Quadratic Forms

Design and Analysis of Experiments (2024)

Kyle Schindl¹ Zach Branson²

Department of Statistics & Data Science, Carnegie Mellon University

¹schindl@stat.cmu.edu

²zach@stat.cmu.edu

Introduction

• We focus on treatment vs. control randomized experiments

- We focus on treatment vs. control randomized experiments
- We are interested in estimating the Average Treatment Effect (ATE)

$$\tau = \mathbb{E}[Y(1) - Y(0)]$$

where Y(1) and Y(0) are the outcomes under treatment and control.

1

- We focus on treatment vs. control randomized experiments
- We are interested in estimating the Average Treatment Effect (ATE)

$$\tau = \mathbb{E}[Y(1) - Y(0)]$$

where Y(1) and Y(0) are the outcomes under treatment and control.

 In completely randomized experiments, estimates of the ATE are unbiased.

1

- We focus on treatment vs. control randomized experiments
- We are interested in estimating the Average Treatment Effect (ATE)

$$\tau = \mathbb{E}[Y(1) - Y(0)]$$

where Y(1) and Y(0) are the outcomes under treatment and control.

- In completely randomized experiments, estimates of the ATE are unbiased.
- Covariate distributions are "balanced" on average.

- We focus on treatment vs. control randomized experiments
- We are interested in estimating the Average Treatment Effect (ATE)

$$\tau = \mathbb{E}[Y(1) - Y(0)]$$

where Y(1) and Y(0) are the outcomes under treatment and control.

- In completely randomized experiments, estimates of the ATE are unbiased.
- Covariate distributions are "balanced" on average.
- We can do better than average.

- We focus on treatment vs. control randomized experiments
- We are interested in estimating the Average Treatment Effect (ATE)

$$\tau = \mathbb{E}[Y(1) - Y(0)]$$

where Y(1) and Y(0) are the outcomes under treatment and control.

- In completely randomized experiments, estimates of the ATE are unbiased.
- Covariate distributions are "balanced" on average.
- We can do better than average.
- By ensuring covariate distributions are closer, we can obtain more precise estimates of the ATE while maintaining unbiasedness.

1

 Covariate imbalances between treatment and control increases standard errors and reduces statistical power when estimating causal effects (Senn 1989; Lachin 1988; Lin, Zhu, and Su 2015; Branson, Li, and Ding 2023).

- Covariate imbalances between treatment and control increases standard errors and reduces statistical power when estimating causal effects (Senn 1989; Lachin 1988; Lin, Zhu, and Su 2015; Branson, Li, and Ding 2023).
- This is why classical experimental designs suggest blocking similar subjects together when randomizing.

- Covariate imbalances between treatment and control increases standard errors and reduces statistical power when estimating causal effects (Senn 1989; Lachin 1988; Lin, Zhu, and Su 2015; Branson, Li, and Ding 2023).
- This is why classical experimental designs suggest blocking similar subjects together when randomizing.
- Another method of reducing covariate imbalance is to rerandomize.

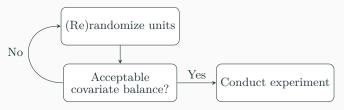


Figure 1: Procedure for implementing rerandomization

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

where

• $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)
 - Sequential designs (Zhou et al. 2018)

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)
 - Sequential designs (Zhou et al. 2018)
 - Clustered experiments (Lu et al. 2023)

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)
 - Sequential designs (Zhou et al. 2018)
 - Clustered experiments (Lu et al. 2023)
 - Bayesian designs (Liu et al. 2023)

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)
 - Sequential designs (Zhou et al. 2018)
 - Clustered experiments (Lu et al. 2023)
 - Bayesian designs (Liu et al. 2023)
- Most of these extensions utilize the Mahalanobis distance.

• (Morgan and Rubin 2012): Randomize until M < a, for some pre-specified a > 0, where

$$M := (\overline{X}_T - \overline{X}_C)^T \Sigma^{-1} (\overline{X}_T - \overline{X}_C).$$

- $\overline{X}_T, \overline{X}_C \in \mathbb{R}^d$ are the covariate means in treatment and control.
- $\Sigma := \operatorname{Cov}(\overline{X}_T \overline{X}_C \mid x)$
- There have been many extensions to other design settings such as
 - 2^K factorial designs (Branson, Dasgupta, and Rubin 2016; Li, Ding, and Rubin 2020)
 - Sequential designs (Zhou et al. 2018)
 - Clustered experiments (Lu et al. 2023)
 - Bayesian designs (Liu et al. 2023)
- Most of these extensions utilize the Mahalanobis distance.
- We'll consider: What distances are optimal for rerandomization?

• As long as $\sum_{i=1}^{n} W_i = \sum_{i=1}^{n} (1 - W_i)$, where $W_i \in \{0, 1\}$ is the treatment assignment, then $\hat{\tau}$ remains unbiased.

- As long as $\sum_{i=1}^{n} W_i = \sum_{i=1}^{n} (1 W_i)$, where $W_i \in \{0, 1\}$ is the treatment assignment, then $\hat{\tau}$ remains unbiased.
- If $\overline{X}_T \overline{X}_C \mid x \sim \mathcal{N}(0, \Sigma)$ then, Mahalanobis Rerandomization applies an equal-percentage variance reduction to all covariates,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, M \le a) = v_a Cov(\overline{X}_T - \overline{X}_C \mid x)$$

where
$$v_a = \mathbb{P}(\chi_{d+2}^2 \le a) / \mathbb{P}(\chi_d^2 \le a) \le 1$$
.

