

Selective peak inference: Unbiased estimation of the effect size at local maxima

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Methods

- \mathcal{V} : set of voxel locations
- Define an **image** to be a map $Z : \mathcal{V} \rightarrow \mathbb{R}$.
- Define a **local maxima** or **peak** of Z to be a voxel $v \in \mathcal{V}$ such that the value that Z takes at that location is larger than the value Z takes at neighbouring voxels

One-Sample Model

Suppose that we have N subjects and for each $n = 1, \dots, N$ a corresponding random image Y_n on \mathcal{V} such that for every voxel $v \in \mathcal{V}$,

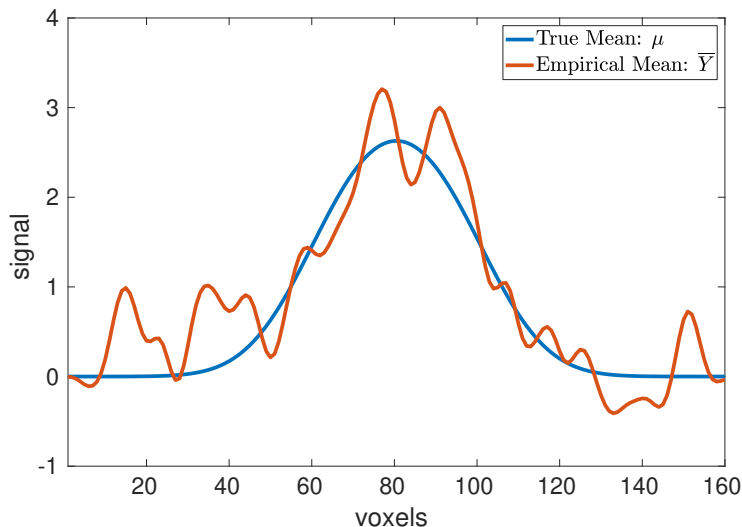
$$Y_n(v) = \mu(v) + \epsilon_n(v).$$

- $\mu(v)$ is the common mean intensity
- $\epsilon_1, \dots, \epsilon_n$ are iid mean zero random images from some unknown multivariate distribution on \mathcal{V}
- Let $\hat{\mu} = \frac{1}{N} \sum_{n=1}^N Y_n$
- let \hat{v}_k be the location of the k th largest local maximum of $\hat{\mu}$

We want to know $\mu(\hat{v}_k)$, but we have $\hat{\mu}(\hat{v}_k)$.

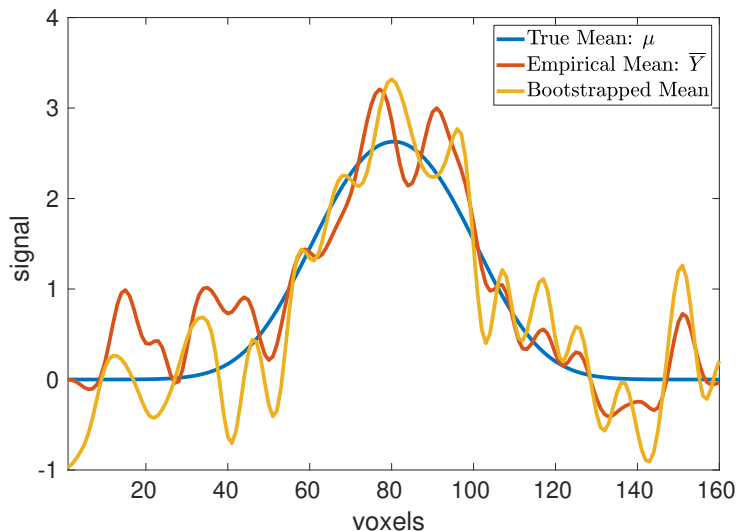
1D Example

20 subjects, $Y_n(t) = \mu(t) + \epsilon_n(t)$, $\hat{\mu} = \bar{Y} = \frac{1}{20} \sum_{n=1}^{20} Y_n$

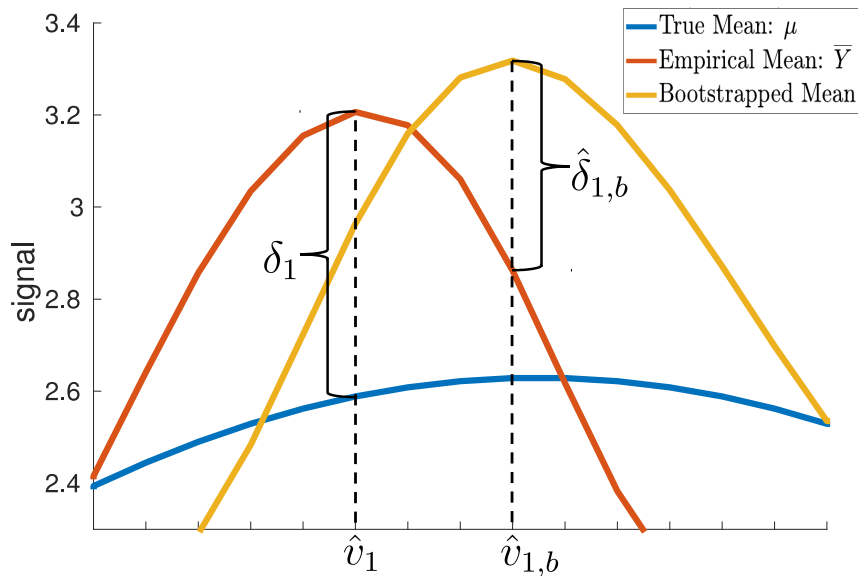


1D Example - Bootstrap Method

20 subjects, $Y_n(t) = \mu(t) + \epsilon_n(t)$, $\hat{\mu} = \bar{Y} = \frac{1}{20} \sum_{n=1}^{20} Y_n$



1D Example - Bootstrap Method



Algorithm 1 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and screening threshold u .
- 2: Let $\hat{\mu} = \frac{1}{N} \sum_{n=1}^N Y_n$ and let K be the number of peaks of $\hat{\mu}$ above u , and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{\mu}$.

Algorithm 2 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and screening threshold u .
- 2: Let $\hat{\mu} = \frac{1}{N} \sum_{n=1}^N Y_n$ and let K be the number of peaks of $\hat{\mu}$ above u , and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{\mu}$.
- 3: **for** $b = 1, \dots, B$ **do**
- 4: Sample $Y_{1,b}^*, \dots, Y_{N,b}^*$ independently with replacement from Y_1, \dots, Y_N .
- 5: Let $\hat{\mu}_b = \frac{1}{N} \sum_{n=1}^N Y_{N,b}^*$ and for $k = 1, \dots, K$, let $\hat{v}_{k,b}$ be the location of the k th largest local maxima of $\hat{\mu}_b$.
- 6: For $k = 1, \dots, K$, let $\hat{\delta}_{k,b} = \hat{\mu}_b(\hat{v}_{k,b}) - \hat{\mu}(\hat{v}_{k,b})$ be an estimate of the bias at the k th largest local maxima.
- 7: **end for**

