

# Almost everything you need to know about PLS

## Part 1: Background, Theory, and Examples

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# PLSC Flavors

# PLSC has many...

## Flavors

- Standard
- Mean-Centered
- Seed
- And many more (not covered today)

## Standard PLSC

# A refresher

## Standard PLSC

- computed as the SVD of the cross-product matrix of  $\mathbf{X}$  and  $\mathbf{Y}$
- $\mathbf{X}$  is a matrix of observations by variables
- $\mathbf{Y}$  is a matrix of observations by (some other) variables

as a reminder

- Today we will cover the most common variations in PLSC
- Terminology differs; we stick to PCA nomenclature
- Stay tuned for more in depth examples in R and Matlab

## An example

# PLSC dataset

ADNI ( $N = 569$ )

- 3 groups of participants
  - $N = 178$  healthy control
  - $N = 275$  late MCI
  - $N = 116$  AD
- 8 neuropsych measures
- 68 cortical thickness estimates (via Freesurfer)

## Data matrices

	BNT	Clock	...RAVLT		R.IFG	L.IFG	... L.Fusi		
Subj <sub>1</sub>	10	5	...	30	Subj <sub>1</sub>	3.24	6.27	...	2.32
Subj <sub>2</sub>	7	4	...	26	Subj <sub>2</sub>	5.89	0.26	...	4.51
Subj <sub>3</sub>	3	0	...	23	Subj <sub>3</sub>	2.84	2.51	...	1.17
⋮	⋮	⋮		⋮	⋮	⋮	⋮	⋮	
Subj <sub>N-1</sub>	2	1	...	18	Subj <sub>N-1</sub>	1.96	8.9	...	3.46
Subj <sub>N</sub>	8	4	...	27	Subj <sub>N</sub>	4.42	7.81	...	1.96

Figure 1: **X** and **Y** matrices in standard PLS

## Standard PLSC scree

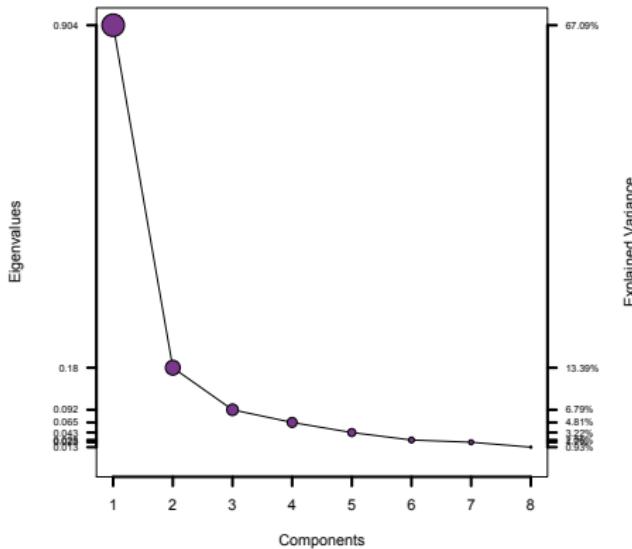


Figure 2

## Beyond descriptive analyses

How many components are “significant”?

- Above expected (or average) contribution
- Permutation
  - scramble the rows of one matrix and recompute PLSC
  - create a null distribution of eigenvalues

For more details on component selection:

- Jackson (1993)
- Peres-Neto et al., (2005)
- Dray (2008)
- Josse and Husson (2011)

## How many components to interpret?

- Again: a mix of art & science
- Use tests, effects sizes, and heuristics

## How many components to interpret?

Component	p.value	eigenvalue	percent.variance
1	0.000	0.904	67.093
2	0.000	0.180	13.388
3	0.009	0.092	6.794
4	0.015	0.065	4.809
5	0.096	0.043	3.223
6	0.453	0.028	2.055
7	0.207	0.023	1.704
8	0.747	0.013	0.934

## How many components to interpret?

- Note: there are no p-values of 0
- Zero means the observed value is outside the distribution
- But the distribution is from number of iterations
- Here: 1000
  - Thus  $p = 0$  is actually  $p < .001$

## How many components to interpret?

Component	p.value	eigenvalue	percent.variance
1	0	0.904	67.093
2	0	0.18	13.388
3	0.009		
4	0.015		
5			
6			
7			
8			