- As long as $\sum_{i=1}^{n} W_i = \sum_{i=1}^{n} (1 W_i)$, where $W_i \in \{0, 1\}$ is the treatment assignment, then $\hat{\tau}$ remains unbiased.
- If $\overline{X}_T \overline{X}_C \mid x \sim \mathcal{N}(0, \Sigma)$ then, Mahalanobis Rerandomization applies an equal-percentage variance reduction to all covariates,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, M \le a) = v_a Cov(\overline{X}_T - \overline{X}_C \mid x)$$

where $v_a = \mathbb{P}(\chi_{d+2}^2 \leq a) / \mathbb{P}(\chi_d^2 \leq a) \leq 1$.

 When the treatment effect is additive, this leads to a variance reduction of

$$\frac{\mathbb{V}(\hat{\tau} \mid x, M < a)}{\mathbb{V}(\hat{\tau} \mid x)} = 1 - (1 - v_a)R^2 \le 1$$

where

$$\hat{\tau} = \frac{1}{n_1} \sum_{i=1}^{n} Y_i(1) W_i - \frac{1}{n_0} \sum_{i=1}^{n} Y_i(0) (1 - W_i)$$

4

- As long as $\sum_{i=1}^{n} W_i = \sum_{i=1}^{n} (1 W_i)$, where $W_i \in \{0, 1\}$ is the treatment assignment, then $\hat{\tau}$ remains unbiased.
- If $\overline{X}_T \overline{X}_C \mid x \sim \mathcal{N}(0, \Sigma)$ then, Mahalanobis Rerandomization applies an equal-percentage variance reduction to all covariates,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, M \le a) = v_a Cov(\overline{X}_T - \overline{X}_C \mid x)$$

where $v_a = \mathbb{P}(\chi_{d+2}^2 \leq a) / \mathbb{P}(\chi_d^2 \leq a) \leq 1$.

 When the treatment effect is additive, this leads to a variance reduction of

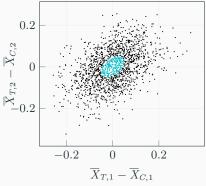
$$\frac{\mathbb{V}(\hat{\tau} \mid x, M < a)}{\mathbb{V}(\hat{\tau} \mid x)} = 1 - (1 - v_a)R^2 \le 1$$

where

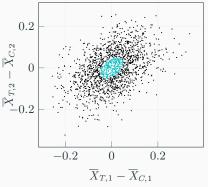
$$\hat{\tau} = \frac{1}{n_1} \sum_{i=1}^{n} Y_i(1) W_i - \frac{1}{n_0} \sum_{i=1}^{n} Y_i(0) (1 - W_i)$$

 Asymptotically, this is true with unequal sample sizes and non-additivity (Li, Ding, and Rubin 2018).

• Geometrically, we can see that Mahalanobis Rerandomization preserves the "shape" of the covariance matrix but reduces the scale.

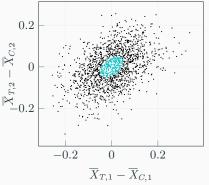


 Geometrically, we can see that Mahalanobis Rerandomization preserves the "shape" of the covariance matrix but reduces the scale.



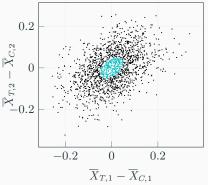
• Each eigenvector of the covariance matrix Σ is scaled by v_a .

 Geometrically, we can see that Mahalanobis Rerandomization preserves the "shape" of the covariance matrix but reduces the scale.



- Each eigenvector of the covariance matrix Σ is scaled by v_a .
- Placing equal priority on all covariates create challenges since $v_a \to 1$ as $d \to \infty$ (no variance reduction).

 Geometrically, we can see that Mahalanobis Rerandomization preserves the "shape" of the covariance matrix but reduces the scale.



- Each eigenvector of the covariance matrix Σ is scaled by v_a .
- Placing equal priority on all covariates create challenges since $v_a \to 1$ as $d \to \infty$ (no variance reduction).
- Mahalanobis Rerandomization may be "wasting" variance reduction on unimportant covariates

Balancing Metrics as Quadratic Forms

 $\bullet\,$ Other recent proposed balance metrics include:

Balancing Metrics as Quadratic Forms

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T (\Sigma + \lambda I_d)^{-1} (\overline{X}_T - \overline{X}_C)$$

Balancing Metrics as Quadratic Forms

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T (\Sigma + \lambda I_d)^{-1} (\overline{X}_T - \overline{X}_C)$$

• The Mahalanobis distance after dropping the bottom d-k principal components (Zhang, Yin, and Rubin 2023):

$$M_k = (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})^T \Sigma_Z^{-1} (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})$$

where Z are the principal components.

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T (\Sigma + \lambda I_d)^{-1} (\overline{X}_T - \overline{X}_C)$$

• The Mahalanobis distance after dropping the bottom d-k principal components (Zhang, Yin, and Rubin 2023):

$$M_k = (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})^T \Sigma_Z^{-1} (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})$$

where Z are the principal components.

