Using Knockoffs to Find Important Variables with Statistical Guarantees

Lucas Janson

Harvard University Department of Statistics



Models, Inference, and Algorithms @ Broad, December 12, 2018

Collaborators: Emmanuel Candès (Stanford), Yingying Fan, Jinchi Lv (USC)

Problem Statement

- Y an outcome of interest (AKA response or dependent variable),
- X_1, \ldots, X_p a set of p potential explanatory variables (AKA covariates, features, or independent variables),

How can we select important explanatory variables with few mistakes?

- Y an outcome of interest (AKA response or dependent variable),
- X_1, \ldots, X_p a set of p potential explanatory variables (AKA covariates, features, or independent variables),

How can we select important explanatory variables with few mistakes?

Applications to:

• Biology/genomics/health care

- Y an outcome of interest (AKA response or dependent variable),
- X_1, \ldots, X_p a set of p potential explanatory variables (AKA covariates, features, or independent variables),

How can we select important explanatory variables with few mistakes?

Applications to:

- Biology/genomics/health care
- Economics/political science

- Y an outcome of interest (AKA response or dependent variable),
- X_1, \ldots, X_p a set of p potential explanatory variables (AKA covariates, features, or independent variables),

How can we select important explanatory variables with few mistakes?

Applications to:

- Biology/genomics/health care
- Economics/political science
- Industry/technology

What is an important variable?

What is an important variable?

We consider X_j to be unimportant if the conditional distribution of Y given X_1, \ldots, X_p does not depend on X_j . Formally, X_j is unimportant if it is conditionally independent of Y given X_{-j} :

$$Y \perp\!\!\!\perp X_j \mid X_{‐j}$$

What is an important variable?

We consider X_j to be unimportant if the conditional distribution of Y given X_1, \ldots, X_p does not depend on X_j . Formally, X_j is unimportant if it is conditionally independent of Y given X_{-j} :

$$Y \perp \!\!\!\perp X_j \mid X_{\text{-}j}$$

Markov Blanket of Y: smallest set S such that $Y \perp X_{-S} \mid X_S$

What is an important variable?

We consider X_j to be unimportant if the conditional distribution of Y given X_1, \ldots, X_p does not depend on X_j . Formally, X_j is unimportant if it is conditionally independent of Y given X_{-j} :

$$Y \perp\!\!\!\perp X_j \mid X_{‐j}$$

Markov Blanket of Y: smallest set S such that $Y \perp X_{-S} \mid X_S$

For GLMs with no stochastically redundant covariates, equivalent to $\{j : \beta_j = 0\}$

What is an important variable?

We consider X_j to be unimportant if the conditional distribution of Y given X_1, \ldots, X_p does not depend on X_j . Formally, X_j is unimportant if it is conditionally independent of Y given X_{-j} :

$$Y \perp\!\!\!\perp X_j \mid X_{‐j}$$

Markov Blanket of Y: smallest set S such that $Y \perp X_{-S} \mid X_S$

For GLMs with no stochastically redundant covariates, equivalent to $\{j : \beta_j = 0\}$

To make sure we do not make too many mistakes, we seek to select a set \hat{S} to control the **false discovery rate (FDR)**:

$$\mathsf{FDR} = \mathbb{E}\left[\frac{\#\{j \text{ in } \hat{S} : X_j \text{ unimportant}\}}{\#\{j \text{ in } \hat{S}\}}\right] \le q \text{ (e.g., 10\%)}$$

"Here is a set of variables \hat{S} , 90% of which I expect to be important"

Insufficient info to select either variable confidently (needed for FDR control)

Insufficient info to select either variable confidently (needed for FDR control)

Single-variable resolution impossible: wrong question

• Group variables with their highly-correlated neighbors: $\bigcup_{k=1}^{m} G_k = \{1, \dots, p\}$

Insufficient info to select either variable confidently (needed for FDR control) Single-variable resolution impossible: **wrong question**

- Group variables with their highly-correlated neighbors: $\bigcup_{k=1}^{m} G_k = \{1, \dots, p\}$
- Redefine null hypothesis on per-group basis: group G_k is unimportant if

$$Y \perp \!\!\!\perp X_{G_k} \mid X_{\text{-}G_k}$$

Insufficient info to select either variable confidently (needed for FDR control) Single-variable resolution impossible: **wrong question**

- Group variables with their highly-correlated neighbors: $\bigcup_{k=1}^{m} G_k = \{1, \dots, p\}$
- Redefine null hypothesis on per-group basis: group G_k is unimportant if

$$Y \perp \!\!\!\perp X_{G_k} \mid X_{\text{-}G_k}$$

• Redefine FDR: for selected set of groups \hat{S}_{G} ,

$$\mathsf{FDR}_G = \mathbb{E}\left[\frac{\#\{k \text{ in } \hat{S}_G : G_k \text{ contains no important variables}\}}{\#\{j \text{ in } \hat{S}_G\}}\right] \le q \quad (\text{e.g., } 10\%)$$

Insufficient info to select either variable confidently (needed for FDR control) Single-variable resolution impossible: **wrong question**

- Group variables with their highly-correlated neighbors: $\bigcup_{k=1}^{m} G_k = \{1, \dots, p\}$
- Redefine null hypothesis on per-group basis: group G_k is unimportant if

$$Y \perp \!\!\!\perp X_{G_k} \mid X_{\text{-}G_k}$$

• Redefine FDR: for selected set of groups \hat{S}_{G} ,

$$\mathsf{FDR}_G = \mathbb{E}\left[\frac{\#\{k \text{ in } \hat{S}_G : G_k \text{ contains no important variables}\}}{\#\{j \text{ in } \hat{S}_G\}}\right] \le q \quad (\text{e.g., } 10\%)$$

Everything in this talk works for this setting! (Dai and Barber, 2016)

Model-X knockoffs uses knowledge of X's distribution to solve the controlled variable selection problem with

- Any model for Y and X_1, \ldots, X_p
- Any dimension (including p > n)
- Finite-sample control (non-asymptotic) of FDR
- Practical performance on real problems

Model-X knockoffs uses knowledge of X's distribution to solve the controlled variable selection problem with

- Any model for Y and X_1, \ldots, X_p
- Any dimension (including p > n)
- Finite-sample control (non-asymptotic) of FDR
- Practical performance on real problems

Analysis of the genetic basis of Crohn's Disease (WTCCC, 2007)

- $\approx 5,000$ subjects ($\approx 40\%$ with Crohn's Disease)
- $\approx 375,000$ single nucleotide polymorphisms (SNPs) for each subject

Model-X knockoffs uses knowledge of X's distribution to solve the controlled variable selection problem with

- Any model for Y and X_1, \ldots, X_p
- Any dimension (including p > n)
- Finite-sample control (non-asymptotic) of FDR
- Practical performance on real problems

Analysis of the genetic basis of Crohn's Disease (WTCCC, 2007)

- $\approx 5,000$ subjects ($\approx 40\%$ with Crohn's Disease)
- $\approx 375,000$ single nucleotide polymorphisms (SNPs) for each subject

Original analysis of the data made **9** discoveries by running marginal tests and selecting p-values to target a FDR of 10%

Model-X knockoffs uses knowledge of X's distribution to solve the controlled variable selection problem with

- Any model for Y and X_1, \ldots, X_p
- Any dimension (including p > n)
- Finite-sample control (non-asymptotic) of FDR
- Practical performance on real problems

Analysis of the genetic basis of Crohn's Disease (WTCCC, 2007)

- $\approx 5,000$ subjects ($\approx 40\%$ with Crohn's Disease)
- $\approx 375,000$ single nucleotide polymorphisms (SNPs) for each subject

