

Principal Components & Multiple Correspondence Analyses

with resampling approaches for stability assessments

Derek Beaton

RRI RTC

May 03, 2019

Where to find everything

- ▶ Generally: <https://github.com/derekbeaton/workshops>

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- ▶ Today: https://github.com/derekbeaton/Workshops/tree/master/RTC/PCA_MCA_Resampling

Some set up

- ▶ Use RStudio (makes it easy)

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 - ▶ Run scripts in order
- ▶ Use of the ADNI data
 - ▶ Via the ‘ADNIMERGE’ package

An advertisement

- ▶ Lots of really cool R & RStudio stuff

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 - ▶ We'll make stuff available

Motivation for today

- ▶ Not everything is a number

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- ▶ Not everything is a number
- ▶ But with care, it can be turned into one

Overview

- ▶ Introduction

Overview

- ▶ Introduction
- ▶ PCA

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- ▶ Introduction
- ▶ PCA
- ▶ CA

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- ▶ Introduction
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Overview

- ▶ Introduction
- ▶ PCA
- ▶ CA
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- ▶ Final notes

Introduction

Prehistory

- ▶ Basis:

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 - ▶ Spearman (1904)

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- ▶ See Lebart's History & Prehistory of CA: http://www.dtmvic.com/doc/About_the_History_of_CA.pdf

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 - ▶ CA makes you an expert with all of them

PCA & CA

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- ▶ So what's the difference?

PCA vs CA

- ▶ PCA: For generally continuous (interval scale) data

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PCA vs CA

- ▶ PCA: For generally continuous (interval scale) data
- ▶ CA: For (almost) everything else
 - ▶ And also for continuous data!

Under the hood

- ▶ The eigenvalue decomposition (EVD)

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 - ▶ Required for CA and fancier PCA-like techniques & extensions

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- ▶ Component scores

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 - ▶ Variance = Sums of squares
- ▶ Magic

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- ▶ Making things look fancy:

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 - ▶ New package for outliers
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- ▶ Making things look fancy:
 - ▶ kable, kableExtra, gridExtra, ggcorrplot

Some alternatives

- ▶ FactoMineR

Some alternatives

- ▶ FactoMineR
- ▶ ade4

Some alternatives

- ▶ FactoMineR
- ▶ ade4
- ▶ ca

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- ▶ FactoMineR
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- ▶ ca
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Some alternatives

- ▶ FactoMineR
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Some alternatives

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- ▶ ade4
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- ▶ So many others

Typology

- ▶ SS Stevens

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 - ▶ Not a boat!

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- ▶ Levels of measurement

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 - ▶ Nominal (categorical)

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 - ▶ Ordinal (ranked, discrete categories)
 - ▶ Interval (continuous, arbitrary 0)

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 - ▶ Nominal (categorical)
 - ▶ Ordinal (ranked, discrete categories)
 - ▶ Interval (continuous, arbitrary 0)
 - ▶ Ratio (continuous, non-arbitrary 0)
- ▶ Excellent examples:
https://en.wikipedia.org/wiki/Level_of_measurement

Today's data

- ▶ Alzheimer's Disease Neuroimaging Initiative (ADNI)

Today's data

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- ▶ Data set:

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 - ▶ 17 variables

Today's data

- ▶ Alzheimer's Disease Neuroimaging Initiative (ADNI)
- ▶ Data set:
 - ▶ 665 observations
 - ▶ 17 variables
- ▶ Walk through this set to tell a whole story

Today's data

	Continuous	Categorical	Ordinal
<i>DX</i>		YES	
<i>AGE</i>	YES		
<i>PTGENDER</i>		YES	
<i>PTEDUCAT</i>			YES
<i>PTETHCAT</i>		YES	
<i>PTRACCAT</i>		YES	
<i>APOE4</i>		YES	YES
<i>FDG</i>	YES		
<i>AV45</i>	YES		
<i>CDRSB</i>			YES
<i>ADAS13</i>			YES
<i>MOCA</i>			YES
<i>WholeBrain</i>	YES		
<i>Hippocampus</i>	YES		
<i>MidTemp</i>	YES		
<i>mPACctrailsB</i>	YES		
<i>HMSCORE</i>		YES	YES