• Each method can be written as $(\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$ for some $A \in \mathbb{R}^{d \times d}$, e.g.,

$$M = (\overline{X}_T - \overline{X}_C)^T \underbrace{\Sigma^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

6

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T \underbrace{(\Sigma + \lambda I_d)^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

• The Mahalanobis distance after dropping the bottom d-k principal components (Zhang, Yin, and Rubin 2023):

$$M_k = (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})^T \Sigma_Z^{-1} (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})$$

where Z are the principal components.

• Each method can be written as $(\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$ for some $A \in \mathbb{R}^{d \times d}$, e.g.,

$$M = (\overline{X}_T - \overline{X}_C)^T \underbrace{\Sigma^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T \underbrace{(\Sigma + \lambda I_d)^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

• The Mahalanobis distance after dropping the bottom d-k principal components (Zhang, Yin, and Rubin 2023):

$$M_k = (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})^T \Sigma_Z^{-1} (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})$$
$$= (\overline{X}_T - \overline{X}_C)^T \underbrace{V \begin{pmatrix} D_k & 0 \\ 0 & 0 \end{pmatrix} V^T (\overline{X}_T - \overline{X}_C)}_{A}$$

where D_k are the singular values from $x = UDV^T$.

• Each method can be written as $(\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$ for some $A \in \mathbb{R}^{d \times d}$, e.g.,

$$M = (\overline{X}_T - \overline{X}_C)^T \underbrace{\Sigma^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

- Other recent proposed balance metrics include:
- A penalized Mahalanobis distance (Branson and Shao 2021):

$$M_{\lambda} = (\overline{X}_T - \overline{X}_C)^T \underbrace{(\Sigma + \lambda I_d)^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

• The Mahalanobis distance after dropping the bottom d-k principal components (Zhang, Yin, and Rubin 2023):

$$M_k = (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})^T \Sigma_Z^{-1} (\overline{Z}_T^{(k)} - \overline{Z}_C^{(k)})$$
$$= (\overline{X}_T - \overline{X}_C)^T \underbrace{V \begin{pmatrix} D_k & 0 \\ 0 & 0 \end{pmatrix} V^T (\overline{X}_T - \overline{X}_C)}_{A}$$

where D_k are the singular values from $x = UDV^T$.

• Each method can be written as $(\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$ for some $A \in \mathbb{R}^{d \times d}$, e.g.,

$$M = (\overline{X}_T - \overline{X}_C)^T \underbrace{\Sigma^{-1}}_{A} (\overline{X}_T - \overline{X}_C)$$

This leads to the question: what is a good choice of A?

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

• We select a by setting an acceptance probability α such that

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

• Some key results we cover:

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

- Some key results we cover:
 - The covariance of $\overline{X}_T \overline{X}_C$ conditioning on any quadratic form.

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

- Some key results we cover:
 - The covariance of $\overline{X}_T \overline{X}_C$ conditioning on any quadratic form.
 - The optimal choice of A for covariate balance

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

- Some key results we cover:
 - The covariance of $\overline{X}_T \overline{X}_C$ conditioning on any quadratic form.
 - The optimal choice of A for covariate balance
 - The optimal choice of A for reducing the variance of $\hat{\tau}$

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

- Some key results we cover:
 - The covariance of $\overline{X}_T \overline{X}_C$ conditioning on any quadratic form.
 - The optimal choice of A for covariate balance
 - The optimal choice of A for reducing the variance of $\hat{\tau}$
- Throughout, we assume A is positive-definite and that $\overline{X}_T \overline{X}_C \mid x \sim \mathcal{N}(0, \Sigma)$.

• In this work, we establish a general framework for randomizing until $Q_A(x) \leq a$ where

$$Q_A(x) := (\overline{X}_T - \overline{X}_C)^T A(\overline{X}_T - \overline{X}_C)$$

$$\mathbb{P}(Q_A(x) \le a) = \alpha$$

- Some key results we cover:
 - The covariance of $\overline{X}_T \overline{X}_C$ conditioning on any quadratic form.
 - The optimal choice of A for covariate balance
 - The optimal choice of A for reducing the variance of $\hat{\tau}$
- Throughout, we assume A is positive-definite and that $\overline{X}_T \overline{X}_C \mid x \sim \mathcal{N}(0, \Sigma)$.
- Also note that we assume the covariates have been standardized.

Theorem 1 (Covariance Reduction)

Suppose that $\Sigma^{1/2}A\Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma(diag\{(q_{j,\eta})_{1 \le j \le d}\}) \Gamma^T \Sigma^{1/2}$$

Theorem 1 (Covariance Reduction)

Suppose that $\sum^{1/2} A \sum^{1/2}$ and \sum share an eigenbasis. Then,

$$Cov\left(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \leq a\right) = \Sigma^{1/2} \Gamma\left(diag\{(q_{j,\eta})_{1 \leq j \leq d}\}\right) \Gamma^T \Sigma^{1/2}$$

• $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ

Theorem 1 (Covariance Reduction)

Suppose that $\Sigma^{1/2}A\Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma(diag\{(q_{j,\eta})_{1 \le j \le d}\}) \Gamma^T \Sigma^{1/2}$$

- $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ
- $q_{i,n}$ is defined as

$$q_{j,\eta} = \mathbb{E}\left[\mathcal{Z}_j^2 \mid x, \sum_{j=1}^d \eta_j \mathcal{Z}_j^2 \le a\right]$$