## Two reliable components

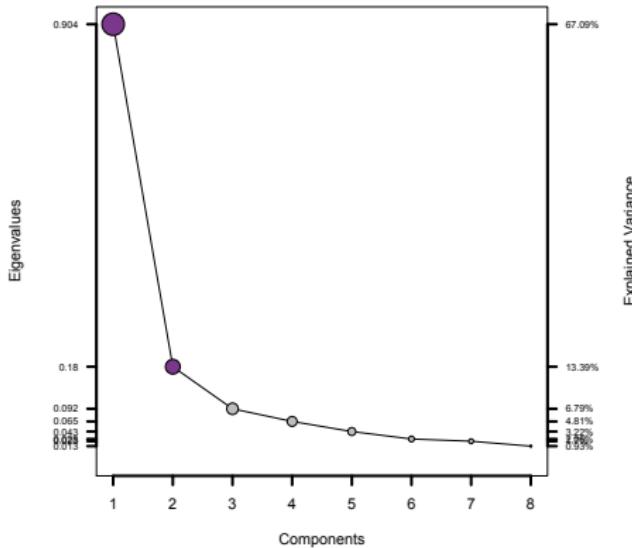


Figure 3

## Latent Variables

# Latent variables

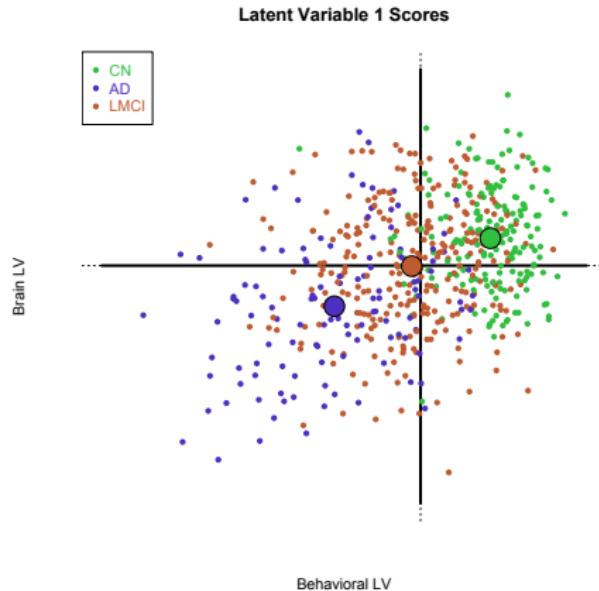


Figure 4

# Latent variables

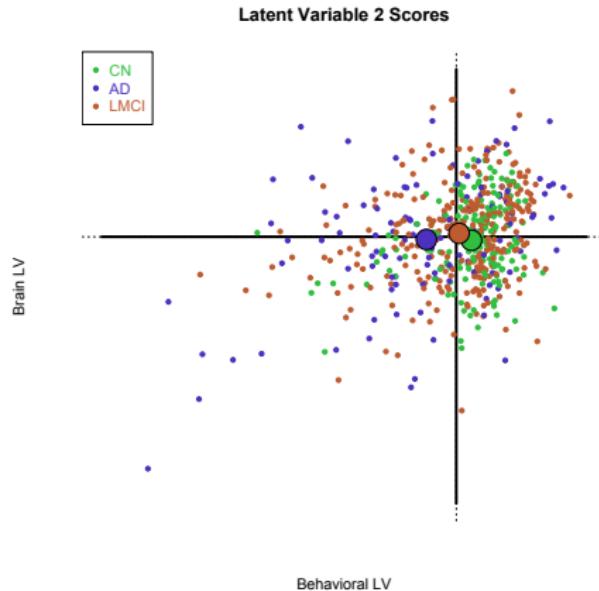


Figure 5

PLSC Flavors  
**Standard PLSC**  
Mean-centered PLSC  
"Seed" PLSC  
And Beyond  
Conclusions