Principal Components Analysis

Let's dive in

- ▶ We'll start with just two variables:

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- ▶ Trails
 - ▶ Neuropsych test
 - ▶ Executive function

Let's dive in

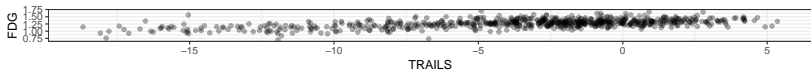
- ▶ We'll start with just two variables:
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- ▶ FDG

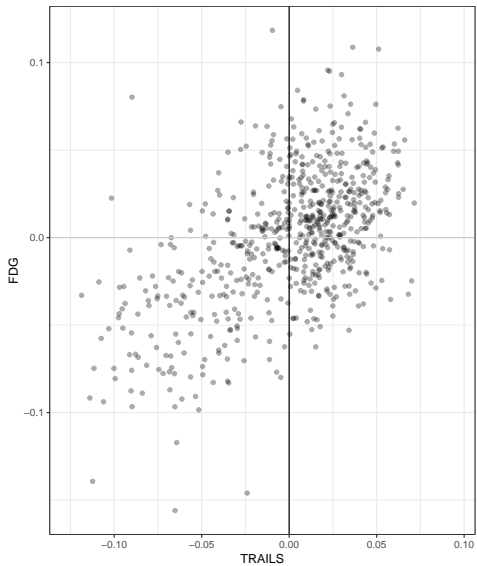
Let's dive in

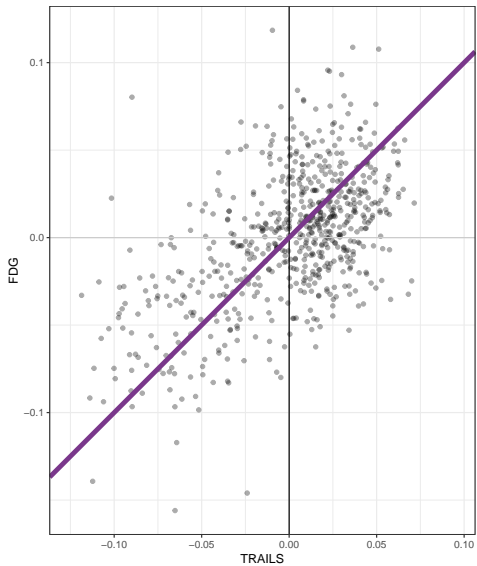
- ▶ We'll start with just two variables:
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 - ▶ Neuropsych test
 - ▶ Executive function
- ▶ FDG
 - ▶ PET imaging; brain function

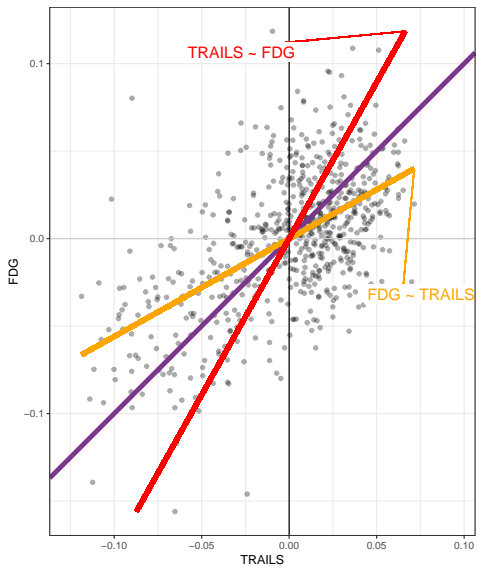
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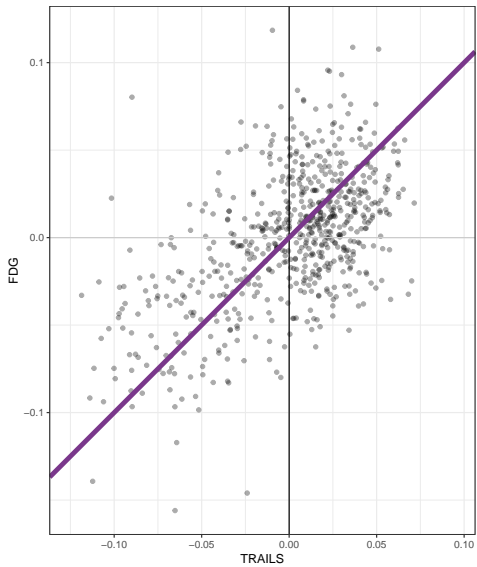
- ▶ We'll start with just two variables:
- ▶ Trails
 - ▶ Neuropsych test
 - ▶ Executive function
- ▶ FDG
 - ▶ PET imaging; brain function
 - ▶ Average of several brain regions