Algorithm 3 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and screening threshold u .
 - 2: Let $\hat{\mu} = \frac{1}{N} \sum_{n=1}^N Y_n$ and let K be the number of peaks of $\hat{\mu}$ above u , and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{\mu}$.
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 - 7: **end for**
 - 8: For $k = 1, \dots, K$, let $\hat{\delta}_k = \frac{1}{B} \sum_{b=1}^B \hat{\delta}_{k,b}$.
 - 9: **return** $(\hat{\mu}(\hat{v}_1) - \hat{\delta}_1, \dots, \hat{\mu}(\hat{v}_K) - \hat{\delta}_K)$.
-

One-Sample t -statistics/Cohen's d

In neuroimaging we are interested in testing

$$H_0(v) : \mu(v) = 0 \text{ versus } H_1(v) : \mu(v) \neq 0$$

using the one-sample t -statistic:

$$t = \frac{\hat{\mu}\sqrt{N}}{\hat{\sigma}}$$

where

$$\hat{\mu} = \frac{1}{N} \sum_{n=1}^N Y_n, \quad \hat{\sigma}^2 = \frac{1}{N-1} \sum_{n=1}^N (Y_n - \hat{\mu})^2.$$

Effect size is measured via

$$\hat{d}(v) = \frac{\hat{\mu}}{\hat{\sigma}}$$

but this is a biased estimator for the population Cohen's d :

$$d(v) = \frac{\mu}{\sigma}.$$

Unbiased Cohen's d Estimation

This t -statistic $\hat{\mu}\sqrt{N}/\hat{\sigma}$ has a non-central t -distribution with non-centrality parameter $\mu\sqrt{N}/\sigma$ and $N - 1$ degrees of freedom. Thus

$$\mathbb{E}\left[\frac{\hat{\mu}\sqrt{N}}{\hat{\sigma}}\right] = \frac{\mu}{\sigma} \sqrt{\frac{N-1}{2}} \frac{\Gamma((N-2)/2)}{\Gamma((N-1)/2)} = C_N \frac{\mu\sqrt{N}}{\sigma}$$

for $N > 2$, where Γ is the gamma function and C_N is a bias correction factor (Hogben, Pinkham, & Wilk, 1961). So we can use

$$\frac{\hat{\mu}}{\hat{\sigma}C_N}$$

as an unbiased of the population Cohen's d .

Algorithm 4 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and threshold u .
- 2: Let K be the number of peaks of t above u and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{d} = \hat{\mu}/\hat{\sigma}$.

Algorithm 5 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and threshold u .
- 2: Let K be the number of peaks of t above u and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{d} = \hat{\mu}/\hat{\sigma}$.
- 3: **for** $b = 1, \dots, B$ **do**
- 4: Sample $Y_{1,b}^*, \dots, Y_{N,b}^*$ independently with replacement from Y_1, \dots, Y_N .
- 5: Let $\hat{\mu}_b = \frac{1}{N} \sum_{n=1}^N Y_{n,b}^*$ and let $\hat{\sigma}_b^2(v) = \frac{1}{N-1} \sum_{n=1}^N (Y_{n,b}^*(v) - \hat{\mu}_b(v))^2$ for each $v \in \mathcal{V}$.

Algorithm 6 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and threshold u .
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- 3: **for** $b = 1, \dots, B$ **do**
- 4: Sample $Y_{1,b}^*, \dots, Y_{N,b}^*$ independently with replacement from Y_1, \dots, Y_N .
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- 6: For $k = 1, \dots, K$, let $\hat{v}_{k,b}$ be the location of the k th largest local maxima of $\hat{d}_b = \hat{\mu}_b/\hat{\sigma}_b$.
- 7: Let $\hat{\delta}_{k,b} = (\hat{d}_b(\hat{v}_{k,b}) - \hat{d}(\hat{v}_{k,b}))/C_N$ be an estimate of the bias.
- 8: **end for**

Algorithm 7 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and threshold u .
 - 2: Let K be the number of peaks of t above u and for $k = 1, \dots, K$, let \hat{v}_k be the location of the k th largest maxima of $\hat{d} = \hat{\mu}/\hat{\sigma}$.
 - 3: **for** $b = 1, \dots, B$ **do**
 - 4: Sample $Y_{1,b}^*, \dots, Y_{N,b}^*$ independently with replacement from Y_1, \dots, Y_N .
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 - 7: Let $\hat{\delta}_{k,b} = (\hat{d}_b(\hat{v}_{k,b}) - \hat{d}(\hat{v}_{k,b}))/C_N$ be an estimate of the bias.
 - 8: **end for**
 - 9: For $k = 1, \dots, K$, let $\hat{\delta}_k = \frac{1}{B} \sum_{b=1}^B \hat{\delta}_{k,b}$
 - 10: **return** $(\hat{d}(\hat{v}_1)/C_N - \hat{\delta}_1, \dots, \hat{d}(\hat{v}_K)/C_N - \hat{\delta}_K)$.
-

To infer on μ instead of μ/σ can just use

$$\hat{\delta}_{k,b} = \hat{\mu}_b(\hat{v}_{k,b}) - \hat{\mu}(\hat{v}_{k,b})$$

- Circular inference estimates are: $\hat{d}(\hat{v}_1)/C_N, \dots, \hat{d}(\hat{v}_K)/C_N$.
- For data-splitting, we first divide the images into two groups: $Y_1, \dots, Y_{N/2}$ and $Y_{N/2+1}, \dots, Y_N$. Then find the peaks using the first half of the subjects and estimate the values at those peaks using the second half of the subjects.

Let Y be an N -dimensional random image such that for each $v \in \mathcal{V}$

$$Y(v) = X\beta(v) + \epsilon(v)$$

- $N \times p$ design matrix X
- parameter vector $\beta(v) \in \mathbb{R}^p$
- $\epsilon(v) = (\epsilon_1(v), \dots, \epsilon_N(v))^T$ is the random image of the noise

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We are interested in testing

$$H_0(v) : C\beta(v) = 0 \text{ versus } H_1(v) : C\beta(v) \neq 0$$

for some contrast matrix $C \in \mathbb{R}^{m \times p}$. We can test this at each voxel with the usual F -test,

$$F(v) = \frac{(C\hat{\beta}(v))^T (C(X^T X)^{-1} C^T)^{-1} (C\hat{\beta}(v)) / m}{\hat{\sigma}(v)^2} \quad (1)$$

where $\hat{\beta}(v) = (X^T X)^{-1} X^T Y$ and $\hat{\sigma}^2(v)$ is the error variance. Under the alternative has a non-central F -distribution.