An example  
Latent Variables  
**Component Scores**

## Component Scores

# Neuropsych component scores

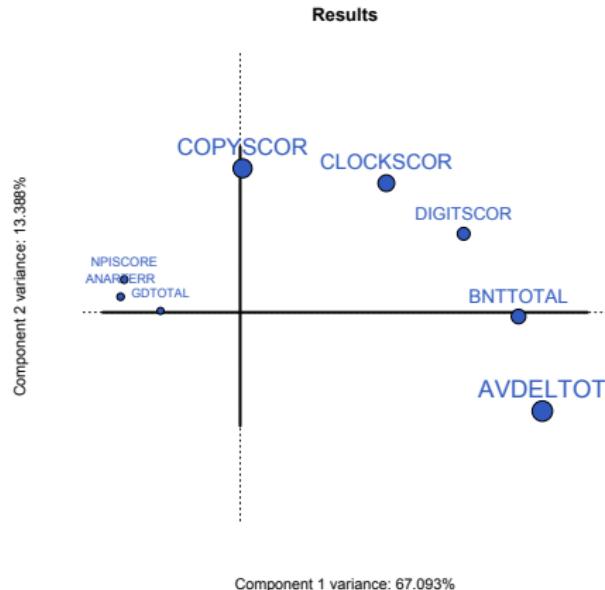


Figure 6

# Structural thickness component scores

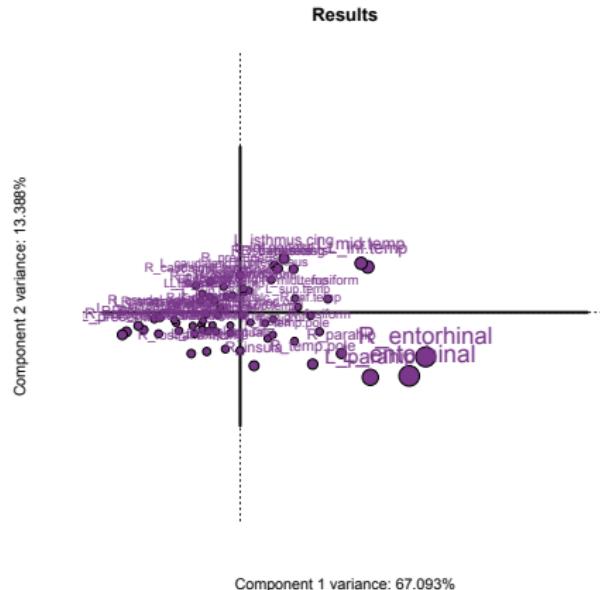


Figure 7

## Beyond descriptive analyses

Which variables significantly contribute to each component?

- Bootstrap resampling (with replacement) builds distributions around our variables
  - Confidence intervals
  - Bootstrap ratios (mean divided by s.d. of distribution)

# Which neuropsych variables significantly contribute?

Table 3: Bootstrap Ratios

	Component.1	Component.2
ANART.ERR	-2.85	0.57
COPY.SCOR	0.12	3.23
CLOCK.SCOR	3.53	3.72
DIGIT.SCOR	5.56	2.44
BNT.TOTAL	6.60	-0.10
AV.DEL.TOT	7.89	-3.50
GD.TOTAL	-1.85	0.02
NPI.SCORE	-2.58	0.80

# Which neuropsych variables significantly contribute?

Table 4: Bootstrap Ratios > +/- 2.5

	Component.1	Component.2
ANART.ERR	-2.85	
COPY.SCOR		3.23
CLOCK.SCOR	3.53	3.72
DIGIT.SCOR	5.56	
BNT.TOTAL	6.6	
AV.DEL.TOT	7.89	-3.5
GD.TOTAL		
NPI.SCORE	-2.58	