Theorem 1 (Covariance Reduction)

Suppose that $\hat{\Sigma}^{1/2}A\Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma\left(diag\{(q_{j,\eta})_{1 \le j \le d}\}\right) \Gamma^T \Sigma^{1/2}$$

- $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ
- $q_{j,\eta}$ is defined as

$$q_{j,\eta} = \mathbb{E}\left[\mathcal{Z}_j^2 \mid x, \sum_{j=1}^d \eta_j \mathcal{Z}_j^2 \le a\right]$$

• $\eta_1 \geq \cdots \geq \eta_d$ are the eigenvalues of $\Sigma^{1/2} A \Sigma^{1/2}$

Theorem 1 (Covariance Reduction)

Suppose that $\Sigma^{1/2}A\Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma(diag\{(q_{j,\eta})_{1 \le j \le d}\}) \Gamma^T \Sigma^{1/2}$$

- $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ
- $q_{i,\eta}$ is defined as

$$q_{j,\eta} = \mathbb{E}\left[\mathcal{Z}_j^2 \mid x, \sum_{j=1}^d \eta_j \mathcal{Z}_j^2 \le a\right]$$

- $\eta_1 \geq \cdots \geq \eta_d$ are the eigenvalues of $\Sigma^{1/2} A \Sigma^{1/2}$
- $\mathcal{Z}_1, \ldots, \mathcal{Z}_d \sim \mathcal{N}(0,1)$

Theorem 1 (Covariance Reduction)

Suppose that $\Sigma^{1/2}A\Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma(diag\{(q_{j,\eta})_{1 \le j \le d}\}) \Gamma^T \Sigma^{1/2}$$

- $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ
- $q_{j,\eta}$ is defined as

$$q_{j,\eta} = \mathbb{E}\left[\mathcal{Z}_j^2 \mid x, \sum_{j=1}^d \eta_j \mathcal{Z}_j^2 \le a\right]$$

- $\eta_1 \geq \cdots \geq \eta_d$ are the eigenvalues of $\Sigma^{1/2} A \Sigma^{1/2}$
- $\mathcal{Z}_1,\ldots,\mathcal{Z}_d \sim \mathcal{N}(0,1)$
- $q_{1,\eta} \leq \cdots \leq q_{d,\eta} \leq 1$.

Theorem 1 (Covariance Reduction)

Suppose that $\hat{\Sigma}^{1/2} A \Sigma^{1/2}$ and Σ share an eigenbasis. Then,

$$Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a) = \Sigma^{1/2} \Gamma(diag\{(q_{j,\eta})_{1 \le j \le d}\}) \Gamma^T \Sigma^{1/2}$$

- $\Gamma \in \mathbb{R}^{d \times d}$ is the orthogonal matrix of eigenvectors of Σ
- $q_{i,\eta}$ is defined as

$$q_{j,\eta} = \mathbb{E}\left[\mathcal{Z}_j^2 \mid x, \sum_{j=1}^d \eta_j \mathcal{Z}_j^2 \le a\right]$$
$$= \frac{p_d}{\eta_j} \det(\Sigma^{1/2} A \Sigma^{1/2})^{1/d} \alpha^{2/d} + o(\alpha^{2/d})$$

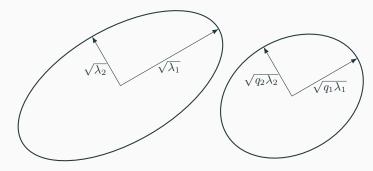
for a sufficiently small α , where $p_d = \frac{2\pi}{d+2} \left(\frac{2\pi^{d/2}}{d\Gamma(d/2)} \right)^{-2/d}$ (Lu et al. 2023).

- $\eta_1 \ge \cdots \ge \eta_d$ are the eigenvalues of $\Sigma^{1/2} A \Sigma^{1/2}$
- $\mathcal{Z}_1, \ldots, \mathcal{Z}_d \sim \mathcal{N}(0, 1)$
- $q_{1,\eta} \leq \cdots \leq q_{d,\eta} \leq 1$.

Geometric Intuition behind Quadratic Form Rerandomization

Geometric Intuition behind Quadratic Form Rerandomization

• The eigenvalues of Cov $(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \leq a)$ are given by $q_{1,\eta}\lambda_1, \ldots, q_{d,\eta}\lambda_d$.



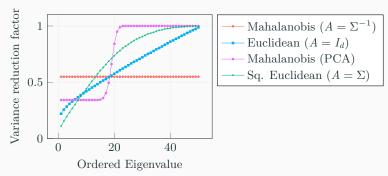
(a) Covariance Matrix before rerandomization

(b) Covariance Matrix after rerandomization

Variance Reduction Factors Visualized

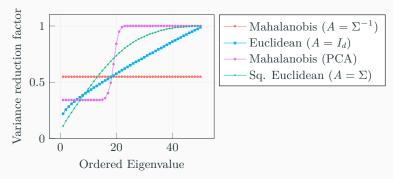
Variance Reduction Factors Visualized

• Depending on the choice of A, there is a different amount of variance reduction applied to each eigenvalue.



Variance Reduction Factors Visualized

• Depending on the choice of A, there is a different amount of variance reduction applied to each eigenvalue.



• We show that different choices of A can lead to significantly different precision levels when estimating $\hat{\tau}$.