Original analysis of the data made **9** discoveries by running marginal tests and selecting p-values to target a FDR of 10%

Knockoffs used the same FDR of 10% and made 18 discoveries, with many of the new discoveries confirmed by a larger meta-analysis

Existing Methods for Controlled Variable Selection

- Marginal p-values
 - Excellent exploratory tool
 - Answer wrong question $Y \perp X_j$ instead of $Y \perp X_j \mid X_{-j}$
 - Can lose power, interpretation, and FDR control when X_j are correlated

Existing Methods for Controlled Variable Selection

- Marginal p-values
 - Excellent exploratory tool
 - Answer wrong question $Y \perp X_j$ instead of $Y \perp X_j \mid X_{-j}$
 - Can lose power, interpretation, and FDR control when X_j are correlated
- Bayesian inference
 - Great way of incorporating prior information
 - Computation constrains to very simple priors which may not match actual prior knowledge
 - Inference (esp. in high dimensions) is sensitive to choice of prior

Existing Methods for Controlled Variable Selection

- Marginal p-values
 - Excellent exploratory tool
 - Answer wrong question $Y \perp X_j$ instead of $Y \perp X_j \mid X_{-j}$
 - Can lose power, interpretation, and FDR control when X_j are correlated
- Bayesian inference
 - Great way of incorporating prior information
 - Computation constrains to very simple priors which may not match actual prior knowledge
 - Inference (esp. in high dimensions) is sensitive to choice of prior
- Machine learning
 - Excellent for prediction
 - Cross-validation comes with no statistical guarantees
 - Statistical analysis exists only for simplest methods (lasso) and makes unrealistic assumptions

Knockoffs

You have:

• n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

Knockoffs allows you to:

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

Knockoffs allows you to:

$$\boldsymbol{y}, \quad \boldsymbol{X}_1, \dots, \boldsymbol{X}_p$$

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

Knockoffs allows you to:

$$egin{array}{ccc} oldsymbol{y}, & oldsymbol{X}_1, \dots, oldsymbol{X}_p & & & \ & & \downarrow & \ & & & Z_1, \dots, Z_p \end{array}$$

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

Knockoffs allows you to:

You have:

- n data samples of Y and X stacked into $oldsymbol{y} \in \mathbb{R}^n$ and $oldsymbol{X} \in \mathbb{R}^{n imes p}$
- Algorithm to compute variable importance measure Z_j of each X_j for Y
 - This need not be based on any statistical model, or have any statistical properties at all
 - For instance, you could fit any machine learning method and use the drop in prediction accuracy when X_j is removed from the data
- Desired FDR level q but no way to use Z_j to control it

Knockoffs allows you to:

$$oldsymbol{y}, oldsymbol{X}_1, \dots, oldsymbol{X}_p$$
 $oldsymbol{\downarrow}$ Variable importances $oldsymbol{Z}_1, \dots, oldsymbol{Z}_p$

$$\stackrel{\mathsf{knockoffs}}{\longrightarrow} \hat{S} \subseteq \{1, \dots, p\} \text{ s.t. } \mathsf{FDR} \leq q$$

(1) Construct knockoffs:

- Artificial versions ("knockoffs") of each variable
- Act as controls for assessing importance of original variables

(1) Construct knockoffs:

- Artificial versions ("knockoffs") of each variable
- Act as controls for assessing importance of original variables

(2) Compute knockoff statistics:

- Compute variable importance measures for all variables and their knockoffs
- For each variable, compute W_j as how much *more* important the original variable is than its knockoff

(1) Construct knockoffs:

- Artificial versions ("knockoffs") of each variable
- Act as controls for assessing importance of original variables

(2) Compute knockoff statistics:

- Compute variable importance measures for all variables and their knockoffs
- For each variable, compute W_j as how much *more* important the original variable is than its knockoff

(3) Find the knockoff threshold:

- Order the variables by decreasing $|W_j|$ and proceed down list
- Select variables with positive W_j until an FDR goes above q

(1) Construct knockoffs:

- Artificial versions ("knockoffs") of each variable
- Act as controls for assessing importance of original variables

(2) Compute knockoff statistics:

- Compute variable importance measures for all variables and their knockoffs
- For each variable, compute W_j as how much *more* important the original variable is than its knockoff

(3) Find the knockoff threshold:

- ullet Order the variables by decreasing $|W_j|$ and proceed down list
- Select variables with positive W_j until an FDR goes above q

Coin-flipping property: The key to knockoffs is that steps (1) and (2) are done specifically to ensure that, conditional on $|W_1|, \ldots, |W_p|$, the signs of the *unimportant/null* W_j are independently ± 1 with probability 1/2

Null distribution of variable importance measure

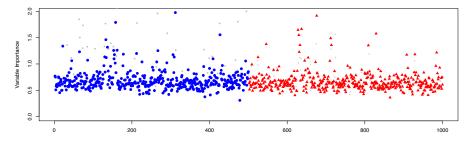


Figure: Variable importances for 500 variables and their knockoffs. Colored points are nulls, grey are non-nulls.

Step (1): Construct Knockoffs

Valid knockoffs are defined by

(1) Swap exchangeability:

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

Valid knockoffs are defined by

(1) Swap exchangeability:

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

(2) Nullity: $\widetilde{X} \perp y \mid X$ (don't look at y when constructing \widetilde{X})

Valid knockoffs are defined by

(1) Swap exchangeability:

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{\widetilde{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{X}_p, \, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{\widetilde{X}}_p \end{bmatrix}$$

(2) Nullity: $\widetilde{X} \perp y \mid X$ (don't look at y when constructing \widetilde{X})

 $\underline{\mathsf{Example}}:\; (X_1,\ldots,X_p) \sim \mathcal{N}(\mathbf{0},\boldsymbol{\Sigma}), \; \mathsf{need}$

$$\operatorname{Cov}(X_1,\ldots,X_p,\widetilde{X}_1,\ldots,\widetilde{X}_p) = \begin{bmatrix} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{bmatrix}$$

Valid knockoffs are defined by

(1) Swap exchangeability:

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

(2) Nullity: $\widetilde{X} \perp y \mid X$ (don't look at y when constructing \widetilde{X})

Example: $(X_1, \ldots, X_p) \sim \mathcal{N}(\mathbf{0}, \Sigma)$, need

$$\operatorname{Cov}(X_1,\ldots,X_p,\widetilde{X}_1,\ldots,\widetilde{X}_p) = \left[\begin{array}{cc} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{array}\right]$$

Semidefinite program construction for s:

$$\begin{array}{ll} \mbox{minimize} & \sum_{j} |\Sigma_{jj} - s_{j}| \\ \mbox{subject to} & s_{j} \geq 0 \\ & \mbox{diag}\{s\} \preceq 2\Sigma, \end{array}$$

ъ **П**

Other Knockoff Constructions

Valid knockoff variables can always be generated:

Algorithm 1 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \widetilde{X}_j \text{ from } \mathcal{L}(X_j | X_{-j}, \widetilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Other Knockoff Constructions

Valid knockoff variables can always be generated:

Algorithm 1 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \widetilde{X}_j \text{ from } \mathcal{L}(X_j | X_{-j}, \widetilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Efficient knockoff constructions for the following X distributions:

- Multivariate Gaussian (Candès et al., 2018)
- Discrete Markov chains (Sesia et al., 2018)
- Hidden Markov models (Sesia et al., 2018)
- Gaussian mixture models (Gimenez et al., 2018)
- General graphical models (ongoing work with Wenshuo and others)

Other Knockoff Constructions

Valid knockoff variables can always be generated:

Algorithm 1 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \widetilde{X}_j \text{ from } \mathcal{L}(X_j | X_{-j}, \widetilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Efficient knockoff constructions for the following X distributions:

- Multivariate Gaussian (Candès et al., 2018)
- Discrete Markov chains (Sesia et al., 2018)
- Hidden Markov models (Sesia et al., 2018)
- Gaussian mixture models (Gimenez et al., 2018)
- General graphical models (ongoing work with Wenshuo and others)

Approximate knockoff constructions (no theoretical guarantees):

- Second-order knockoffs (tend to work well with regression-based statistics) (Candès et al., 2018)
- Deep learning, including GANs (empirically valid in low-dimensions n > p) (Romano et al., 2018; Liu and Zheng, 2018; Anonymous, 2018)

Knockoffs

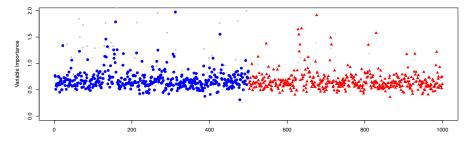


Figure: Variable importances for 500 variables and their knockoffs. Colored points are nulls, grey are non-nulls.

i.i.d. Gaussians

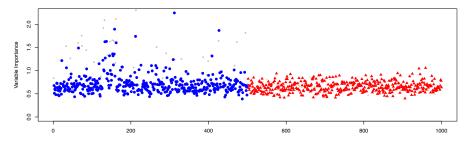


Figure: Variable importances for 500 variables and their knockoffs. Colored points are nulls, grey are non-nulls.

Permutations

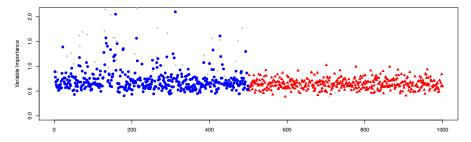
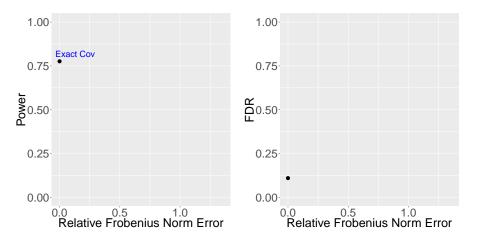
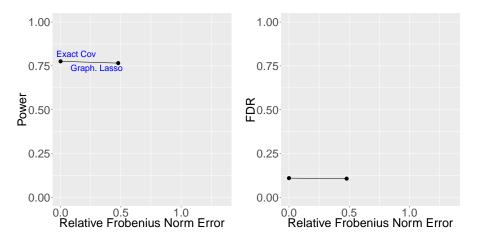
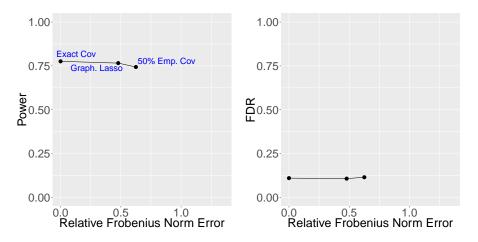
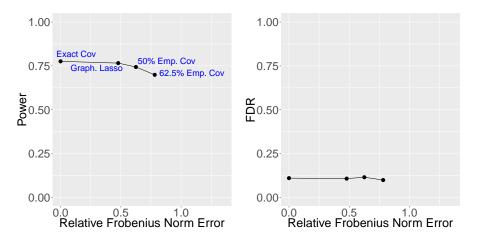


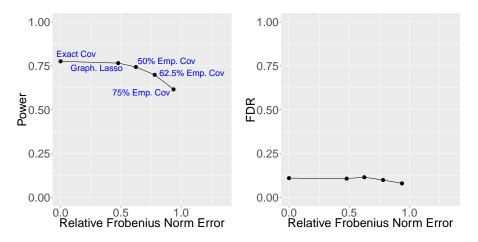
Figure: Variable importances for 500 variables and their knockoffs. Colored points are nulls, grey are non-nulls.

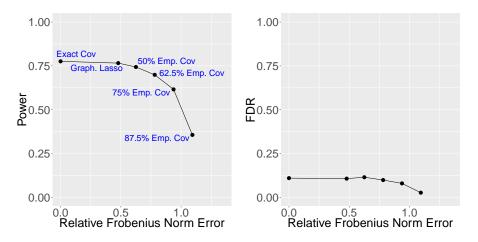


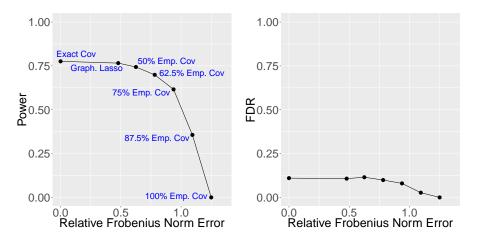












Step (2): Compute Knockoff Statistics

• Variable importance measures for all original and knockoff variables

 $Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$

• Variable importance measures for all original and knockoff variables

$$Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$$

• Antisymmetric function $f_j: \mathbb{R}^2 \to \mathbb{R}$, i.e., $f_j(a,b) = -f_j(b,a)$

• Variable importance measures for all original and knockoff variables

$$Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$$

• Antisymmetric function $f_j: \mathbb{R}^2 \to \mathbb{R}$, i.e., $f_j(a,b) = -f_j(b,a)$ • $W_j = f_j(Z_j, \widetilde{Z}_j)$

• Variable importance measures for all original and knockoff variables

$$Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$$

Antisymmetric function f_j : ℝ² → ℝ, i.e., f_j(a, b) = -f_j(b, a)
W_i = f_i(Z_i, Ž_i)

Example 1:

• Z is magnitude of fitted coefficient β from a lasso regression of ${\pmb y}$ on $[{\pmb X}\widetilde{\pmb X}]$ • $f_j(a,b)=a-b$

• Variable importance measures for all original and knockoff variables

$$Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$$

Antisymmetric function f_j : ℝ² → ℝ, i.e., f_j(a, b) = -f_j(b, a)
W_i = f_i(Z_i, Ž_i)

Example 1:

- Z is magnitude of fitted coefficient β from a lasso regression of y on $[X\widetilde{X}]$ • $f_j(a,b) = a - b$
 - $W_j = |\beta_j| |\tilde{\beta}_j|$ (Lasso Coefficient Difference statistic)

• Variable importance measures for all original and knockoff variables

$$Z_1,\ldots,Z_p,\widetilde{Z}_1,\ldots,\widetilde{Z}_p$$

Antisymmetric function f_j : ℝ² → ℝ, i.e., f_j(a,b) = -f_j(b,a)
W_i = f_i(Z_i, Ž_i)

Example 1:

• Z is magnitude of fitted coefficient β from a lasso regression of y on $[X\widetilde{X}]$ • $f_j(a,b) = a - b$

 $W_j = |\beta_j| - |\tilde{\beta}_j|$ (Lasso Coefficient Difference statistic)

Example 2:

- Fit machine learning method (e.g., deep learning) to y with features [XX]
- Z is increase in cross-validation error when variable is dropped; same f_j

$$(\mathsf{CV}_{-j} - \mathsf{CV}) - (\widetilde{\mathsf{CV}}_{-j} - \mathsf{CV}) = \mathsf{CV}_{-j} - \widetilde{\mathsf{CV}}_{-j}$$

Adaptivity and Prior Information in \boldsymbol{Z}

Adaptivity

• Z can be **any** variable importance measure

- Z can be **any** variable importance measure
- Higher-level adaptivity: CV to choose best-fitting model for inference

- $\bullet~Z$ can be any variable importance measure
- Higher-level adaptivity: CV to choose best-fitting model for inference
 - E.g., fit random forest and ℓ_1 -penalized regression; derive feature importance from whichever has lower CV error—still strict FDR control