# Which neuropsych variables significantly contribute?

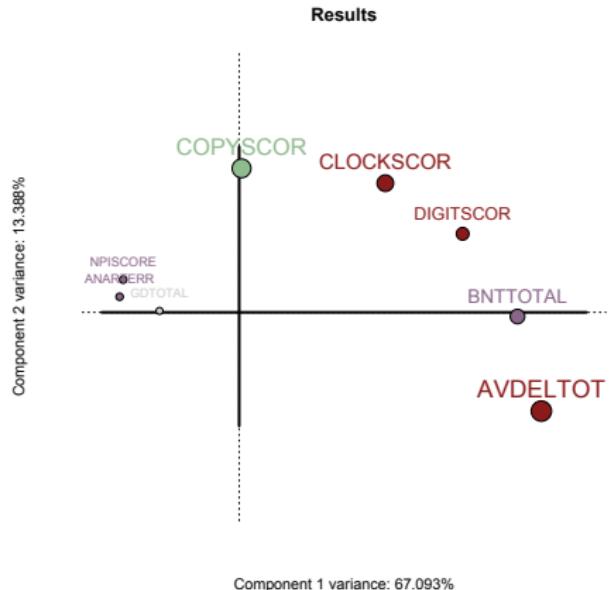


Figure 8

# Which structural regions significantly contribute?

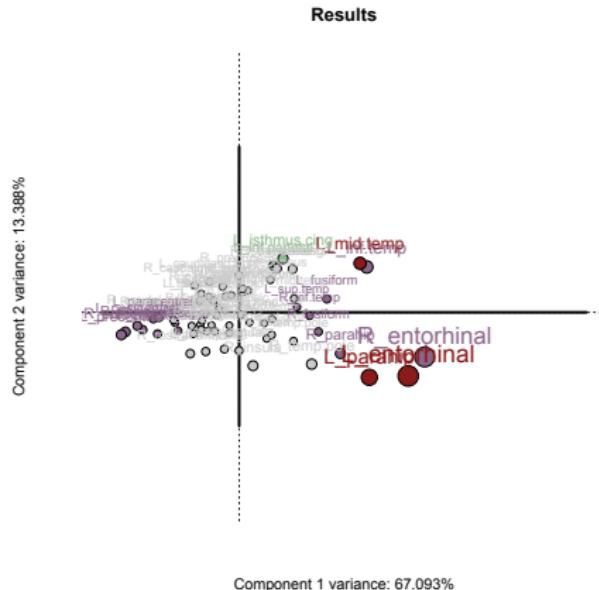


Figure 9

# Putting all the pieces together

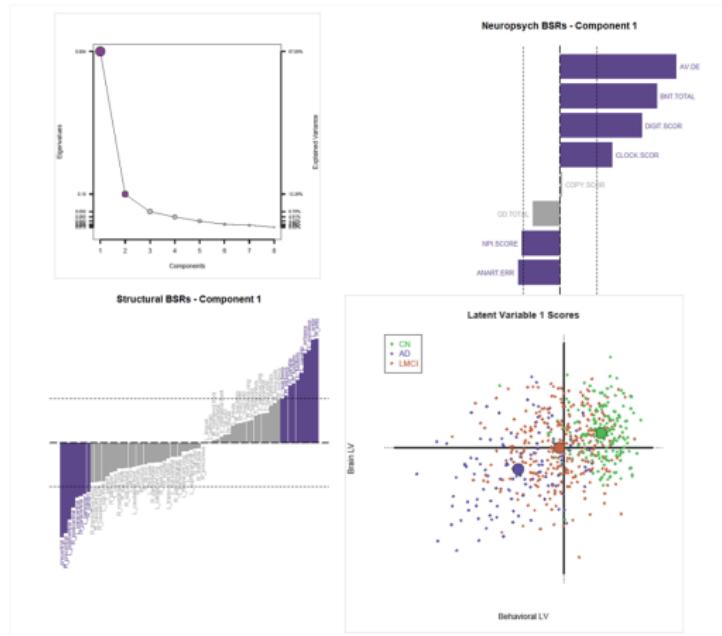


Figure 10

## Mean-centered PLSC

# Mean-centered PLSC

A.k.a.

- Barycentric discriminant analysis
- Between-groups PLSC
- Discriminant PLSC

# Mean-centered PLSC

- used when observations are structured into groups or conditions
- $\mathbf{X}$  is a matrix of observations by variables
- $\mathbf{Y}$  is a dummy matrix that codes for experimental groups or conditions

## An example

## ADNI structural thickness data by diagnostic group

	R.IFG	L.IFG	R.STG	...	L.Fusi		CN	MCI	AD
Subj <sub>1</sub>	3.24	6.27	2.31	...	2.32	Subj <sub>1</sub>	1	0	0
X = Subj <sub>2</sub>	5.89	0.26	3.99	...	4.51	Subj <sub>2</sub>	0	1	0
Subj <sub>3</sub>	2.84	2.51	5.88	...	1.17	Subj <sub>3</sub>	0	1	0
⋮	⋮	⋮	⋮		⋮	⋮	⋮	⋮	⋮
Subj <sub>N-1</sub>	1.96	8.9	7.19	...	3.46	Subj <sub>N-1</sub>	0	0	1
Subj <sub>N</sub>	4.42	7.81	5.99	...	1.96	Subj <sub>N</sub>	1	0	0