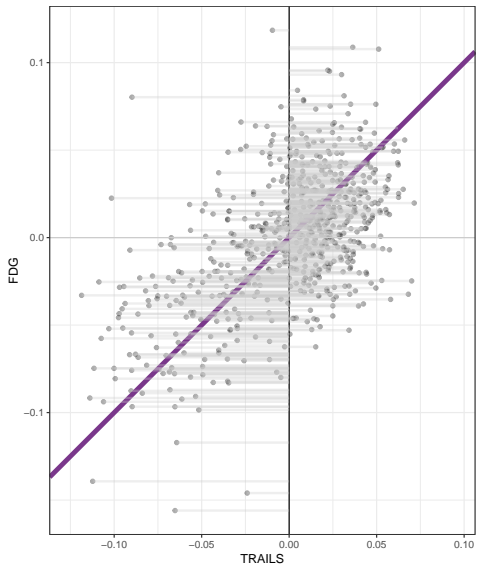


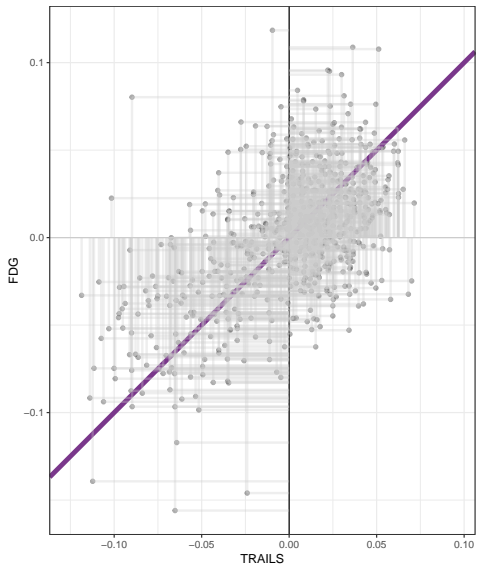


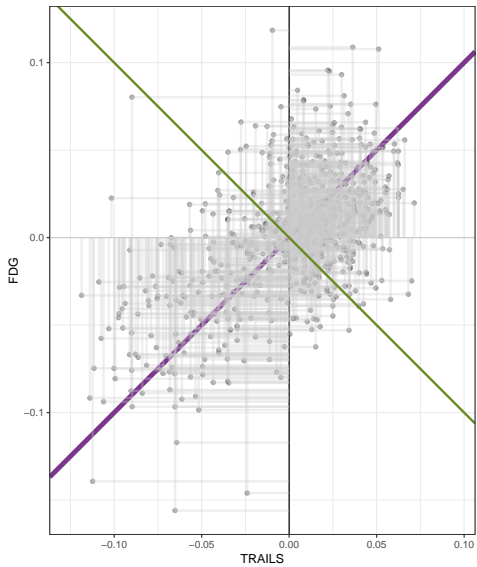




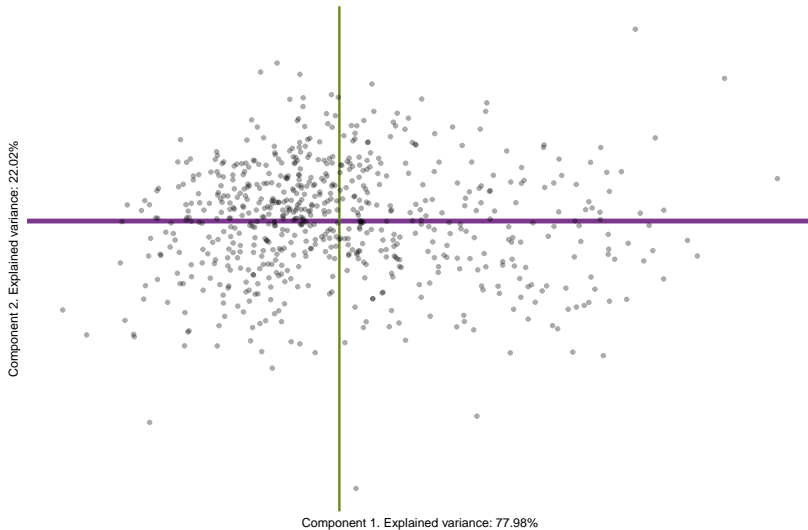




