Robust Outlier Detection

for Low and High-Dimensional Neuroimaging Data with Principal Components Analysis and Split-Half Resampling

Derek Beaton

Rotman Research Institute, Baycrest Health Sciences

Twitter: @derek___beaton (that's two underscores!)
Github for today: http://github.com/derekbeaton/ours

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The other authors

Kelly M Sunderland, Abiramy Uthirakumaran, Stephen R Arnott, Robert Bartha, Sandra E Black, Leanne Casaubon, Morris Freedman, Richard H Swartz, Sean Symons, ONDRI Investigators, Malcolm A Binns, and Stephen C Strother

Acknowledgements



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Acknowledgements

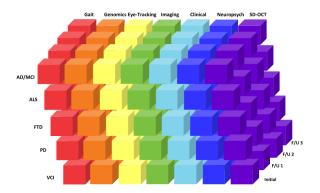


Ontario neurodegenerative disease research initiative (ONDRI)

ONDRI Outlier detection

1

Introduction



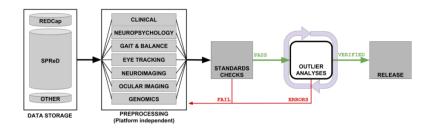
The ONDRI "cube" (Farhan et al., 2017). Ontario-wide, multi-site, longitudinal, multi-cohort, "deep-phenotyping". Today's focus: Alzheimer's (AD/MCI) and Vascular Cognitive Impairment (VCI)

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- Almost everything multivariate with varying complexities
- How to ensure data are of highest quality?



The ONDRI preprocessing to release pipeline

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- Why a new approach?

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- Why a new approach?
 - There are substantial limitations of existing methods

Distances PCA Resampling

2

PCA+SHR background

- New framework:
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 - Principal components analysis (PCA) plus
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- The BIG goal
 - Provide flexible & robust multivariate outlier detection

Distances PCA Resampling

Mahalanobis distances (MD)

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- Orthogonal distances

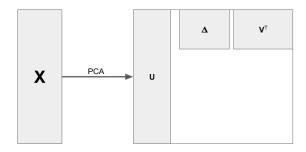
- Mahalanobis distances (MD)
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 - MD scaled by explained variance per component
- Orthogonal distances
 - We'll bring these up later

Distances PCA Resampling

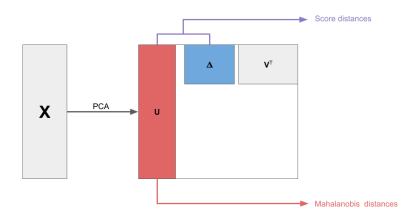
PCA

X

PCA



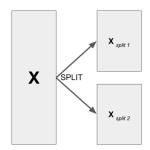
PCA



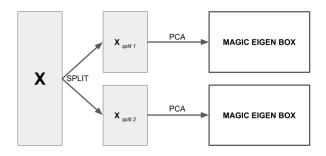
PCA



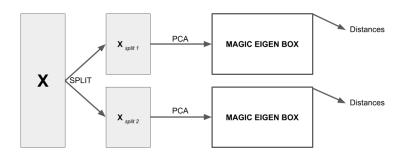
Single split PCA



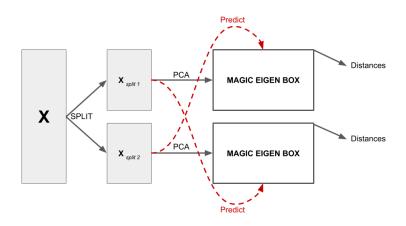
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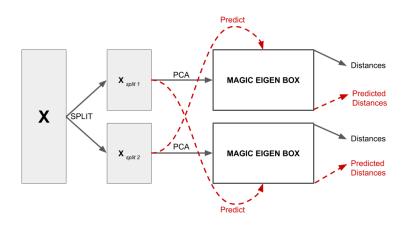
Single split PCA



Prediction from split PCAs



Prediction from split PCAs



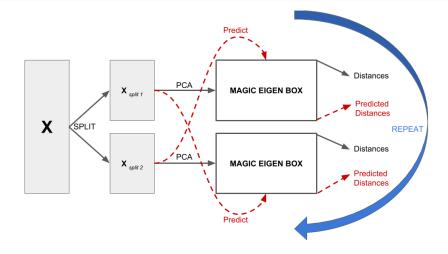
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 - Standard MD cannot be computed
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 - Predicted distances can always be computed
- But: just one pass at split PCA is not enough



Introduction
PCA+SHR background
PCA+SHR walkthrough
Discussion
Bonus material

Distances PCA Resampling

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- Estimates of reproducibility
 - Will come up later with orthogonal distances (OD).