Figure 11: **X** and **Y** matrices in mean-centered PLSC

# Group component scores

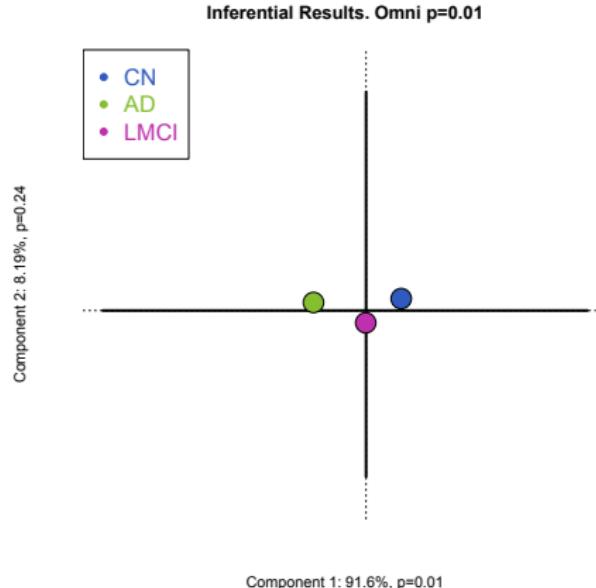


Figure 12

# One set of LV scores

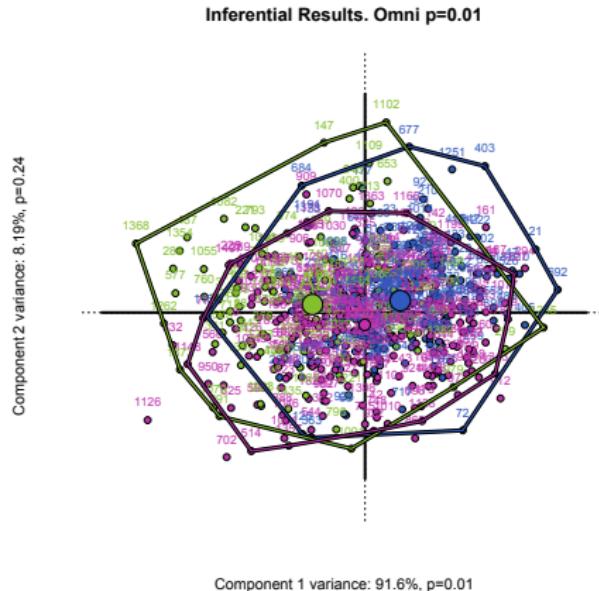


Figure 13

## Fixed effects classification accuracy

Because mc-PLSC is about groups we can assess classification accuracy

## Fixed effects classification accuracy

Table 5: 267 of 569 correctly classified. 46.92 % accuracy

	CN.actual	LMCI.actual	AD.actual
CN.predicted	117	108	25
LMCI.predicted	39	76	17
AD.predicted	22	91	74

## Inference Testing

# Bootstrapping group component scores

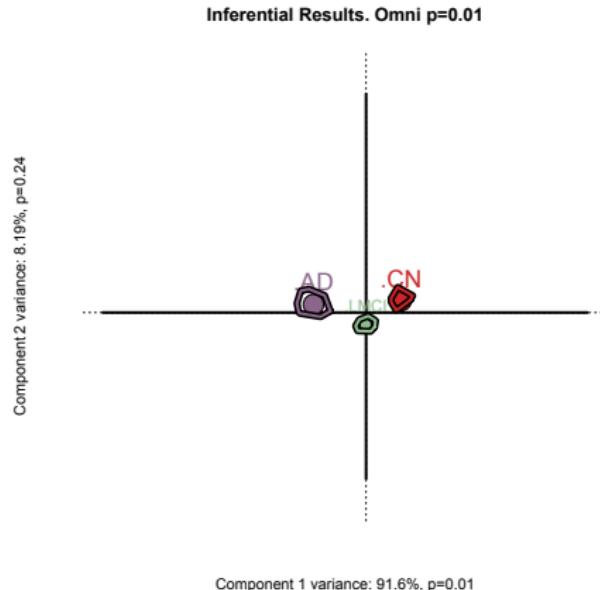


Figure 14

# Bootstrapping structural thickness estimates



Figure 15

## Leave-one-out cross validation

Table 6: 238 of 569 correctly classified. 41.83 % accuracy

	CN.actual	LMCI.actual	AD.actual
CN.predicted	107	111	26
LMCI.predicted	47	65	24
AD.predicted	24	99	66

## “Seed” PLSC

# Seed PLSC

- used to examine “connectivity” between two sets of variables
- name comes from “connectivity”
- broadly it is correlation (or covariance) between
  - $\mathbf{X}_A$  for  $A$  set of variables
  - $\mathbf{X}_B$  for  $B$  set of variables
- Also referred to as “Burt bands”

# Seed PLSC

- **X** is a matrix of observations by variables
- **Y** is a subset of variables (removed from **X**)
  - subset of voxels (i.e., ROI)