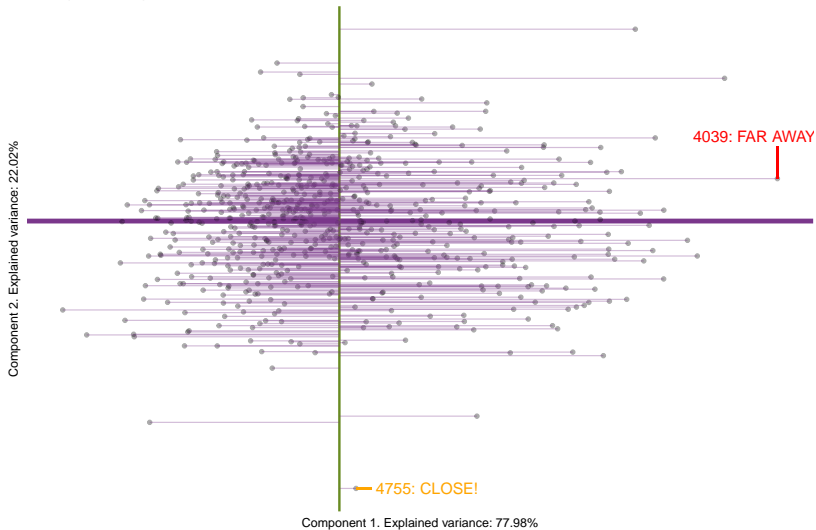




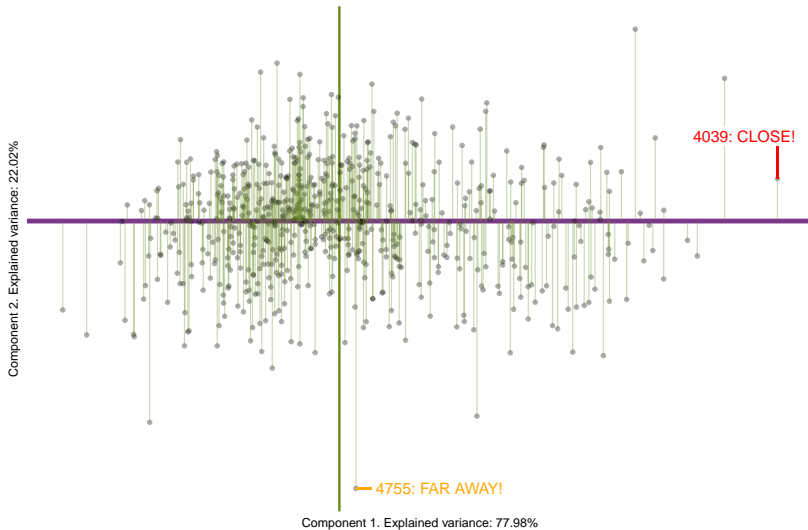
PCA of Trails & FDG:
Participants' Component Scores