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Low dimensional PCA+SHR High dimensional PCA+SHR

3

PCA+SHR walkthrough

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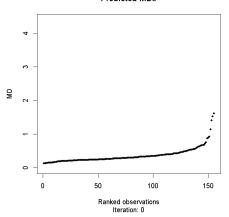
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- N = 1000 iterations

Predicted MD distributions

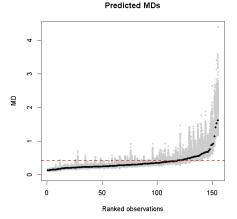




Predicted MD distributions

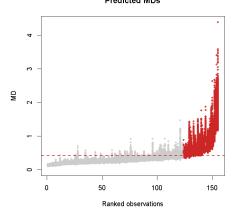
MD distribution threshold



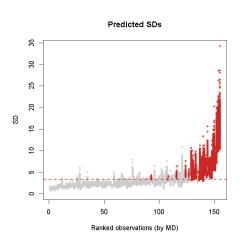


Mahalanobis distribution outliers





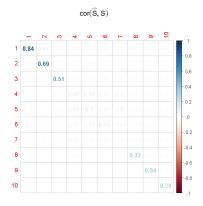
Score distribution outliers



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 - Find reproducible & robust subspace (components)



Median R^2 between split scores and predicted scores over 1000 iterations

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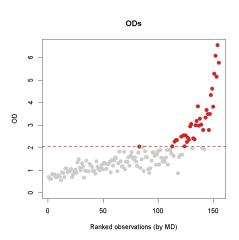
- We can use those to rebuild a robust version of X:
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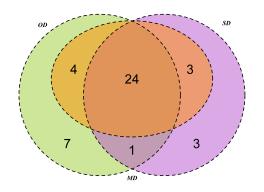
- We can use those to rebuild a robust version of X:
 - X original data
 - X' reconstructed from just first three components
- How much do observations change between X and robust X'?

Orthogonal distance (OD)

ullet Distance between observation and itself for ${f X}$ and robust ${f X}'$

Orthogonal distance outliers





I = 161, Outliers = 42.

Summary of PCA+SHR

Build (predictive) distributions

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- Find a reproducible subspace

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- Build (predictive) distributions
- Find a reproducible subspace
- Three types of distances & outliers

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- High dimensional, low sample size
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- PCA+SHR works the same regardless of size

• I = 109 observations from the AD/MCI cohort

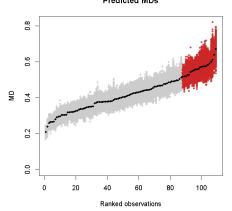
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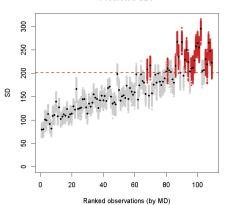
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 - Resting state fMRI
 - "Seed" voxel used to estimate whole-brain connectivity
 - OPPNI pipeline (Churchill et al., 2015)



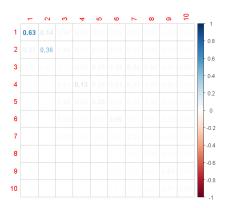


Predicted SDs

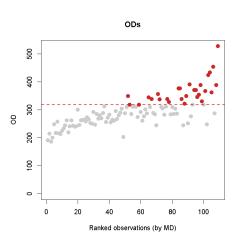


Subspace

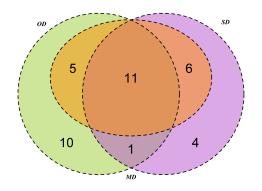




Only the first 10 of 108 Components shown



OD computed from two components



$$I = 109$$
, Outliers = 37.

4

Discussion

Conclusions

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- Different distances provide different perspectives
- Thresholds play a role in (non) overlap
 - High: Tends to intersect, find most outlying individuals
 - Low: Find most outlying & unique outliers to each type

Benefits

• PCA+SHR is flexible, overcomes limits of other methods

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- PCA+SHR is flexible, overcomes limits of other methods
 - Most methods don't work for high dimensional/low sample size data
- A lot of information available
 - Multiple distances, distributions to help make decisions

Limitations

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- Lots of options
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 - Distributions, point, spread?
 - But we provide defaults in the software that work well
- Can be slow

Current & future work

• Speed up (e.g., parallelization)

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- Rolling this out within ONDRI

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- PCA+SHR easily extends to virutally any data type
 - Continuous, categorical, ordinal, or mixed
 - Beaton et al., (2018) extended MCD [1] to any data type

References

- [1] Beaton, D., Sunderland, K. M., Levine, B., Mandzia, J., Masellis, M., Swartz, R. H., . . . & Strother, S. C. (2018). Generalization of the minimum covariance determinant algorithm for categorical and mixed data types. bioRxiv, 333005.
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- [4] Hubert, M., & Debruyne, M. (2010). Minimum covariance determinant. Wiley interdisciplinary reviews: Computational statistics, 2(1), 36-43.
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Questions, comments, complaints?