## An example

# Structural connectivity in ADNI

$$\mathbf{X} = \begin{matrix} & \text{R.IFG} & \text{L.IFG} & \text{R.STG} & \text{L.STG} & \text{R.MTG} & \dots & \text{L.Fusi} \\ \text{Subj}_1 & 3.24 & 6.27 & 2.31 & 2.67 & 2.59 & \dots & 2.32 \\ \text{Subj}_2 & 5.89 & 0.26 & 3.99 & 2.71 & 1.02 & \dots & 4.51 \\ \text{Subj}_3 & 2.84 & 2.51 & 5.88 & 3.61 & 4.56 & \dots & 1.17 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \dots & \vdots \\ \text{Subj}_{N-1} & 1.96 & 8.9 & 7.19 & 5.36 & 3.01 & \dots & 3.46 \\ \text{Subj}_N & 4.42 & 7.81 & 5.99 & 6.3 & 0.23 & \dots & 1.96 \end{matrix}$$

Figure 16:  $\mathbf{X}$  matrix of Freesurfer thickness estimates

## Structural connectivity in ADNI

	R.IFG	L.IFG	R.STG	L.STG	R.MTG	...	L.Fusi
Subj <sub>1</sub>	3.24	6.27	2.31	2.67	2.59	...	2.32
Subj <sub>2</sub>	5.89	0.26	3.99	2.71	1.02	...	4.51
Subj <sub>3</sub>	2.84	2.51	5.88	3.61	4.56	...	1.17
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮
Subj <sub>N-1</sub>	1.96	8.9	7.19	5.36	3.01	...	3.46
Subj <sub>N</sub>	4.42	7.81	5.99	6.3	0.23	...	1.96

Figure 17: Selecting lateral temporal regions as a seed

# Structural connectivity in ADNI

	R.IFG	L.IFG	...	L.Fusi		R.STG	L.STG	R.MTG	....	TempPole
Subj <sub>1</sub>	3.24	6.27	...	2.32	Subj <sub>1</sub>	2.31	2.67	2.59	...	2.59
Subj <sub>2</sub>	5.89	0.26	...	4.51	Subj <sub>2</sub>	3.99	2.71	1.02	...	1.02
Subj <sub>3</sub>	2.84	2.51	...	1.17	Subj <sub>3</sub>	5.88	3.61	4.56	...	4.56
⋮	⋮	⋮		⋮	⋮	⋮	⋮	⋮	⋮	⋮
Subj <sub>N-1</sub>	1.96	8.9	...	3.46	Subj <sub>N-1</sub>	7.19	5.36	3.01	...	3.01
Subj <sub>N</sub>	4.42	7.81	...	1.96	Subj <sub>N</sub>	5.99	6.3	0.23	...	0.23

Figure 18: **X & Y** matrices in seed PLS

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An example  
**Connectivity Matrices**  
Seed PLSC results  
PLSC wrap-up