Variance Reduction Visualized

Variance Reduction Visualized

• Depending on the relationship between the potential outcomes and x, and the eigenstructure of Σ , different choices of A yield different levels of precision.

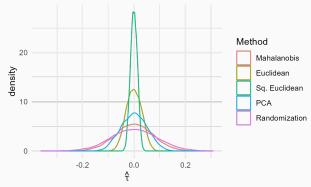


Figure 3: Distribution of $\hat{\tau}$

Variance Reduction Visualized

• Depending on the relationship between the potential outcomes and x, and the eigenstructure of Σ , different choices of A yield different levels of precision.

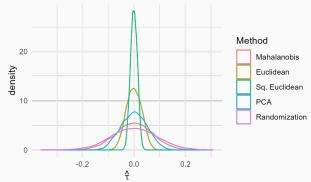


Figure 3: Distribution of $\hat{\tau}$

• Given the clear differences between each of these methods, the question arises: which choice of A is optimal?

• First we consider optimality in terms of covariate balance.

- First we consider optimality in terms of covariate balance.
- $\bullet~$ We consider two targets for quantifying covariate balance:

- First we consider optimality in terms of covariate balance.
- We consider two targets for quantifying covariate balance:
 - The choice of A that maximizes the variance reduction applied across all covariates,

$$\max_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} (1 - q_{j,\eta}) = \min_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} q_{j,\eta}$$

- First we consider optimality in terms of covariate balance.
- We consider two targets for quantifying covariate balance:
 - The choice of A that maximizes the variance reduction applied across all covariates,

$$\max_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} (1 - q_{j,\eta}) = \min_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} q_{j,\eta}$$

 The choice of A that constrains the covariance matrix after rerandomization the most, which we define as

$$\min_{A \in \mathbb{R}^{d \times d}} || \operatorname{Cov} \left(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a \right) ||_F^2$$

where $||\cdot||_F^2 = \sum_{j=1}^d \sigma_j(\cdot)$ and $\sigma_j(\cdot)$ are the singular values of the matrix.

- First we consider optimality in terms of covariate balance.
- We consider two targets for quantifying covariate balance:
 - The choice of A that maximizes the variance reduction applied across all covariates,

$$\max_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} (1 - q_{j,\eta}) = \min_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} q_{j,\eta}$$

• The choice of A that constrains the covariance matrix after rerandomization the most, which we define as

$$\min_{A \in \mathbb{R}^{d \times d}} || \operatorname{Cov} \left(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a \right) ||_F^2$$

where $||\cdot||_F^2 = \sum_{j=1}^d \sigma_j(\cdot)$ and $\sigma_j(\cdot)$ are the singular values of the matrix.

• Equivalently, $\sigma_j^2(\cdot)$ are the eigenvalues of the sample covariance matrix.

- First we consider optimality in terms of covariate balance.
- We consider two targets for quantifying covariate balance:
 - The choice of A that maximizes the variance reduction applied across all covariates,

$$\max_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} (1 - q_{j,\eta}) = \min_{A \in \mathbb{R}^{d \times d}} \sum_{j=1}^{d} q_{j,\eta}$$

• The choice of A that constrains the covariance matrix after rerandomization the most, which we define as

$$\min_{A \in \mathbb{R}^{d \times d}} || \operatorname{Cov} \left(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a \right) ||_F^2$$

where $||\cdot||_F^2 = \sum_{j=1}^d \sigma_j(\cdot)$ and $\sigma_j(\cdot)$ are the singular values of the matrix.

- Equivalently, $\sigma_j^2(\cdot)$ are the eigenvalues of the sample covariance matrix.
- Later, we investigate how this impacts the precision of $\hat{\tau}$.

Theorem 2 (Covariate Balance)

For all positive-definite matrices $A \in \mathbb{R}^{d \times d}$

• $A = I_d$ minimizes the Frobenius norm:

$$||\operatorname{Cov}(\overline{X}_T - \overline{X}_C \mid x, Q_I(x) \leq a)||_F \leq ||\operatorname{Cov}(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \leq a)||_F + o(\alpha^{2/d})$$

• $A = \Sigma^{-1}$ maximizes the total variance reduction applied

$$\sum_{j=1}^{d} v_a \le \sum_{j=1}^{d} q_{j,\eta} + o(\alpha^{2/d})$$

Theorem 2 (Covariate Balance)

For all positive-definite matrices $A \in \mathbb{R}^{d \times d}$

• $A = I_d$ minimizes the Frobenius norm:

$$||\operatorname{Cov}(\overline{X}_T - \overline{X}_C \mid x, Q_I(x) \leq a)||_F \leq ||\operatorname{Cov}(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \leq a)||_F + o(\alpha^{2/d})$$

• $A = \Sigma^{-1}$ maximizes the total variance reduction applied

$$\sum_{j=1}^{d} v_a \le \sum_{j=1}^{d} q_{j,\eta} + o(\alpha^{2/d})$$

 These two methods represent opposite sides of the rerandomization spectrum:

Theorem 2 (Covariate Balance)

For all positive-definite matrices $A \in \mathbb{R}^{d \times d}$

• $A = I_d$ minimizes the Frobenius norm:

$$||Cov(\overline{X}_T - \overline{X}_C \mid x, Q_I(x) \le a)||_F \le ||Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a)||_F + o(\alpha^{2/d})$$

• $A = \Sigma^{-1}$ maximizes the total variance reduction applied

$$\sum_{j=1}^{d} v_a \le \sum_{j=1}^{d} q_{j,\eta} + o(\alpha^{2/d})$$

- These two methods represent opposite sides of the rerandomization spectrum:
 - Mahalanobis Rerandomization $(A = \Sigma^{-1})$ leaves the shape of the covariance matrix unchanged by applying an equal variance reduction to all eigenvectors.