Alternative F statistic - General Linear Hypothesis

Another (common) way to define the F -statistic is as follows. Let Ω denote the overall model and let $\omega \subset \Omega$ denote some sub-model with p_0 degrees of freedom. Then

$$F = \frac{(\text{RSS}_\omega - \text{RSS}_\Omega)/m}{\text{RSS}_\Omega/N - p}$$

where $m = p - p_0$ and

$$\text{RSS}_\Omega = \sum_{n=1}^N (Y_n - X\hat{\beta})^2 \text{ and } \text{RSS}_\omega = \sum_{n=1}^N (Y_n - X\hat{\beta}_0)^2$$

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where $m = p - p_0$ and

$$\text{RSS}_\Omega = \sum_{n=1}^N (Y_n - X\hat{\beta})^2 \text{ and } \text{RSS}_\omega = \sum_{n=1}^N (Y_n - X\hat{\beta}_0)^2$$

Theorem (non-obvious!): Taking $\omega = \{\beta : C\beta = 0\}$ these F -statistic forms are equivalent. This is known as the general linear hypothesis. Great stackexchange post on proving this: <https://stats.stackexchange.com/questions/17207/general-linear-hypothesis-test-statistic-equivalence-of-two-expressions>

$$R^2 = 1 - \frac{\text{RSS}_\Omega}{\text{RSS}_\omega}.$$

which is commonly reported in papers.

$$F = \frac{(\text{RSS}_\omega - \text{RSS}_\Omega)/m}{\text{RSS}_\Omega/(N-p)} = \frac{N-p}{m} \left(\frac{\text{RSS}_\omega}{\text{RSS}_\Omega} - 1 \right) = \frac{N-p}{m} \left(\frac{1}{1-R^2} - 1 \right)$$

which implies that

$$R^2 = 1 - \left(\frac{m}{N-p} F + 1 \right)^{-1} = 1 - \frac{N-p}{mF + N-p} = \frac{mF}{mF + N-p}.$$

This gives us an easy way of computing the R^2 value in terms of the F -statistic.

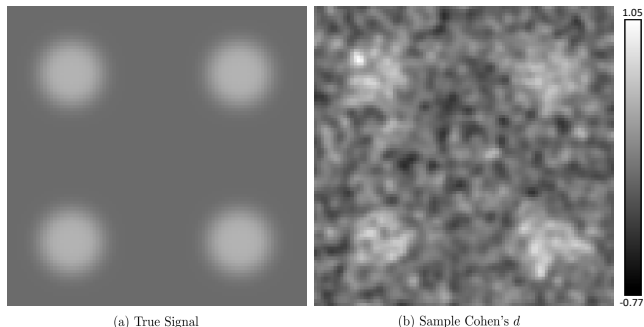
Algorithm 8 Non-Parametric Bootstrap Bias Calculation

- 1: **Input:** Images Y_1, \dots, Y_N , the number of bootstrap samples B and threshold u .
- 2: Let $\hat{\beta} = \hat{\beta}(X, Y) = (X^T X)^{-1} X^T Y$ and let $\hat{\epsilon} = Y - X\hat{\beta}$ be the residuals.
- 3: For each $n = 1, \dots, N$, let $r_n = \hat{\epsilon}_n / \sqrt{1 - p_n}$ be the modified residuals, where $p_n = (X(X^T X)^{-1} X^T)_{nn}$. Let $\bar{r} = \frac{1}{N} \sum_{n=1}^N r_i$ be their mean.
- 4: **for** $b = 1, \dots, B$ **do**
- 5: Sample $\epsilon_{1,b}^*, \dots, \epsilon_{N,b}^*$ independently with replacement from $r_1 - \bar{r}, \dots, r_N - \bar{r}$ and let $\epsilon_b^* = (\epsilon_{1,b}^*, \dots, \epsilon_{N,b}^*)^T$ and set $Y_b^* = X\hat{\beta} + \epsilon_b^*$.
- 6: Let F_b^* be the bootstrapped F -statistic image computed using Y_b^* . Let R_b^2 be the bootstrapped partial R^2 image and set $\hat{\delta}_{k,b} = R_b^2(\hat{v}_{k,b}) - R^2(\hat{v}_{k,b})$ to be the estimate of the bias.
- 7: **end for**
- 8: For $k = 1, \dots, K$, let $\hat{\delta}_k = \frac{1}{B} \sum_{b=1}^B \hat{\delta}_{k,b}$.
- 9: **return** $(R^2(\hat{v}_1) - \hat{\delta}_1, \dots, R^2(\hat{v}_K) - \hat{\delta}_K)$.

Simulations

Simulations - Cohen's d

All simulations generated using code from the RFTtoolbox
<https://github.com/BrainStatsSam/RFTtoolbox> (avoiding edge problems)



- Panel (a) illustrates a slice through the true signal (actually 9 peaks only 4 shown).
- Panel (b) illustrates the same slice through the one sample Cohen's d for 50 subjects. Noise: Gaussian random field with FWHM 6.

Traditionally, one estimates a common θ with estimators $\hat{\theta}_1, \dots, \hat{\theta}_K$ however we have estimators $\hat{\theta}_1, \dots, \hat{\theta}_K$ of parameters $\theta_1, \dots, \theta_K$ where K is the number of significant peaks that are found over all realizations. As such we instead define

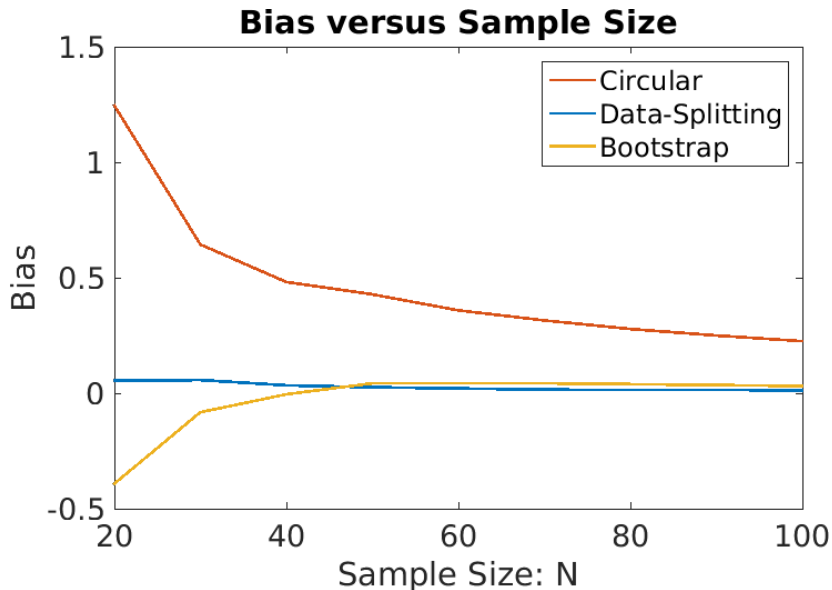
$$\tilde{\theta}_k = \hat{\theta}_k - \theta_k$$

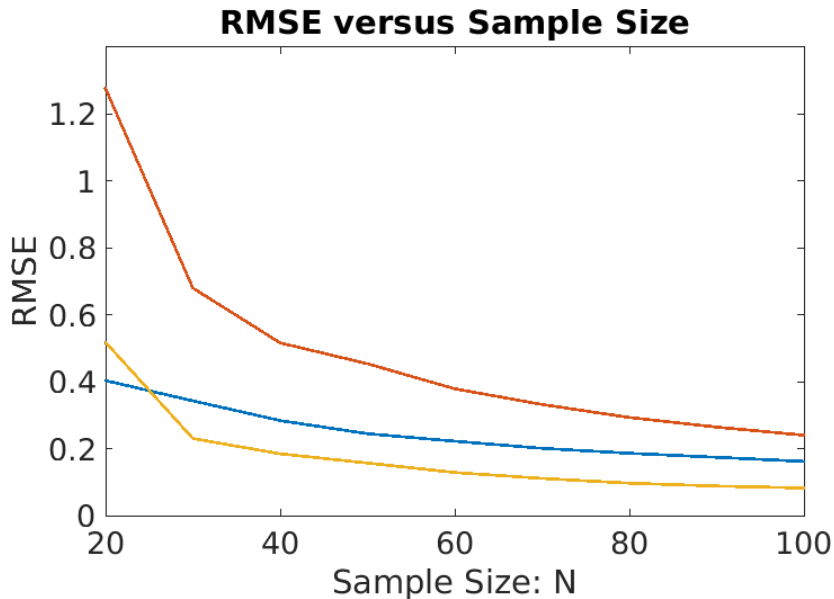
and use the fact that the noise-free value of $\tilde{\theta}_k$ is 0 for each k .