## Connectivity Matrices

# Structural connectivity in ADNI

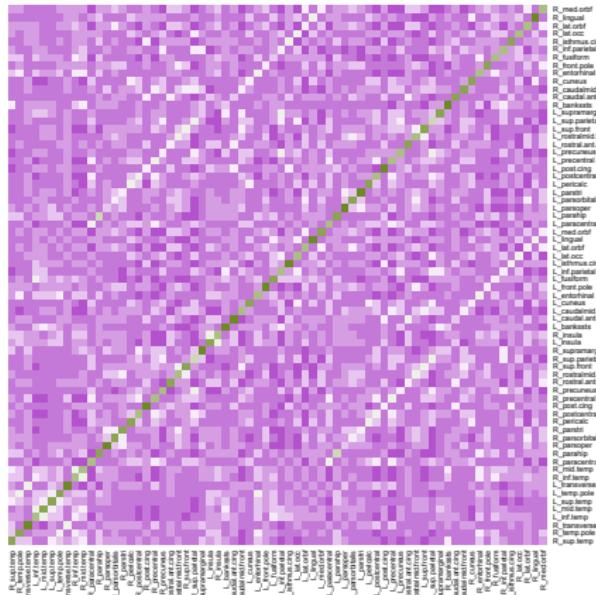


Figure 19

# Structural connectivity in ADNI

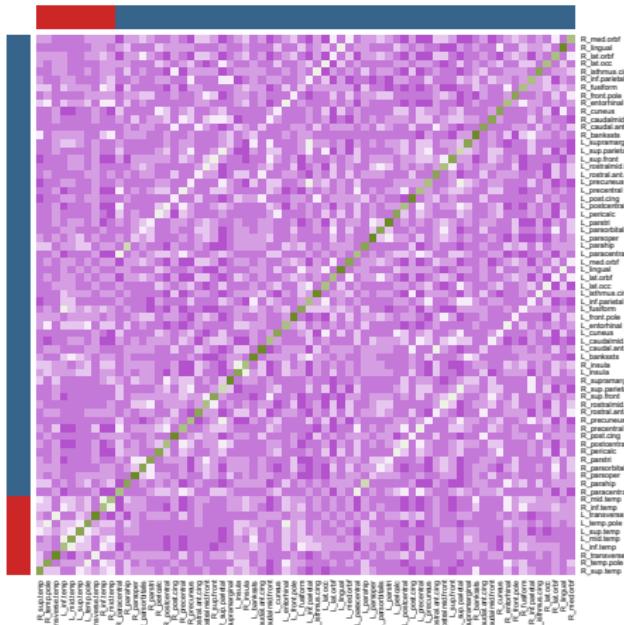


Figure 20

# Structural connectivity in ADNI

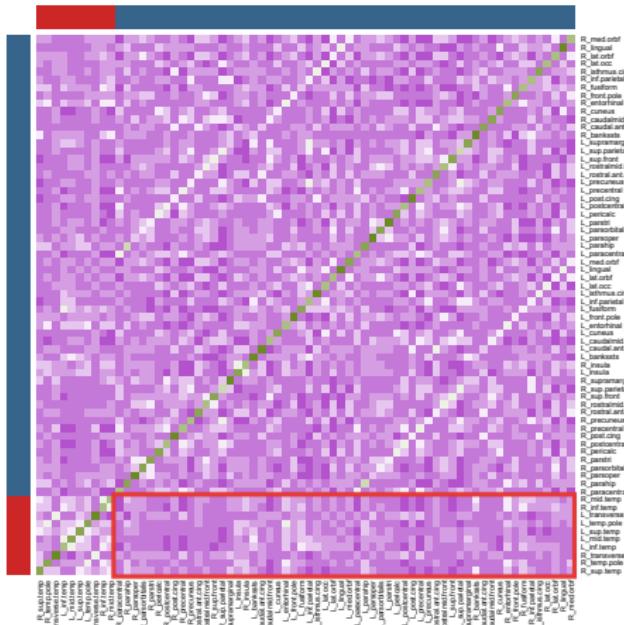


Figure 21

## Seed PLSC results

# Seed PLSC scree

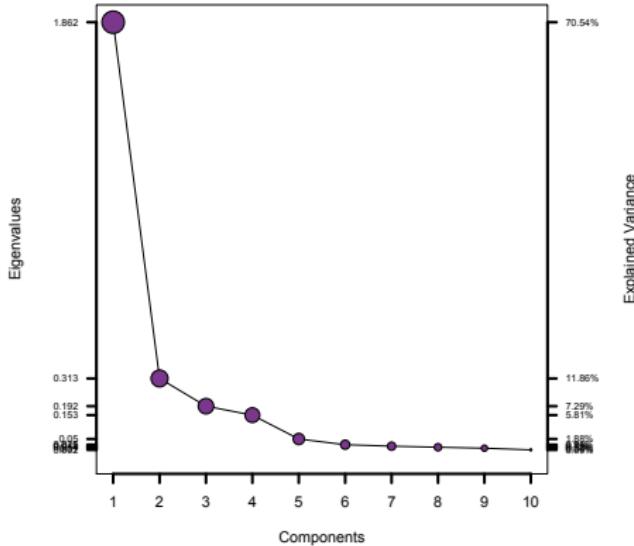


Figure 22

# Structural connectivity latent variable

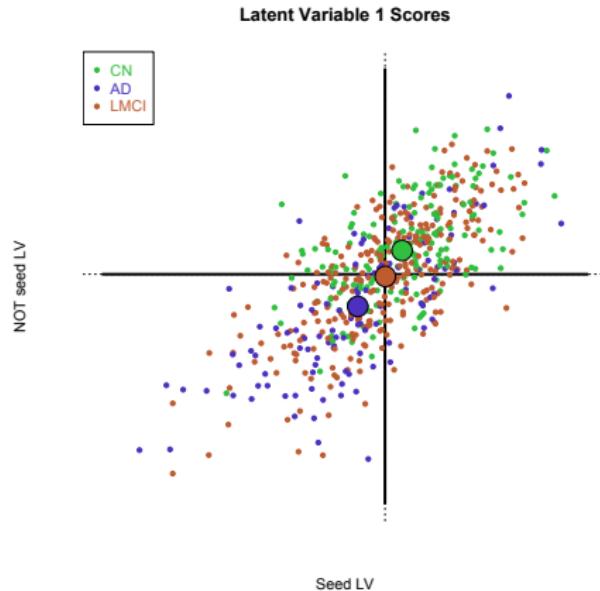
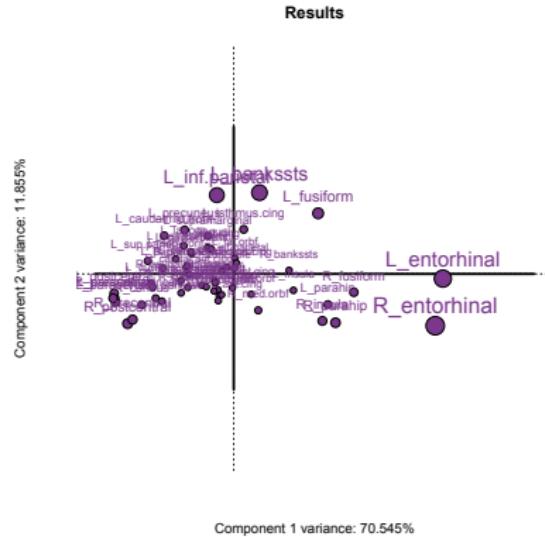
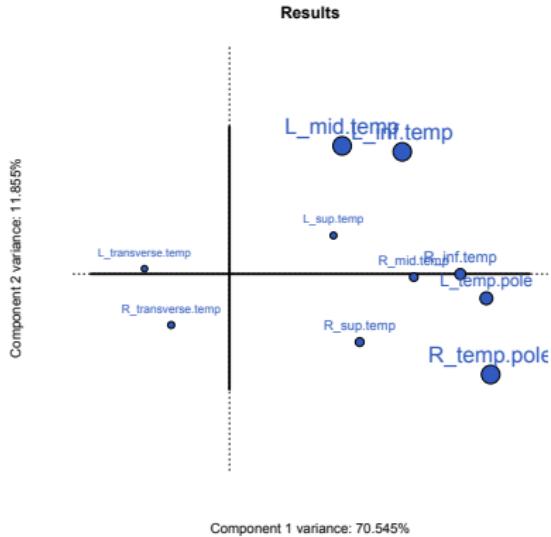


Figure 23

# Seed and Not Seed component scores



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## PLSC wrap-up

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# PLSC

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Friends of PLS  
More inference & issues

## And Beyond

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## Friends of PLS

## RRR & CCA

### Reduced Rank Regression (RRR) and Canonical Correlation Analysis (CCA)

- PLS is about covariance:  $\mathbf{X}^T \mathbf{Y}$
- RRR is like OLS:
  - $\mathbf{R}_{RRR} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y} = \mathbf{U} \Delta \mathbf{V}^T$
- CCA is about correlation:
  - $\mathbf{R}_{CCA} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y} (\mathbf{Y}^T \mathbf{Y})^{-1} = \mathbf{U} \Delta \mathbf{V}^T$