- $\bullet~Z$ can be any variable importance measure
- Higher-level adaptivity: CV to choose best-fitting model for inference
 - E.g., fit random forest and ℓ_1 -penalized regression; derive feature importance from whichever has lower CV error—still strict FDR control
- Can even let analyst look at (masked version of) data to choose Z function

- Z can be **any** variable importance measure
- Higher-level adaptivity: CV to choose best-fitting model for inference
 - E.g., fit random forest and ℓ_1 -penalized regression; derive feature importance from whichever has lower CV error—still strict FDR control
- Can even let analyst look at (masked version of) data to choose Z function

Prior information

• **Bayesian approach**: choose prior and model, and Z_j could be the posterior probability that X_j contributes to the model

- $\bullet~Z$ can be any variable importance measure
- Higher-level adaptivity: CV to choose best-fitting model for inference
 - E.g., fit random forest and ℓ_1 -penalized regression; derive feature importance from whichever has lower CV error—still strict FDR control
- Can even let analyst look at (masked version of) data to choose Z function

Prior information

- **Bayesian approach**: choose prior and model, and Z_j could be the posterior probability that X_j contributes to the model
- Still strict FDR control, even if wrong prior or MCMC has not converged

Why Does it Work?

Recall swap exchangeability property: for any j,

$$\begin{split} & [\boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{\widetilde{X}}_p] \\ \stackrel{\mathcal{D}}{=} [\boldsymbol{X}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{X}_p, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{\widetilde{X}}_p] \end{split}$$

Why Does it Work?

Recall swap exchangeability property: for any j,

$$\begin{split} & [\boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{\widetilde{X}}_p] \\ \stackrel{\mathcal{D}}{=} [\boldsymbol{X}_1, \cdots, \boldsymbol{\widetilde{X}}_j, \cdots, \boldsymbol{X}_p, \boldsymbol{\widetilde{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{\widetilde{X}}_p] \end{split}$$

Coin-flipping property for W_j :

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\left(Z_j, \widetilde{Z}_j\right) := \left(Z_j\left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right]\right), \quad \widetilde{Z}_j\left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right]\right)\right)$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix}$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \end{pmatrix}$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j, Z_j \end{pmatrix}$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j, Z_j \end{pmatrix}$$

$$W_j = f_j(Z_j, \widetilde{Z}_j) \stackrel{\mathcal{D}}{=} f_j(\widetilde{Z}_j, Z_j)$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j, Z_j \end{pmatrix}$$

$$W_j = f_j(Z_j, \widetilde{Z}_j) \stackrel{\mathcal{D}}{=} f_j(\widetilde{Z}_j, Z_j) = -f_j(Z_j, \widetilde{Z}_j) = -W_j$$

Recall swap exchangeability property: for any j,

$$\begin{bmatrix} \boldsymbol{X}_1, \cdots, \boldsymbol{X}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$
$$\stackrel{\mathcal{D}}{=} \begin{bmatrix} \boldsymbol{X}_1, \cdots, \widetilde{\boldsymbol{X}}_j, \cdots, \boldsymbol{X}_p, \, \widetilde{\boldsymbol{X}}_1, \cdots, \boldsymbol{X}_j, \cdots, \widetilde{\boldsymbol{X}}_p \end{bmatrix}$$

$$\begin{pmatrix} Z_j, \widetilde{Z}_j \end{pmatrix} := \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ \stackrel{\mathcal{D}}{=} \begin{pmatrix} Z_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right), & \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \widetilde{\boldsymbol{X}}_j \cdots \boldsymbol{X}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right), & Z_j \left(\boldsymbol{y}, \left[\cdots \boldsymbol{X}_j \cdots \widetilde{\boldsymbol{X}}_j \cdots \right] \right) \end{pmatrix} \\ = \begin{pmatrix} \widetilde{Z}_j, \left[\widetilde{Z}_j, \widetilde{Z}_j \right] \end{pmatrix}$$

$$W_j \stackrel{\mathcal{D}}{=} -W_j$$

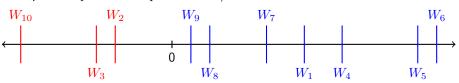
Step (3): Find the Knockoff Threshold

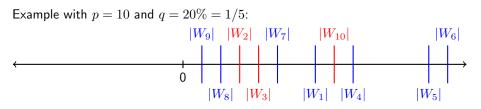
Example with p = 10 and q = 20% = 1/5:

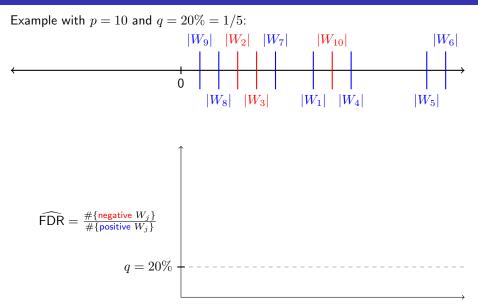
Example with p = 10 and q = 20% = 1/5:

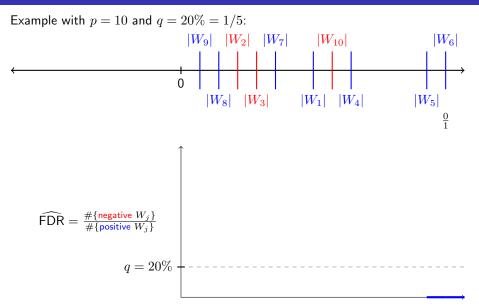


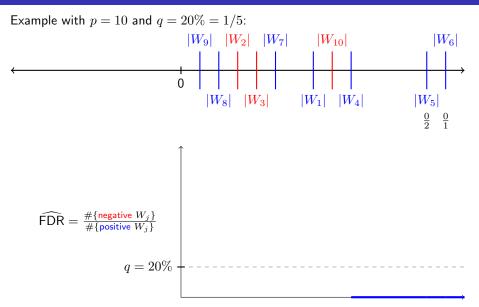
Example with p = 10 and q = 20% = 1/5:

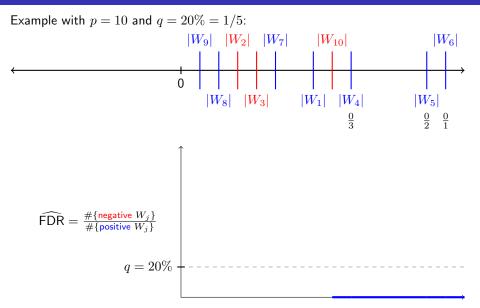


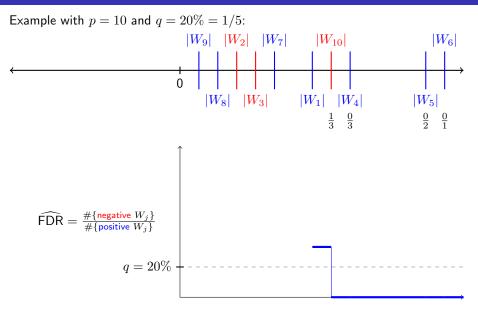


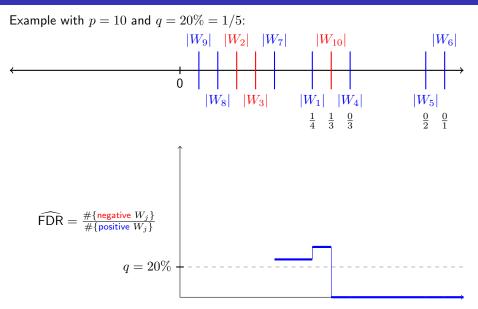


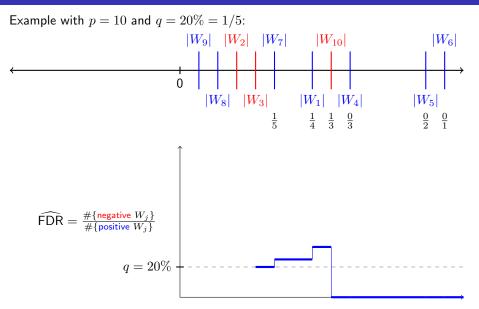


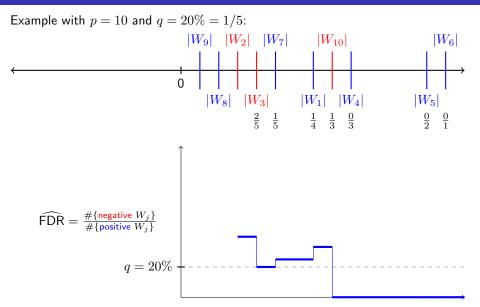


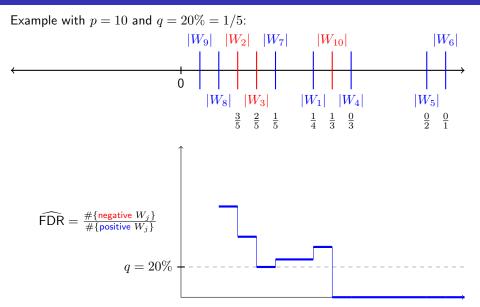


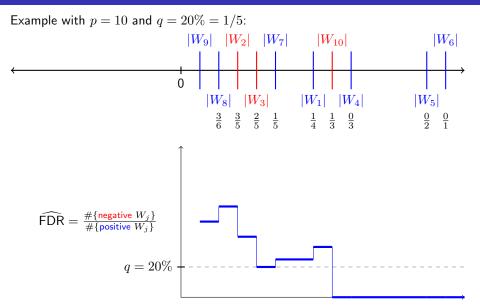


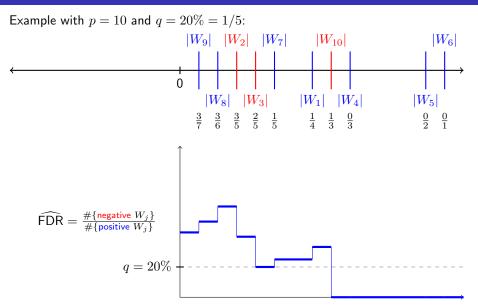


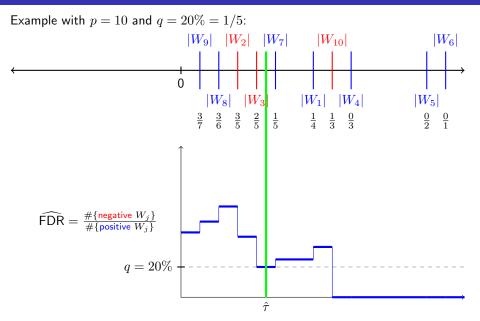


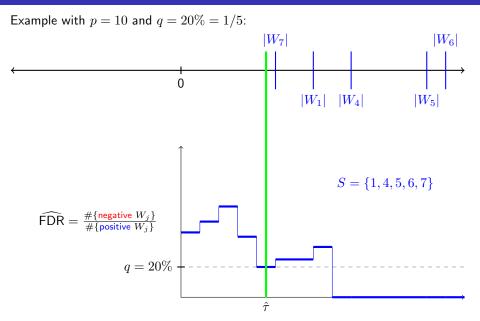












$$\mathsf{FDR} = \mathbb{E}\left[\frac{\#\{\mathsf{null} \ \boldsymbol{X}_j \ \mathsf{selected}\}}{\#\{\mathsf{total} \ \boldsymbol{X}_j \ \mathsf{selected}\}}\right]$$

$$\begin{aligned} \mathsf{FDR} \ &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \boldsymbol{X}_j\ \mathsf{selected}\}}{\#\{\mathsf{total}\ \boldsymbol{X}_j\ \mathsf{selected}\}}\right] \\ &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \mathsf{positive}\ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive}\ |W_j| > \hat{\tau}\}}\right] \end{aligned}$$

$$\begin{aligned} \mathsf{FDR} \ &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \boldsymbol{X}_j\ \mathsf{selected}\}}{\#\{\mathsf{total}\ \boldsymbol{X}_j\ \mathsf{selected}\}}\right] \\ &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \mathsf{positive}\ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive}\ |W_j| > \hat{\tau}\}}\right] \\ &\approx \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \mathsf{negative}\ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive}\ |W_j| > \hat{\tau}\}}\right] \end{aligned}$$

$$\begin{aligned} \mathsf{FDR} \ &= \mathbb{E} \left[\frac{\#\{\mathsf{null} \ X_j \ \mathsf{selected}\}}{\#\{\mathsf{total} \ X_j \ \mathsf{selected}\}} \right] \\ &= \mathbb{E} \left[\frac{\#\{\mathsf{null} \ \mathsf{positive} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \\ &\approx \mathbb{E} \left[\frac{\#\{\mathsf{null} \ \mathsf{negative} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \\ &\leq \mathbb{E} \left[\frac{\#\{\mathsf{negative} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \end{aligned}$$

$$\begin{aligned} \mathsf{FDR} &= \mathbb{E} \left[\frac{\#\{\mathsf{null} \ X_j \ \mathsf{selected}\}}{\#\{\mathsf{total} \ X_j \ \mathsf{selected}\}} \right] \\ &= \mathbb{E} \left[\frac{\#\{\mathsf{null} \ \mathsf{positive} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \\ &\approx \mathbb{E} \left[\frac{\#\{\mathsf{null} \ \mathsf{negative} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \\ &\leq \mathbb{E} \left[\frac{\#\{\mathsf{negative} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \\ &= \mathbb{E} \left[\frac{\#\{\mathsf{negative} \ |W_j| > \hat{\tau}\}}{\#\{\mathsf{positive} \ |W_j| > \hat{\tau}\}} \right] \end{aligned}$$

Numerical Results

Simulations in Low-Dimensional Linear Model

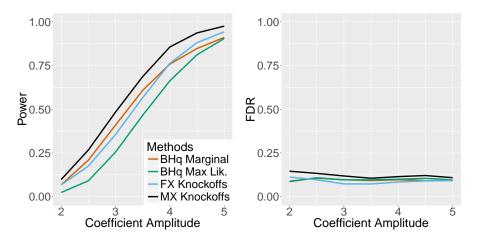


Figure: Power and FDR (target is 10%) for knockoffs and alternative procedures. The design matrix is i.i.d. $\mathcal{N}(0, 1/n)$, n = 3000, p = 1000, and y comes from a Gaussian linear model with 60 nonzero regression coefficients having equal magnitudes and random signs. The noise variance is 1.

Simulations in Low-Dimensional Nonlinear Model

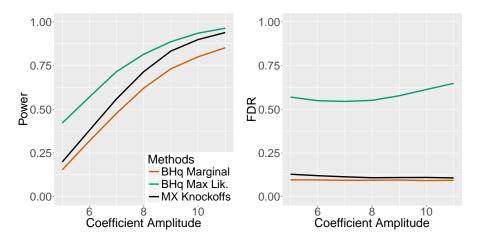


Figure: Power and FDR (target is 10%) for knockoffs and alternative procedures. The design matrix is i.i.d. $\mathcal{N}(0, 1/n)$, n = 3000, p = 1000, and y comes from a binomial linear model with logit link function, and 60 nonzero regression coefficients having equal magnitudes and random signs.