Theorem 2 (Covariate Balance)

For all positive-definite matrices $A \in \mathbb{R}^{d \times d}$

• $A = I_d$ minimizes the Frobenius norm:

$$||Cov(\overline{X}_T - \overline{X}_C \mid x, Q_I(x) \le a)||_F \le ||Cov(\overline{X}_T - \overline{X}_C \mid x, Q_A(x) \le a)||_F + o(\alpha^{2/d})$$

• $A = \Sigma^{-1}$ maximizes the total variance reduction applied

$$\sum_{j=1}^{d} v_a \le \sum_{j=1}^{d} q_{j,\eta} + o(\alpha^{2/d})$$

- These two methods represent opposite sides of the rerandomization spectrum:
 - Mahalanobis Rerandomization (A = Σ⁻¹) leaves the shape of the covariance matrix unchanged by applying an equal variance reduction to all eigenvectors.
 - Euclidean Rerandomization ($A=I_d$) scales each eigenvector by a different factor such that their magnitude is the same.

Quantifying the variance reduction to $\hat{\tau}$

Quantifying the variance reduction to $\hat{\tau}$

• When the treatment effect is additive, it follows that

$$Y_i(W_i) = \beta_0 + Z_i \beta_Z + \tau W_i + \varepsilon_i$$

where $\beta_0 + Z_i\beta_Z$ represent the linear projection of Y_i onto the principal components Z = XV and ε_i is a residual that encompasses deviations from the projection.

Quantifying the variance reduction to $\hat{\tau}$

• When the treatment effect is additive, it follows that

$$Y_i(W_i) = \beta_0 + Z_i\beta_Z + \tau W_i + \varepsilon_i$$

where $\beta_0 + Z_i\beta_Z$ represent the linear projection of Y_i onto the principal components Z = XV and ε_i is a residual that encompasses deviations from the projection.

Theorem 3 (Variance Reduction)

Suppose the treatment effect is additive. Then,

$$\mathbb{V}(\widehat{\tau} \mid x) - \mathbb{V}(\widehat{\tau} \mid x, Q_A(x) \le a) = \sum_{j=1}^d \beta_{Z,j}^2 \lambda_j (1 - q_{j,\eta}) \ge 0$$

where $\lambda_1, \ldots, \lambda_d$ are the eigenvalues of Σ and $\hat{\tau}$ is the mean-difference estimator.

The Optimal Choice of A for variance reduction to $\hat{\tau}$

The Optimal Choice of A for variance reduction to $\hat{\tau}$

Theorem 4 (Optimal A) Suppose that the treatment effect is additive. Then for all positive-definite matrices $A \in \mathbb{R}^{d \times d}$,

$$A^* = \Gamma \begin{pmatrix} \beta_{Z,1}^2 & 0 \\ & \ddots & \\ 0 & & \beta_{Z,d}^2 \end{pmatrix} \Gamma^T$$

minimizes the variance of the mean-differences estimator, i.e.,

$$\mathbb{V}(\widehat{\tau} \mid x, Q_{A^*}(x) \le a) \le \mathbb{V}(\widehat{\tau} \mid x, Q_A(x) \le a) + o(\alpha^{2/d})$$

where Γ is the orthogonal matrix of eigenvectors of Σ .

• For any choice of A there exists some β_Z such that A is optimal.

- For any choice of A there exists some β_Z such that A is optimal.
- For example:
 - When $\beta_{Z,j}^2=\frac{1}{\lambda_j},$ $A=\Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.

- For any choice of A there exists some β_Z such that A is optimal.
- For example:
 - When $\beta_{Z,j}^2 = \frac{1}{\lambda_i}$, $A = \Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = 1$, $A = I_d$ (Euclidean Rerandomization) is optimal.

- For any choice of A there exists some β_Z such that A is optimal.
- · For example:
 - When $\beta_{Z,j}^2 = \frac{1}{\lambda_i}$, $A = \Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = 1$, $A = I_d$ (Euclidean Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = \lambda_j$, the choice $A = \Sigma$ is optimal.

- For any choice of A there exists some β_Z such that A is optimal.
- For example:
 - When $\beta_{Z,j}^2 = \frac{1}{\lambda_j}$, $A = \Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = 1$, $A = I_d$ (Euclidean Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = \lambda_j$, the choice $A = \Sigma$ is optimal.
- When the top eigenvectors and potential outcomes are closely related, choices like $A = I_d, \Sigma$ or PCA rerandomization tend to be better

- For any choice of A there exists some β_Z such that A is optimal.
- For example:
 - When $\beta_{Z,j}^2 = \frac{1}{\lambda_j}$, $A = \Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = 1$, $A = I_d$ (Euclidean Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = \lambda_j$, the choice $A = \Sigma$ is optimal.
- When the top eigenvectors and potential outcomes are closely related, choices like $A = I_d, \Sigma$ or PCA rerandomization tend to be better
- When the bottom eigenvectors and potential outcomes are closely related, methods like $A = \Sigma^{-1}$ are better.