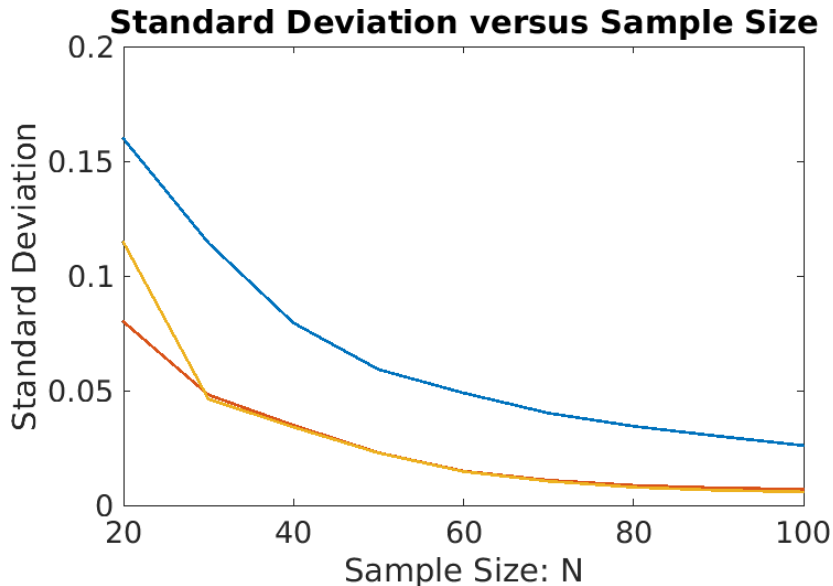
$$\begin{aligned} \text{MSE} &= \frac{1}{K} \sum_{k=1}^K (\tilde{\theta}_k - 0)^2 \\ &= \frac{1}{K} \sum_{k=1}^K \left(\tilde{\theta}_k - \frac{1}{K} \sum_{k=1}^K \tilde{\theta}_k \right)^2 + \left(\frac{1}{K} \sum_{k=1}^K \tilde{\theta}_k \right)^2 \end{aligned}$$

Simulation Evaluation and Thresholding

- We evaluate our methods for $N = \{20, 30, \dots, 100\}$.
- For each N we generate 1,000 realizations and compare the performance of the three methods across realizations.
- we generate 5,000 null t_{N-1} random fields take the 95% quantile of the distribution of the maximum to provide a voxelwise threshold.

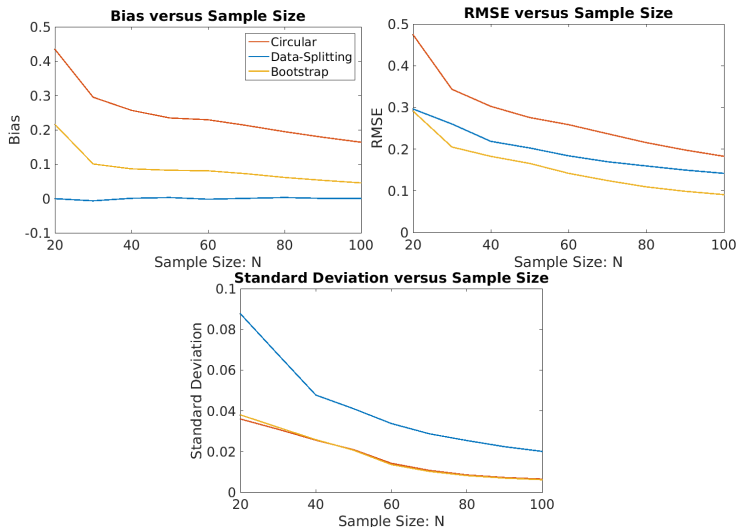






Results - Mean estimation - simulations

Thresholding using Cohen's d but estimating using μ gives similar results (see below) and the results for GLM simulations are also similar.



Big Data Validation

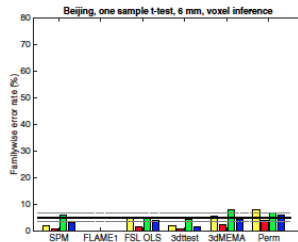
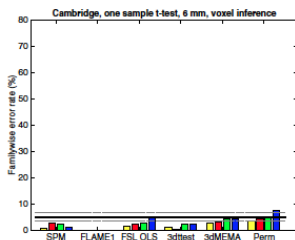
Data and Testing

- 8940 total subjects from the UK biobank. We have task fMRI and VBM data from all subjects
- We test the one-sample methods using the task fMRI data and the GLM methods using the VBM data (as the R^2 effect sizes are very small for the task fMRI data sets)
- For the task-fMRI data we estimate Cohen's d or μ .
- For the VBM data we regress against age, sex and an intercept and compute the partial R^2 for age.
- Set aside 4000 subjects to compute a ground truth and divide the rest into 4940/ N groups of size $N = 20, 50, 100$.
- Actually for the VBM data we take $N = 50, 100, 150$ as the effect size is lower

We recommend this type of testing framework for all statistical methods.

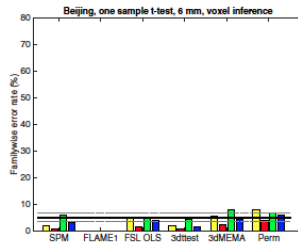
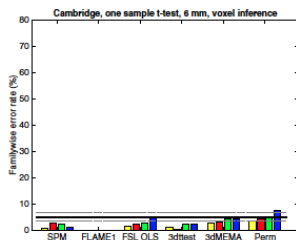
Thresholding

- We threshold using voxelwise RFT. This doesn't have the same problems as clusterwise inference as it doesn't make the same assumptions.



Thresholding

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- Our method independent of the threshold.
- For the big data analysis we do permutation is very computational so is not practical.
- But permutation testing can be used to compute the voxelwise threshold when doing a general analysis.