Simulations in High Dimensions

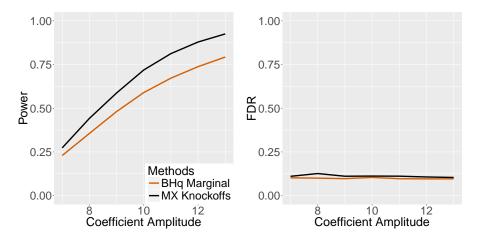


Figure: Power and FDR (target is 10%) for knockoffs and alternative procedures. The design matrix is i.i.d. $\mathcal{N}(0, 1/n)$, n = 3000, p = 6000, and y comes from a binomial linear model with logit link function, and 60 nonzero regression coefficients having equal magnitudes and random signs.

Simulations in High Dimensions with Dependence

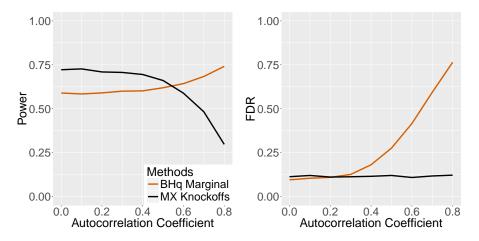


Figure: Power and FDR (target is 10%) for knockoffs and alternative procedures. The design matrix has AR(1) columns, and marginally each $X_j \sim \mathcal{N}(0, 1/n)$. n = 3000, p = 6000, and y follows a binomial linear model with logit link function, and 60 nonzero coefficients with random signs and randomly selected locations.

Lucas Janson (Harvard Statistics)

Knockoffs for Controlled Variable Selection

Genetic Analysis of Crohn's Disease

2007 case-control study by WTCCC

• $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)
- Entire analysis took 6 hours of serial computation time; 1 hour in parallel

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)
- Entire analysis took 6 hours of serial computation time; 1 hour in parallel
- Knockoffs made twice as many discoveries as original analysis

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)
- Entire analysis took 6 hours of serial computation time; 1 hour in parallel
- Knockoffs made twice as many discoveries as original analysis
 - Some new discoveries confirmed in larger study

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)
- Entire analysis took 6 hours of serial computation time; 1 hour in parallel
- Knockoffs made twice as many discoveries as original analysis
 - Some new discoveries confirmed in larger study
 - Some corroborated by work on nearby genes: promising candidates

- $n \approx 5,000$, $p \approx 375,000$; preprocessing mirrored original analysis
- **Strong spatial structure**: second-order knockoffs generated on genetic covariance estimate (Wen and Stephens, 2010)
- Entire analysis took 6 hours of serial computation time; 1 hour in parallel
- Knockoffs made twice as many discoveries as original analysis
 - Some new discoveries confirmed in larger study
 - Some corroborated by work on nearby genes: promising candidates
- Similar result obtained with X model taken from existing genomic imputation software (Sesia et al., 2018)

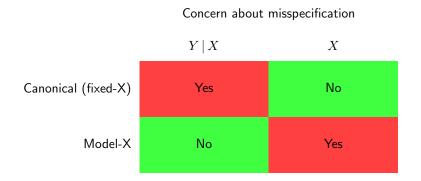
• R, Python, and Matlab packages available depending on knockoff construction; link on my website

- R, Python, and Matlab packages available depending on knockoff construction; link on my website
- $\bullet\,$ Knockoff construction algorithms generally scale linearly in p and n

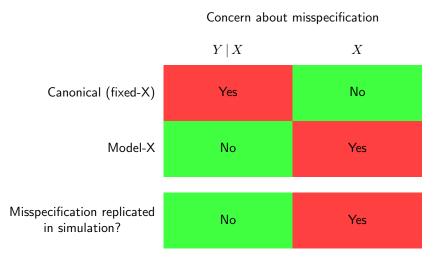
- R, Python, and Matlab packages available depending on knockoff construction; link on my website
- $\bullet\,$ Knockoff construction algorithms generally scale linearly in p and n
- Given variable importances $Z_1, \ldots, Z_p, \widetilde{Z}_1, \ldots, \widetilde{Z}_p$, computation trivial

- R, Python, and Matlab packages available depending on knockoff construction; link on my website
- $\bullet\,$ Knockoff construction algorithms generally scale linearly in p and n
- Given variable importances $Z_1, \ldots, Z_p, \widetilde{Z}_1, \ldots, \widetilde{Z}_p$, computation trivial
- Need to compute $Z_1, \ldots, Z_p, \widetilde{Z}_1, \ldots, \widetilde{Z}_p$
 - Just compute variable importances for twice as many variables
 - Generally only constant slower than computing variable importances without knockoffs

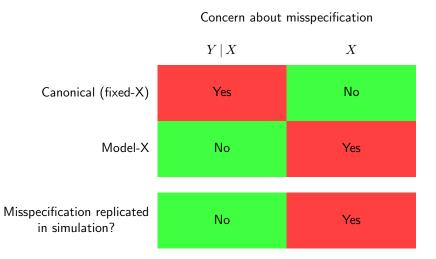
Checking Sensitivity to Misspecification Error



Checking Sensitivity to Misspecification Error



Checking Sensitivity to Misspecification Error



Model-X: can actually check sensitivity to misspecification error!

Lucas Janson (Harvard Statistics)

Robustness on Real Data

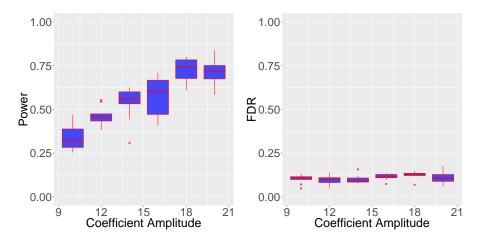


Figure: Power and FDR (target is 10%) for knockoffs applied to subsamples of a chromosome 1 of real genetic design matrix; $n \approx 1,400$.

The controlled variable selection problem is ubiquitous in modern science, and knockoffs is a powerful, flexible, and robust solution

The controlled variable selection problem is ubiquitous in modern science, and knockoffs is a powerful, flexible, and robust solution

My group is extremely interested in **making knockoffs work for you**. Please reach out if you think it could help with your research.

The controlled variable selection problem is ubiquitous in modern science, and knockoffs is a powerful, flexible, and robust solution

My group is extremely interested in **making knockoffs work for you**. Please reach out if you think it could help with your research.

Thank you!

http://lucasjanson.fas.harvard.edu ljanson@fas.harvard.edu

Appendix

References

- Anonymous (2018). KnockoffGAN: generating knockoffs for feature selection using generative adversarial networks. *Submitted to International Conference on Learning Representations*.
- Barber, R. F. and Candès, E. J. (2015). Controlling the false discovery rate via knockoffs. *Ann. Statist.*, 43(5):2055–2085.
- Candès, E., Fan, Y., Janson, L., and Lv, J. (2018). Panning for gold: 'model-X' knockoffs for high dimensional controlled variable selection. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 80(3):551–577.
- Dai, R. and Barber, R. F. (2016). The knockoff filter for FDR control in group-sparse and multitask regression. In *Proceedings of the 33nd International Conference on Machine Learning (ICML 2016).*
- Gimenez, J. R., Ghorbani, A., and Zou, J. (2018). Knockoffs for the mass: new feature importance statistics with false discovery guarantees. *arXiv preprint arXiv:1807.06214*.
- Liu, Y. and Zheng, C. (2018). Auto-encoding knockoff generator for fdr controlled variable selection. *arXiv preprint arXiv:1809.10765*.
- Romano, Y., Sesia, M., and Candès, E. J. (2018). Deep knockoffs. arXiv preprint arXiv:1811.06687.