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 - Higher level overview of ONDRI curation-through-release pipeline

5

Bonus material

PCA

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The SVD of a matrix **X** of size $I \times J$ (at least column-wise centered, i.e., covariance)

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 (1)

The SVD of a matrix \mathbf{X} of size $I \times J$ (at least column-wise centered, i.e., covariance)

$$\mathbf{X} = \mathbf{U} \mathbf{\Delta} \mathbf{V}^{\mathsf{T}} \tag{1}$$

with the following properties

• Rank is L where $L \leq \min(I, J)$

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- **U** is $I \times L$ (left singular vectors; rows of **X**)
- **V** is $J \times L$ (right singular vectors; columns of **X**)
- **U** and **V** are orthonormal: $\mathbf{U}^{\mathsf{T}}\mathbf{U} = \mathbf{I} = \mathbf{V}^{\mathsf{T}}\mathbf{V}$

Score distances

Given
$$\mathbf{X} = \mathbf{U} \mathbf{\Delta} \mathbf{V}^{\mathsf{T}}$$

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Given
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• Sum of squared component (factor) scores

$$S = \operatorname{diag}\{(U\Delta)(U\Delta)^{T}\} = \operatorname{diag}\{(XV)(XV)^{T}\}$$
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- Score distances (SD) are defined as:

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Mahalanobis

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Sum of squared singular vectors

$$\mathbf{M} = \operatorname{diag}\{\mathbf{U}\mathbf{U}^{T}\} = \operatorname{diag}\{(\mathbf{X}\mathbf{V}\boldsymbol{\Delta}^{-1})(\mathbf{X}\mathbf{V}\boldsymbol{\Delta}^{-1})^{T}\}$$
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(3)

Single split PCA

For some subset H and its complement \bar{H} we have two PCAs:

$$\mathbf{X}_{H} = \mathbf{U}_{H} \mathbf{\Delta}_{H} \mathbf{V}_{H}^{T}$$

$$\mathbf{X}_{\bar{H}} = \mathbf{U}_{\bar{H}} \mathbf{\Delta}_{\bar{H}} \mathbf{V}_{\bar{H}}^{T}$$

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$$(4)$$

Size of H could be $\alpha=.5$ (split half) or e.g., $\alpha=.9$ (90-10)

Predicted distances

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Predicted SD:

$$\widehat{\mathbf{S}}_{H} = \operatorname{diag}\{(\mathbf{X}_{H}\mathbf{V}_{\bar{H}})(\mathbf{X}_{H}\mathbf{V}_{\bar{H}})^{T}\}$$

$$\widehat{\mathbf{S}}_{\bar{H}} = \operatorname{diag}\{(\mathbf{X}_{\bar{H}}\mathbf{V}_{H})(\mathbf{X}_{\bar{H}}\mathbf{V}_{H})^{T}\}$$
(5)

Predicted distances

Predicted SD:

$$\widehat{\mathbf{S}}_{H} = \operatorname{diag}\{(\mathbf{X}_{H}\mathbf{V}_{\bar{H}})(\mathbf{X}_{H}\mathbf{V}_{\bar{H}})^{T}\}$$

$$\widehat{\mathbf{S}}_{\bar{H}} = \operatorname{diag}\{(\mathbf{X}_{\bar{H}}\mathbf{V}_{H})(\mathbf{X}_{\bar{H}}\mathbf{V}_{H})^{T}\}$$
(5)

Predicted MD:

$$\widehat{\mathbf{M}}_{H} = \operatorname{diag}\{(\widehat{\mathbf{S}}_{H} \boldsymbol{\Delta}_{\bar{H}})(\widehat{\mathbf{S}}_{H} \boldsymbol{\Delta}_{\bar{H}})^{T}\}$$

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• How much do observations change between X and robust X'?

Orthogonal distance

Given two commensurate matrices, \mathbf{X} and \mathbf{X}' , orthogonal distances (OD) are defined as:

$$\mathbf{O} = \| \mathbf{X} - \mathbf{X}' \| \tag{7}$$