- For any choice of A there exists some β_Z such that A is optimal.
- For example:
 - When $\beta_{Z,j}^2 = \frac{1}{\lambda_i}$, $A = \Sigma^{-1}$ (Mahalanobis Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = 1$, $A = I_d$ (Euclidean Rerandomization) is optimal.
 - When $\beta_{Z,j}^2 = \lambda_j$, the choice $A = \Sigma$ is optimal.
- When the top eigenvectors and potential outcomes are closely related, choices like $A = I_d, \Sigma$ or PCA rerandomization tend to be better
- When the bottom eigenvectors and potential outcomes are closely related, methods like $A = \Sigma^{-1}$ are better.
- However, different choices may be less risky than others.

The minimax optimal choice of A

Theorem 5 (Minimax A)

The choice $A = I_d$ minimizes the maximum difference between the optimal quadratic form and any other choice of A, given by

$$\min_{A \in \mathbb{R}^{d \times d} \mid |\beta_Z| \mid_2 \le c} \max_{|A| \in \mathbb{R}^{d \times d}} |\mathbb{V}(\widehat{\tau} \mid x, Q_{A^*}(x) \le a) - \mathbb{V}(\widehat{\tau} \mid x, Q_A(x) \le a)|.$$

The minimax optimal choice of A

Theorem 5 (Minimax A)

The choice $A = I_d$ minimizes the maximum difference between the optimal quadratic form and any other choice of A, given by

$$\min_{A \in \mathbb{R}^{d \times d} ||\beta_Z||_2 \le c} \max_{2 \le c} |\mathbb{V}(\widehat{\tau} \mid x, Q_{A^*}(x) \le a) - \mathbb{V}(\widehat{\tau} \mid x, Q_A(x) \le a)|.$$

• Under $A = I_d$, the precision of $\hat{\tau}$ is never too far from the optimal choice.

The minimax optimal choice of A

Theorem 5 (Minimax A)

The choice $A = I_d$ minimizes the maximum difference between the optimal quadratic form and any other choice of A, given by

$$\min_{A \in \mathbb{R}^{d \times d} ||\beta_{\mathcal{I}}||_2 \le c} \max |\mathbb{V}(\widehat{\tau} \mid x, Q_{A^*}(x) \le a) - \mathbb{V}(\widehat{\tau} \mid x, Q_A(x) \le a)|.$$

- Under $A = I_d$, the precision of $\hat{\tau}$ is never too far from the optimal choice.
- For intuition, compare the variance reduction factors between the two:

$$q_{j,\lambda} = \mathbb{E}\left[\mathcal{Z}_{j}^{2} \mid x, \sum_{j=1}^{d} \lambda_{j} \mathcal{Z}_{j}^{2} \leq a\right]$$
$$q_{j,\beta\lambda}^{*} = \mathbb{E}\left[\mathcal{Z}_{j}^{2} \mid x, \sum_{j=1}^{d} \beta_{Z,j}^{2} \lambda_{j} \mathcal{Z}_{j}^{2} \leq a\right]$$

• Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.

- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{j=1}^{d} \lambda_j = d = \operatorname{tr}(\Sigma)$

- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{i=1}^{d} \lambda_i = d = \operatorname{tr}(\Sigma)$
- We simulate n = 500 observations where 250 units are sent to treatment and 250 units are sent to control.

- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{i=1}^{d} \lambda_i = d = \operatorname{tr}(\Sigma)$
- We simulate n = 500 observations where 250 units are sent to treatment and 250 units are sent to control.
- Potential outcomes $Y_i(0)$ and $Y_i(1)$ are defined as

$$Y_i(0) = \mathcal{N}(Z\beta_Z, 1)$$

$$Y_i(1) = Y_i(0) + \tau$$

where $\tau = 1$, $\beta_Z \in \mathbb{R}^d$, and Z = XV are the principal components of X.

- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{j=1}^{d} \lambda_j = d = \operatorname{tr}(\Sigma)$
- We simulate n = 500 observations where 250 units are sent to treatment and 250 units are sent to control.
- Potential outcomes $Y_i(0)$ and $Y_i(1)$ are defined as

$$Y_i(0) = \mathcal{N}(Z\beta_Z, 1)$$

$$Y_i(1) = Y_i(0) + \tau$$

where $\tau = 1$, $\beta_Z \in \mathbb{R}^d$, and Z = XV are the principal components of X.

• In this presentation we consider two choices of β_Z :

- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{i=1}^{d} \lambda_i = d = \operatorname{tr}(\Sigma)$
- We simulate n = 500 observations where 250 units are sent to treatment and 250 units are sent to control.
- Potential outcomes $Y_i(0)$ and $Y_i(1)$ are defined as

$$Y_i(0) = \mathcal{N}(Z\beta_Z, 1)$$

$$Y_i(1) = Y_i(0) + \tau$$

where $\tau = 1$, $\beta_Z \in \mathbb{R}^d$, and Z = XV are the principal components of X.