Sesia, M., Sabatti, C., and Candès, E. J. (2018). Gene hunting with hidden

Existing Methods: Low-Dimensional Linear Model

Suppose we assume that our data:

• follows a linear model:

$$Y = X_1\beta_1 + \dots + X_p\beta_p + \varepsilon, \qquad \varepsilon \sim \mathcal{N}(0, \sigma^2),$$

• has more observations that variables: $n \ge p$ (low-dimensional).

Existing Methods: Low-Dimensional Linear Model

Suppose we assume that our data:

• follows a linear model:

$$Y = X_1\beta_1 + \dots + X_p\beta_p + \varepsilon, \qquad \varepsilon \sim \mathcal{N}(0, \sigma^2),$$

• has more observations that variables: $n \ge p$ (low-dimensional).

Classical problem:

- Ordinary least squares (OLS) theory gives exact p-values for testing whether each β_j = 0 or not (under very mild assumptions, β_j = 0 ⇔ Y ⊥ X_j | X_{-j})
- The Benjamini-Hochberg procedure (BHq) applied to the p-values will essentially control the FDR

Existing Methods: Low-Dimensional Linear Model

Suppose we assume that our data:

• follows a linear model:

$$Y = X_1\beta_1 + \dots + X_p\beta_p + \varepsilon, \qquad \varepsilon \sim \mathcal{N}(0, \sigma^2),$$

• has more observations that variables: $n \ge p$ (low-dimensional).

Classical problem:

- Ordinary least squares (OLS) theory gives exact p-values for testing whether each β_j = 0 or not (under very mild assumptions, β_j = 0 ⇔ Y ⊥ X_j | X_{-j})
- The Benjamini-Hochberg procedure (BHq) applied to the p-values will essentially control the FDR

Minor caveats:

- FDR control not exact (but good enough in practice)
- Sparsity not used (reduces power to find important variables)

Nonlinearity and High Dimensions

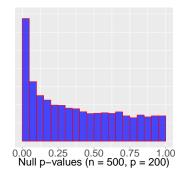
Low-dimensional $(n \ge p)$ generalized linear model

• Apply BHq to asymptotic p-values

Nonlinearity and High Dimensions

Low-dimensional $(n \ge p)$ generalized linear model

- Apply BHq to asymptotic p-values
- Can be far from valid in practice

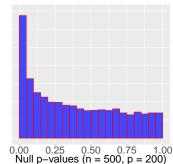


Nonlinearity and High Dimensions

Low-dimensional $(n \ge p)$ generalized linear model

- Apply BHq to asymptotic p-values
- Can be far from valid in practice

High-dimensional (n < p) generalized linear models



- Apply BHq to p-values from
 - Debiased lasso, e.g., Zhang and Zhang (2014), Javanmard and Montanari (2014), van de Geer et al. (2014), Cai and Guo (2015)
 - Causal inference, e.g., Belloni et al. (2014), Athey et al. (2016), Farrell (2015)
 - Inference after selection, e.g., Berk et al. (2013), Lee et al. (2016), Fithian et al. (2014)
- Asymptotic, require sparsity and random design assumptions

Algorithm 2 Sequential Conditional Independent Pairs

 $\begin{array}{l} \text{for } j = \{1, \ldots, p\} \text{ do} \\ \mid & \text{Sample } \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{\text{-}j}, \, \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \\ \text{end} \end{array}$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, \dots, p\}$ do $| Sample \ \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{-j}, \ \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

• Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \tilde{X}_j \text{ from } \mathcal{L}(X_j | X_{-j}, \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

- Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$
- Conditional PMF of $\tilde{X}_j | X_{1:p}, \tilde{X}_{1:j-1}$ is

$$\frac{\mathcal{L}(X_{\cdot j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\cdot j}, u, \tilde{X}_{1:j-1})}.$$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \ \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{-j}, \ \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

- Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$
- Conditional PMF of $\tilde{X}_j | X_{1:p}, \tilde{X}_{1:j-1}$ is

$$\frac{\mathcal{L}(X_{\cdot j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\cdot j}, u, \tilde{X}_{1:j-1})}.$$

$$\frac{\mathcal{L}(X_{\text{-}j}, X_j, \tilde{X}_{1:j-1})\mathcal{L}(X_{\text{-}j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\text{-}j}, u, \tilde{X}_{1:j-1})}$$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \ \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{-j}, \ \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

- Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$
- Conditional PMF of $\tilde{X}_j | X_{1:p}, \tilde{X}_{1:j-1}$ is

$$\frac{\mathcal{L}(X_{\cdot j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\cdot j}, u, \tilde{X}_{1:j-1})}.$$

$$\frac{\mathcal{L}(X_{\text{-}j}, \frac{X_j}{X_j}, \tilde{X}_{1:j-1})\mathcal{L}(X_{\text{-}j}, \frac{\tilde{X}_j}{X_j}, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\text{-}j}, u, \tilde{X}_{1:j-1})}$$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, ..., p\}$ do $| Sample \ \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{-j}, \ \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

- Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$
- Conditional PMF of $\tilde{X}_j | X_{1:p}, \tilde{X}_{1:j-1}$ is

$$\frac{\mathcal{L}(X_{\cdot j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\cdot j}, u, \tilde{X}_{1:j-1})}.$$

$$\frac{\mathcal{L}(X_{\text{-}j}, \tilde{X}_j, \tilde{X}_{1:j-1})\mathcal{L}(X_{\text{-}j}, X_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\text{-}j}, u, \tilde{X}_{1:j-1})}$$

Algorithm 2 Sequential Conditional Independent Pairs

for $j = \{1, \dots, p\}$ do $| Sample \ \tilde{X}_j \text{ from } \mathcal{L}(X_j \,|\, X_{-j}, \ \tilde{X}_{1:j-1}) \text{ conditionally independently of } X_j \text{ end}$

Proof sketch (discrete case):

- Denote PMF of $(X_{1:p}, \tilde{X}_{1:j-1})$ by $\mathcal{L}(X_{\cdot j}, X_j, \tilde{X}_{1:j-1})$
- Conditional PMF of $\tilde{X}_j | X_{1:p}, \tilde{X}_{1:j-1}$ is

$$\frac{\mathcal{L}(X_{\cdot j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\cdot j}, u, \tilde{X}_{1:j-1})}.$$

$$\frac{\mathcal{L}(X_{\text{-}j}, X_j, \tilde{X}_{1:j-1})\mathcal{L}(X_{\text{-}j}, \tilde{X}_j, \tilde{X}_{1:j-1})}{\sum_u \mathcal{L}(X_{\text{-}j}, u, \tilde{X}_{1:j-1})}$$

 $\operatorname{Cov}(X_1,\ldots,X_p) = \Sigma$, need:

$$\operatorname{Cov}(X_1,\ldots,X_p,\tilde{X}_1,\ldots,\tilde{X}_p) = \begin{bmatrix} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{bmatrix}$$

 $\operatorname{Cov}(X_1,\ldots,X_p) = \Sigma$, need:

$$\operatorname{Cov}(X_1,\ldots,X_p,\tilde{X}_1,\ldots,\tilde{X}_p) = \begin{bmatrix} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{bmatrix}$$

• Equicorrelated (EQ) (fast, less powerful): $s_j^{\mathsf{EQ}} = 2\lambda_{\min}(\Sigma) \wedge 1$ for all j

 $\operatorname{Cov}(X_1,\ldots,X_p) = \Sigma$, need:

$$\operatorname{Cov}(X_1,\ldots,X_p,\tilde{X}_1,\ldots,\tilde{X}_p) = \begin{bmatrix} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{bmatrix}$$