Cohen's d ground truth

Computing the ground truth is difficult due to memory constraints. So you have load images sequentially. Let \mathcal{D} be the set of all possible voxels. Typically \mathcal{D} is a $91 \times 109 \times 91$ grid. Define

$$M_n(v) = \begin{cases} 1 & \text{if subject } n \text{ has data at } v \\ 0 & \text{otherwise} \end{cases}$$

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Take $\mathcal{S} \subset \{1, \dots, 8940\}$ of size 4000 and let

$$\mu(v) = \frac{\sum_{n \in \mathcal{S}} Y_n(v) M_n(v)}{\sum_{n \in \mathcal{S}} M_n(v)} \times \mathbf{1}(M_n(v) = 1 \text{ for at least } 100 \text{ } n \in \mathcal{S})$$

$$\sigma^2(v) = \frac{\sum_{n \in \mathcal{S}} (Y_n - \mu(v))^2 M_n(v)}{\sum_{n \in \mathcal{S}} M_n(v) - 1} \times \mathbf{1}(M_n(v) = 1 \text{ for at least } 100 \text{ } n \in \mathcal{S}),$$

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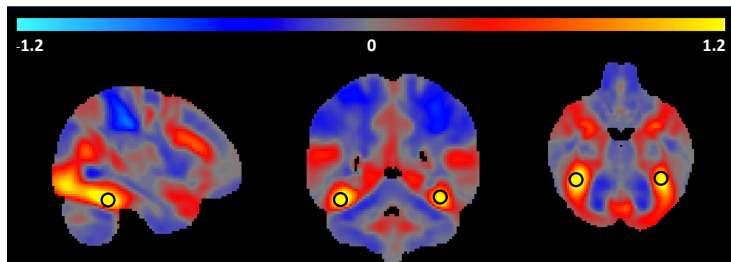
$$\sigma^2(v) = \frac{\sum_{n \in \mathcal{S}} (Y_n - \mu(v))^2 M_n(v)}{\sum_{n \in \mathcal{S}} M_n(v) - 1} \times \mathbf{1}(M_n(v) = 1 \text{ for at least } 100 \text{ } n \in \mathcal{S}),$$

and the **ground truth Cohen's d** estimate as

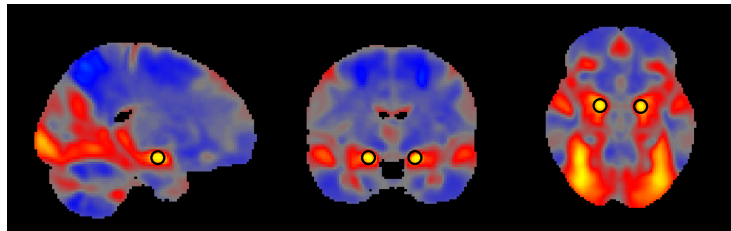
$$d(v) = \frac{\mu(v)}{\sigma(v)}.$$

Finally each of these are additionally masked with the 2mm MNI brain mask.

Cohen's d Ground Truth Slices



(a) Top 2 peaks



(b) 3rd and 4th Highest Peaks

Illustrating the Winner's Curse

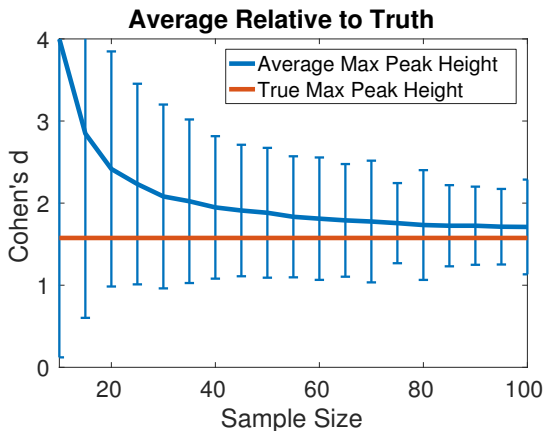


Figure 2: Comparing the maximum values at small sample Cohen's d (over $4940/N$ groups) to the max ground truth value.

GLM ground truth

For now assume that no data is missing and that we have

- $N_{\text{all}} = 4000$ subjects
- an $N_{\text{all}} \times p$ design matrix $X = (x_1, \dots, x_{N_{\text{all}}})^T$
- V is the number of voxels in each subject image Y_n
- Y be the $N_{\text{all}} \times V$ matrix of all the subject images

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For $Y = X\beta + \epsilon$, we want to compute

$$\hat{\beta} = (X^T X)^{-1} X^T Y,$$

at each voxel. For each $v \in \mathcal{V}$,

$$X^T Y(v) = (x_1, \dots, x_{N_{\text{all}}}) \begin{pmatrix} Y_1(v) \\ \vdots \\ Y_{N_{\text{all}}}(v) \end{pmatrix} = \sum_{n=1}^{N_{\text{all}}} Y_n(v) x_n,$$

$$\hat{\sigma}^2 = (N_{\text{all}} - p)^{-1} \sum_{n=1}^{N_{\text{all}}} (Y_n - x_n^T \hat{\beta})^2.$$

and this allows F and R^2 to be calculated

For each $v \in \mathcal{V}$,

$$X^T Y(v) = (x_1, \dots, x_{N_{\text{all}}}) \begin{pmatrix} Y_1(v) \\ \vdots \\ Y_{N_{\text{all}}}(v) \end{pmatrix} = \sum_{n=1}^{N_{\text{all}}} Y_n(v) x_n,$$

Can compute $\hat{\beta} = (X^T X)^{-1} X^T Y$ from this and estimate

$$\hat{\sigma}^2 = (N_{\text{all}} - p)^{-1} \sum_{n=1}^{N_{\text{all}}} (Y_n - x_n^T \hat{\beta})^2.$$

and this allows F and R^2 to be calculated.

GLM ground truth with missingness

Let $C(v) := \{n : M_n(v) = 1\}$. Then for each voxel v we need to compute the complete case estimate

$$\hat{\beta}(v) = (X_{C(v)}^T X_{C(v)})^{-1} X_{C(v)}^T Y_{C(v)}.$$

The first and second parts of this expression can be computed as

$$(X_{C(v)}^T X_{C(v)})^{-1} = \left(\sum_{n=1}^{N_{\text{all}}} M_n(v) x_n x_n^T \right)^{-1}$$

and

$$X_{C(v)}^T Y_{C(v)} = \sum_{n=1}^{N_{\text{all}}} M_n(v) Y_n(v) x_n$$

$\hat{\sigma}^2$, F and R^2 can similarly be computed.

Missingness Assumption

Theorem

At each voxel, suppose that $Y = X\beta + \epsilon$ for some zero mean random vector ϵ and that R is the missingness information of Y and that no X variables are missing. If $R \perp\!\!\!\perp Y|X$ then

$$\hat{\beta} = \left(X_{O(R)}^T X_{O(R)} \right)^{-1} X_{O(R)}^T Y_{O(R)}$$

is an unbiased estimate of β .

Proof.

We have: $Y_{O(R)} = X_{O(R)}\beta + \epsilon_{O(R)}$. Integrating we find that,

$$\begin{aligned} \int \hat{\beta} d\pi(Y, R, X) &= \int \hat{\beta} d\pi(Y_{O(R)}, R, X) = \int \hat{\beta} d\pi(Y_{O(R)}|R, X) d\pi(R, X) \\ &= \int \beta d\pi(R, X) = \beta. \end{aligned}$$

Missingness Assumption

Proof.