- In this presentation we consider two choices of β_Z :
 - β_Z such that Mahalanobis is optimal $(A = \Sigma^{-1})$

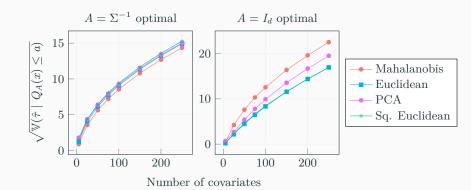
- Because the optimal choice of A depends on β_Z as well as the distribution of eigenvalues, we simulate $\lambda_1, \ldots, \lambda_d \sim d \cdot \text{Dirichlet}(\gamma)$.
- That way $\sum_{i=1}^{d} \lambda_i = d = \operatorname{tr}(\Sigma)$
- We simulate n = 500 observations where 250 units are sent to treatment and 250 units are sent to control.
- Potential outcomes $Y_i(0)$ and $Y_i(1)$ are defined as

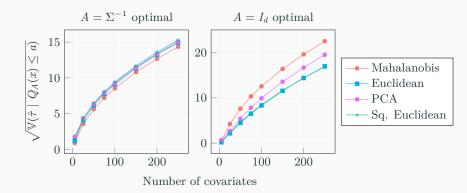
$$Y_i(0) = \mathcal{N}(Z\beta_Z, 1)$$

$$Y_i(1) = Y_i(0) + \tau$$

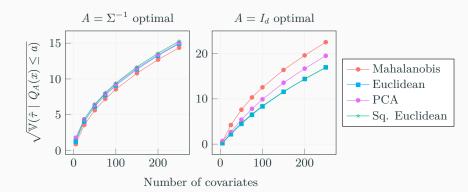
where $\tau = 1$, $\beta_Z \in \mathbb{R}^d$, and Z = XV are the principal components of X.

- In this presentation we consider two choices of β_Z :
 - β_Z such that Mahalanobis is optimal $(A = \Sigma^{-1})$
 - β_Z such that Euclidean is optimal $(A = I_d)$





 \bullet In these settings, Euclidean is either the best or second best choice of A



- \bullet In these settings, Euclidean is either the best or second best choice of A
- We tend to find asymmetry: Mahalanobis is either the best in settings favorable to it, or it is the worst.

 $\bullet~$ We derived general results for rerandomization using any quadratic form.

- We derived general results for rerandomization using any quadratic form.
- This allows researchers to quickly and easily derive new rerandomization schemes not previously considered.

- We derived general results for rerandomization using any quadratic form.
- This allows researchers to quickly and easily derive new rerandomization schemes not previously considered.
- We find that $A = \Sigma^{-1}$ and $A = I_d$ optimize different measurements of covariate balance.

- We derived general results for rerandomization using any quadratic form.
- This allows researchers to quickly and easily derive new rerandomization schemes not previously considered.
- We find that $A = \Sigma^{-1}$ and $A = I_d$ optimize different measurements of covariate balance.
- However, $A = I_d$ is never too far away from the optimal choice, and tends to be more precise on average.

- We derived general results for rerandomization using any quadratic form.
- This allows researchers to quickly and easily derive new rerandomization schemes not previously considered.
- We find that $A = \Sigma^{-1}$ and $A = I_d$ optimize different measurements of covariate balance.
- However, $A = I_d$ is never too far away from the optimal choice, and tends to be more precise on average.
- In future work, we plan to extend this framework to sequential experiments as well as multi-valued treatments.

Thank you!

• Questions?

References

Branson, Zach, Tirthankar Dasgupta, and Donald B Rubin (2016). "Improving covariate balance in 2 K factorial designs via rerandomization with an application to a New York City Department of Education High School Study". In.

Branson, Zach, Xinran Li, and Peng Ding (May 2023). "Power and sample size calculations for rerandomization". In: *Biometrika* 111.1,

pp. 355-363. ISSN: 1464-3510. eprint: https://academic.oup.com/biomet/articlepdf/111/1/355/56665596/asad027.pdf.

Branson, Zach and Stephane Shao (2021). "Ridge rerandomization: An experimental design strategy in the presence of covariate collinearity". In: Journal of Statistical Planning and Inference 211, pp. 287–314.

References ii

- Lachin, John M (1988). "Statistical properties of randomization in clinical trials". In: Controlled clinical trials 9.4, pp. 289–311.
- Li, Xinran, Peng Ding, and Donald B Rubin (2018). "Asymptotic theory of rerandomization in treatment–control experiments". In: Proceedings of the National Academy of Sciences 115.37, pp. 9157–9162.
- i (2020). "Rerandomization in 2^K factorial experiments". In.
 - Lin, Yunzhi, Ming Zhu, and Zheng Su (2015). "The pursuit of balance: an overview of covariate-adaptive randomization techniques in clinical trials". In: Contemporary clinical trials 45, pp. 21–25.
- Liu, Zhaoyang et al. (2023). "Bayesian Criterion for Re-randomization". In: arXiv preprint arXiv:2303.07904.
- Lu, Xin et al. (2023). "Design-based theory for cluster rerandomization". In: *Biometrika* 110.2, pp. 467–483.
- Morgan, Kari Lock and Donald B Rubin (2012). "Rerandomization to improve covariate balance in experiments". In.

References iii

- Senn, SJ (1989). "Covariate imbalance and random allocation in clinical trials". In: Statistics in medicine 8.4, pp. 467–475.
- Zhang, Hengtao, Guosheng Yin, and Donald B Rubin (2023). "PCA rerandomization". In: Canadian Journal of Statistics.
- Zhou, Quan et al. (2018). "Sequential rerandomization". In: *Biometrika* 105.3, pp. 745–752.