- Equicorrelated (EQ) (fast, less powerful): $s_j^{\sf EQ} = 2\lambda_{\sf min}(\Sigma) \wedge 1$ for all j
- Semidefinite program (SDP) (slower, more powerful):

minimize subject to

$$\begin{split} & \sum_{j} |1 - s_{j}^{\mathsf{SDP}}| \\ & s_{j}^{\mathsf{SDP}} \geq 0 \\ & \mathsf{diag}\{s^{\mathsf{SDP}}\} \preceq 2 \mathbf{\Sigma} \end{split}$$

 $\operatorname{Cov}(X_1,\ldots,X_p) = \Sigma$, need:

$$\operatorname{Cov}(X_1,\ldots,X_p,\tilde{X}_1,\ldots,\tilde{X}_p) = \begin{bmatrix} \Sigma & \Sigma - \operatorname{diag}\{s\} \\ \Sigma - \operatorname{diag}\{s\} & \Sigma \end{bmatrix}$$

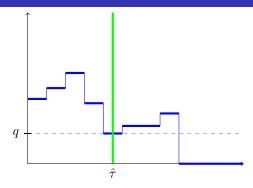
- Equicorrelated (EQ) (fast, less powerful): $s_j^{\mathsf{EQ}} = 2\lambda_{\min}(\mathbf{\Sigma}) \wedge 1$ for all j
- Semidefinite program (SDP) (slower, more powerful):

 $\begin{array}{ll} \mbox{minimize} & \sum_{j} |1 - s_{j}^{\rm SDP}| \\ \mbox{subject to} & s_{j}^{\rm SDP} \geq 0 \\ & \mbox{diag}\{s^{\rm SDP}\} \preceq 2\mathbf{\Sigma}, \end{array}$

• (New) Approximate SDP:

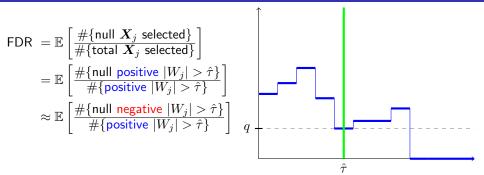
- ullet Approximate Σ as block diagonal so that SDP separates
- Bisection search scalar multiplier of solution to account for approximation
- faster than SDP, more powerful than EQ, and easily parallelizable

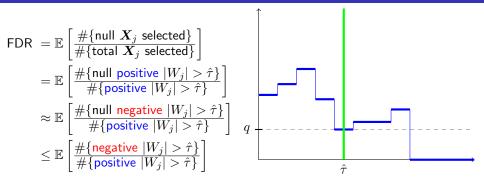
$$\mathsf{FDR} = \mathbb{E}\left[rac{\#\{\mathsf{null} \ oldsymbol{X}_j \ \mathsf{selected}\}}{\#\{\mathsf{total} \ oldsymbol{X}_j \ \mathsf{selected}\}}
ight]$$

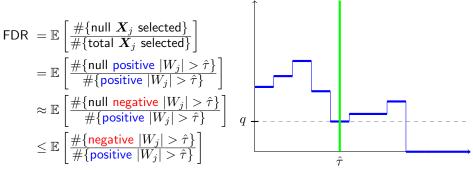


$$FDR = \mathbb{E} \left[\frac{\#\{\text{null } X_j \text{ selected}\}}{\#\{\text{total } X_j \text{ selected}\}} \right]$$
$$= \mathbb{E} \left[\frac{\#\{\text{null positive } |W_j| > \hat{\tau}\}}{\#\{\text{positive } |W_j| > \hat{\tau}\}} \right]$$
$$q$$
$$\hat{\tau}$$

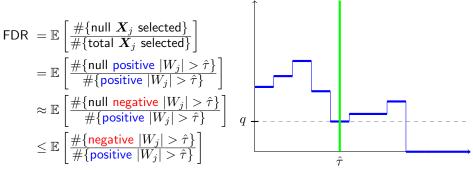
 \mathbf{T}



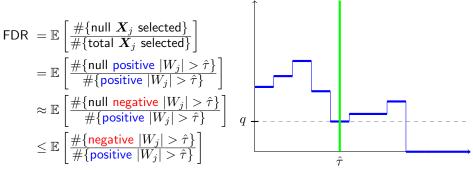




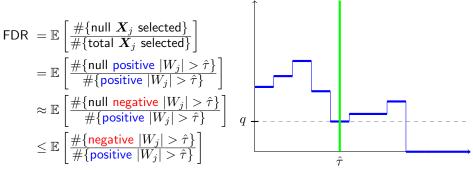
$$\mathsf{mFDR} = \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \boldsymbol{X}_j \; \mathsf{selected}\}}{q^{-1} + \#\{\mathsf{total}\; \boldsymbol{X}_j \; \mathsf{selected}\}}\right] = \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \mathsf{positive}\; |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}\; |W_j| > \hat{\tau}\}}\right]$$



$$\begin{split} \mathsf{mFDR} &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \boldsymbol{X}_j\ \mathsf{selected}\}}{q^{-1} + \#\{\mathsf{total}\ \boldsymbol{X}_j\ \mathsf{selected}\}}\right] = \mathbb{E}\left[\frac{\#\{\mathsf{null}\ \mathsf{positive}\ |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}\ |W_j| > \hat{\tau}\}}\right] \\ &= \mathbb{E}\left(\frac{\#\{\mathsf{null}\ \mathsf{positive}\ |W_j| > \hat{\tau}\}}{1 + \#\{\mathsf{null}\ \mathsf{negative}\ |W_j| > \hat{\tau}\}} \cdot \frac{1 + \#\{\mathsf{null}\ \mathsf{negative}\ |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}\ |W_j| > \hat{\tau}\}}\right) \end{split}$$



$$\begin{split} \mathsf{mFDR} &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \boldsymbol{X}_j \; \mathsf{selected}\}}{q^{-1} + \#\{\mathsf{total}\; \boldsymbol{X}_j \; \mathsf{selected}\}}\right] = \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \mathsf{positive}\; |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}\; |W_j| > \hat{\tau}\}}\right] \\ &= \mathbb{E}\left(\frac{\#\{\mathsf{null}\; \mathsf{positive}\; |W_j| > \hat{\tau}\}}{1 + \#\{\mathsf{null}\; \mathsf{negative}\; |W_j| > \hat{\tau}\}} \cdot \underbrace{\frac{1 + \#\{\mathsf{null}\; \mathsf{negative}\; |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}|W_j| > \hat{\tau}\}}}_{\leq q \; \mathsf{by}\; \mathsf{definition}\; \mathsf{of}\; \hat{\tau}} \end{split}$$



$$\begin{split} \mathsf{mFDR} &= \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \boldsymbol{X}_j \; \mathsf{selected}\}}{q^{-1} + \#\{\mathsf{total}\; \boldsymbol{X}_j \; \mathsf{selected}\}}\right] = \mathbb{E}\left[\frac{\#\{\mathsf{null}\; \mathsf{positive}\; |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}\; |W_j| > \hat{\tau}\}}\right] \\ &= \mathbb{E}\left(\underbrace{\frac{\#\{\mathsf{null}\; \mathsf{positive}\; |W_j| > \hat{\tau}\}}{1 + \#\{\mathsf{null}\; \mathsf{negative}\; |W_j| > \hat{\tau}\}}}_{\mathsf{Supermartingale}\; \leq \; 1} \\ \cdot \underbrace{\frac{1 + \#\{\mathsf{null}\; \mathsf{negative}\; |W_j| > \hat{\tau}\}}{q^{-1} + \#\{\mathsf{positive}|W_j| > \hat{\tau}\}}}_{\leq \; q \; \mathsf{by\; definition\; of}\; \hat{\tau}} \right) \end{split}$$