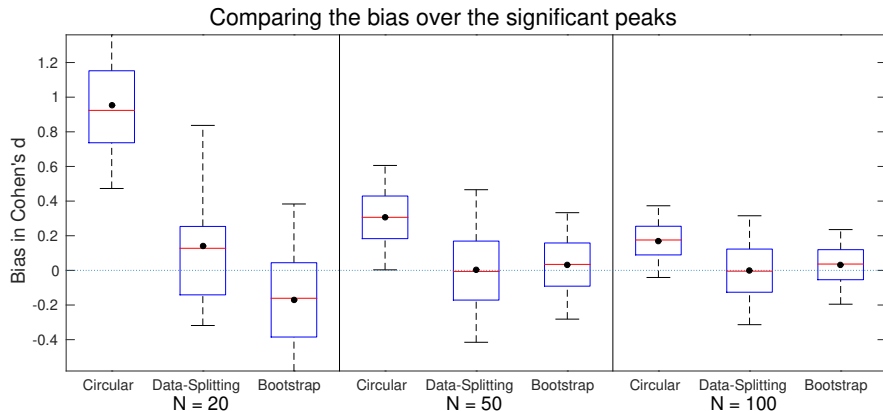
This follows as

$$\begin{aligned}\int \hat{\beta} d\pi(Y_{O(R)}|R, X) &= \int \left(X_{O(R)}^T X_{O(R)}\right)^{-1} X_{O(R)}^T Y_{O(R)} d\pi(Y_{O(R)}|R, X) \\ &= \left(X_{O(R)}^T X_{O(R)}\right)^{-1} X_{O(R)}^T \int Y_{O(R)} d\pi(Y_{O(R)}|R, X) \\ &= \left(X_{O(R)}^T X_{O(R)}\right)^{-1} X_{O(R)}^T \int Y_{O(R)} d\pi(Y_{O(R)}|X) \\ &= \left(X_{O(R)}^T X_{O(R)}\right)^{-1} X_{O(R)}^T X_{O(R)} \beta = \beta.\end{aligned}$$

where the third equality uses the fact that $R \perp\!\!\!\perp Y|X$. □

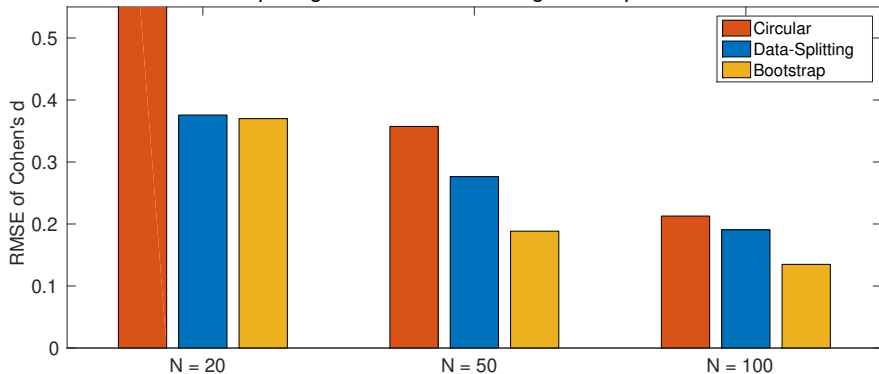
Results

One Sample Cohen's d - Bias

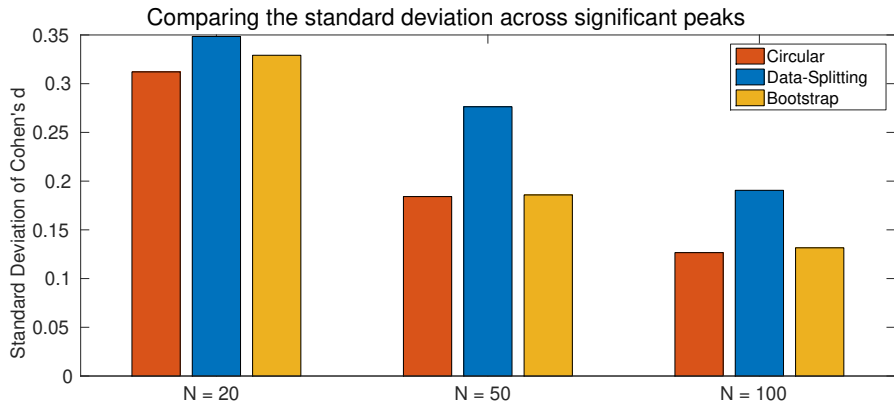


One Sample Cohen's d - RMSE

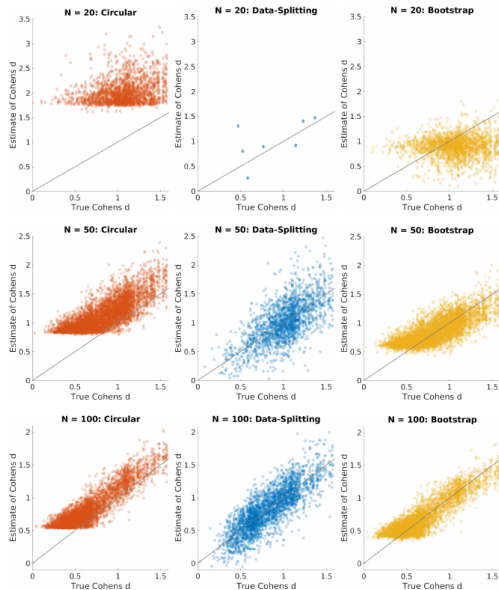
Comparing the RMSE across significant peaks



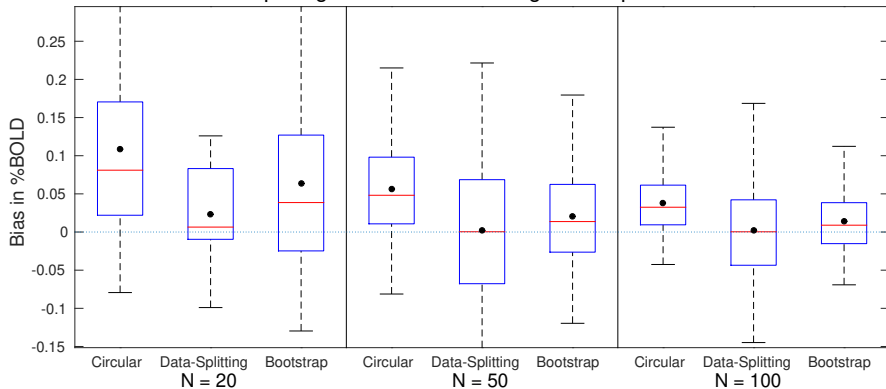
One Sample Cohen's d - Standard Deviation



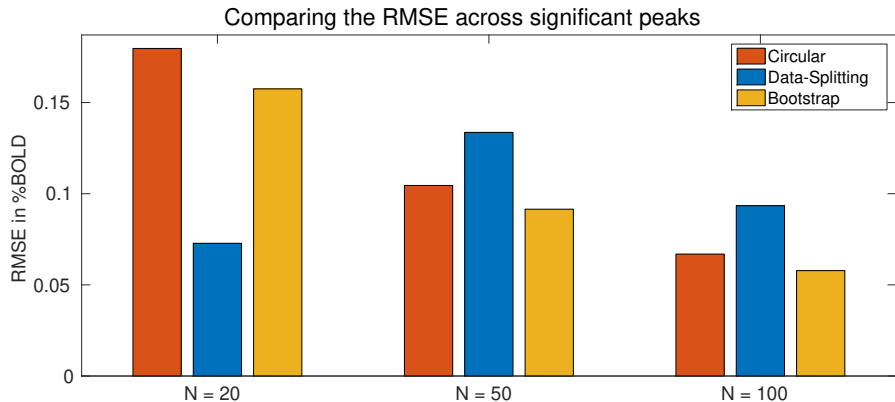
One Sample Cohen's d - Estimates vs Ground truth