- These look very similar
  - But they really are not
  - See McIntosh & Misic (2013)

# PLS-Correspondence analysis

- What if you have non-quantitative data?
  - For both **X** and **Y**
  - For just **X** or **Y**
  - Mixed within **X** or **Y**
- PLS-CA handles mixed data in a PLSC framework
  - Generalizes PLSC
  - See Beaton et al., (2016)

## More inference & issues

## Resampling & Inference

- See <https://github.com/derekbeaton/Workshops/tree/master/RTC/Apr2017> for prequel to today's workshop
- We've covered
  - Bootstrap
  - Permutation
  - Leave-one-out (a bit)
- We did not cover
  - Variations of the above
  - Split-half
- There are some known issues to be aware of
  - Kovacevic et al., (2013)
  - Churchill et al., (2013)

## Conclusions

## Major points

- If you know the SVD you know
  - "Almost everything you need to know about [blank]"
- PLS is a large family of techniques
  - In our fields we focus on PLSC
- PLSC has many flavors
  - We've covered a few
  - We'll cover a few more in November
- Today's material will be at:
  - <https://github.com/derekbeaton/Workshops/tree/master/RTC/Oct2017>

## Before next time

- We will send out reading material
- Almost everything you need:
  - McIntosh & Lobaugh (2004)
  - Krishan et al., (2011)

Fin

- Questions?
- Comments?
- Complaints?