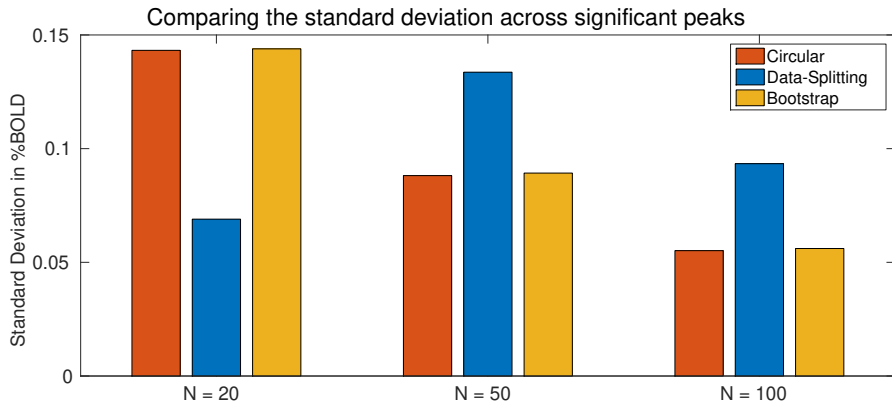


Comparing the bias over the significant peaks

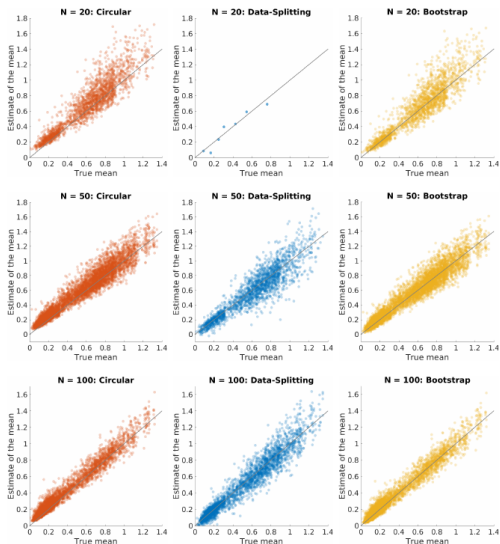


Mean estimation - RMSE

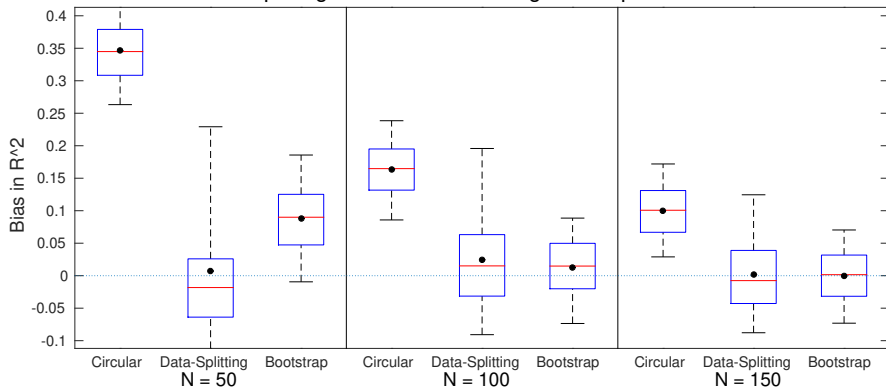




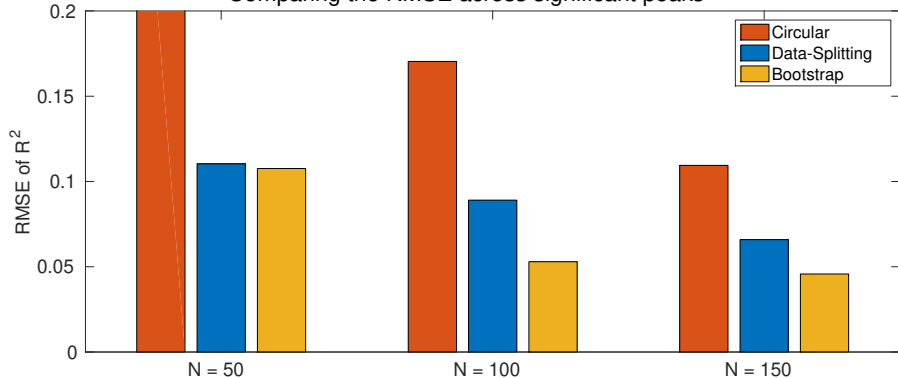
Mean estimation - Estimates versus Ground truth



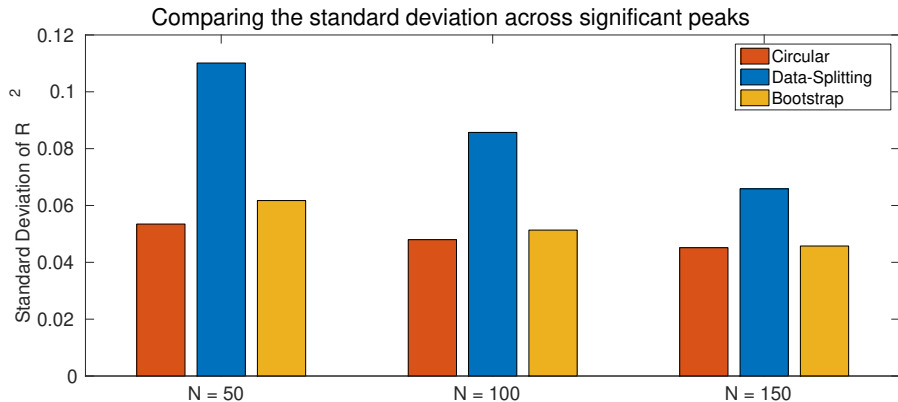
Comparing the bias over the significant peaks



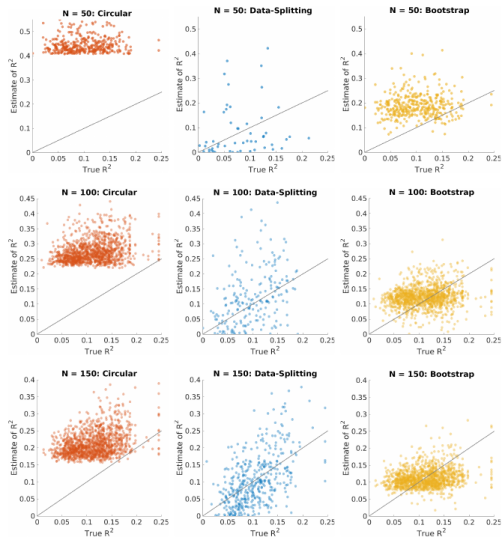
Comparing the RMSE across significant peaks



R^2 - Standard Deviation



R^2 - Estimates versus Ground truth



Power Analyses

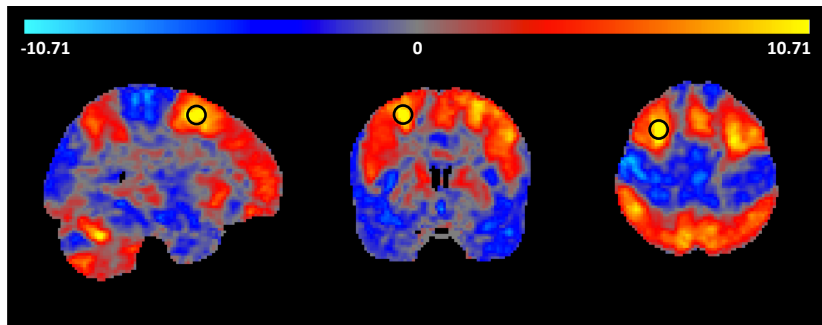
One-Sample t-statistic power

Given a potential future sample size N' and an estimate of the non-centrality parameter: λ , the power is:

$$\mathbb{P}(T_{N'-1,\lambda} > t_{1-\alpha,N'-1})$$

where $t_{1-\alpha,N'-1}$ is chosen such that $\mathbb{P}(T_{N'-1,0} > t_{1-\alpha,N'-1}) = \alpha$ and $T_{N'-1,\lambda}$ has a non-central T distribution with $N' - 1$ degrees of freedom and non-centrality parameter λ .

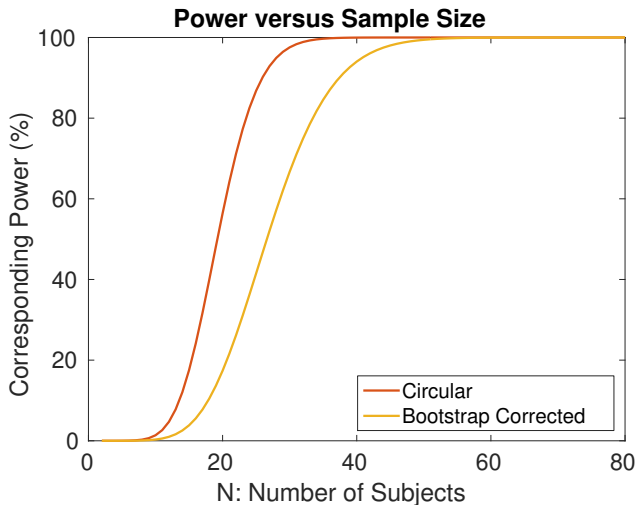
Working Memory Example



- One-sample t -statistic for 80 subjects from the HCP.
- Activation in the Medial Frontal Gyrus.
- At the maximum the observed (circular) Cohen's d is 1.519, while the bootstrap-corrected value is 1.161
- The observed %BOLD change there is %0.450 and corrected estimate is %0.433.

Power graph

At the maximum the observed (circular) Cohen's d is 1.519, while the bootstrap-corrected value is 1.161. So we can generate a power graph:



$$F = \frac{(C\hat{\beta})^T (C(X^T X)^{-1}C^T)^{-1} (C\hat{\beta})/m}{\hat{\sigma}^2}$$

has a non-central F distribution with non-centrality parameter

$$(C\beta)^T (C(X^T X)^{-1}C^T)^{-1} (C\beta)/\sigma^2$$

and degrees of freedom m and $N - p$.

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and degrees of freedom m and $N - p$. Define Cohen's f^2 to be

$$f^2 := \frac{R^2}{1 - R^2} = \frac{m}{N - p} F = \frac{(C\hat{\beta})^T (C(\frac{1}{N-p}X^T X)^{-1}C^T)^{-1} (C\hat{\beta})}{\hat{\sigma}^2}.$$

Suppose

$$Y_N = X_N\beta + \epsilon^N$$

where $X_N = [x_1, \dots, x_N]^T$ is the design matrix, for $\{x_n\}_{n \in \mathbb{N}} \in \mathbb{R}^p$ a sequence of finite variance, iid random vectors (independent of the noise process) each with multivariate distribution D .

Also suppose that the noise $\epsilon^N = (\epsilon_1, \dots, \epsilon_N)^T$ has finite variance.

Theorem

$$\begin{aligned}\frac{1}{N} X_N^T X_N &\xrightarrow{a.s.} \mathbb{E}[x_1 x_1^T] \\ \hat{\sigma}_N^2 &\xrightarrow{a.s.} \sigma \\ \hat{\beta}_N &\xrightarrow{a.s.} \beta\end{aligned}$$

Proof.

See the supplementary material of (Davenport & Nichols, 2019). □

Let f_N^2 be Cohen's f^2 for the N th model. Then combining the above results,

$$\begin{aligned}f_N^2 &= \frac{(C\hat{\beta}_N)^T (C(\frac{1}{N-p} X^T X)^{-1} C^T)^{-1} (C\hat{\beta}_N)}{\hat{\sigma}_N^2} \\ &\xrightarrow{a.s.} f_p^2 := \frac{(C\beta)^T (C(\mathbb{E}[x_1 x_1^T])^{-1} C^T)^{-1} (C\beta)}{\sigma^2}\end{aligned}$$

as $N \rightarrow \infty$. This also implies almost sure convergence of R^2 .

Power in the GLM

Given N' subjects with design matrix X' whose rows are iid with distribution D . then for N' is sufficiently large, we can obtain reasonable estimates of the power.

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$$\frac{(C\beta)^T (C(X'^T X')^{-1} C^T)^{-1} (C\beta)}{\sigma^2} =$$
$$N' \frac{(C\beta)^T (C(\frac{1}{N'} X'^T X')^{-1} C^T)^{-1} (C\beta)}{\sigma^2} \approx N' f_p^2 \approx N' f^2$$

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Then the power is:

$$\mathbb{P}(F_{m, N'-p, \lambda} > f_{1-\alpha, m, N'-p})$$

where $f_{1-\alpha, m, N'-p}$ is chosen such that $\mathbb{P}(F_{m, N'-p, 0} > f_{1-\alpha, m, N'-p}) = \alpha$ and where $F_{m, N'-p, \lambda}$ has a non-central F distribution with m and $N' - p$ degrees of freedom and non-centrality parameter λ .

Conclusion

Conclusion and Future Work

- We provide a method for dealing with the winner's curse which outperforms existing methods in terms of RMSE.
- Can also be used to estimate the maximum rather than the maximum at a given location.
- Would be cool to develop an RFT method potentially using stuff from (Cheng & Schwartzman, 2015), but this is probably quite difficult!
- Other cool method: (Benjamini & Meir, 2014) which works for voxelwise inference. So far only developed in certain settings but its not hard to extend.
- Interesting to work on an estimate for the cluster mass statistic which is also commonly reported.

- Code and scripts to reproduce figures available in SIbootstrap toolbox.
- Simulations and thresholding were performed using RFTtoolbox available at
- Can find both at `sjdavenport.github.io/software`.

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- Cheng, D., & Schwartzman, A. (2015). Distribution of the height of local maxima of Gaussian random fields. *Extremes*, 18(2), 213–240. doi: 10.1007/s10687-014-0211-z
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- Hogben, D., Pinkham, R. S., & Wilk, M. B. (1961). The moments of the non-central t-distribution. *Biometrika*, 465–468.