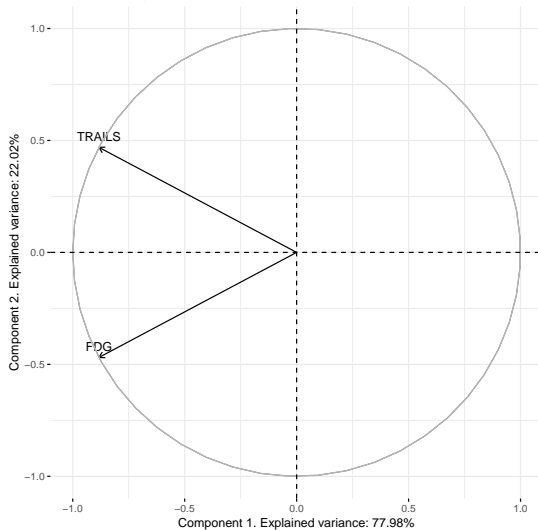
PCA of Trails & FDG:
Participants' Component Scores



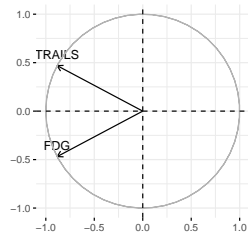
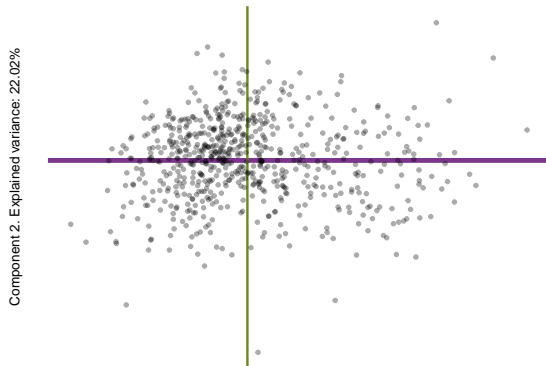
PCA of Trails & FDG:
Participants' Component Scores



PCA:
Variable-Component Correlations

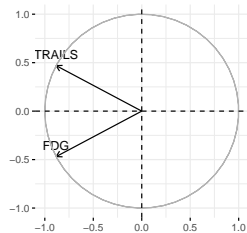
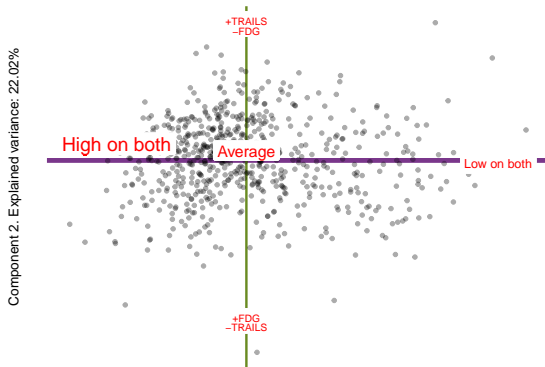


PCA:
Trails & FDG



Component 1. Explained variance: 77.98%

PCA:
Trails & FDG



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Scaling up

- ▶ Scale up: MORE DATA!

Scaling up

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- ▶ All of the continuous variables

Scaling up

	AGE	FDG	AV45	WholeBrain	Hippocampus	MidTemp	mPACtrailsB
5023	63.9	1.29	1.03	1057350.97	7904	21306	1.81
5026	70.5	1.08	1.44	1023057.28	8051	16501	-1.45
5027	75.5	1.06	1.44	986723.65	6534	17437	-17.27
5028	61.9	1.13	1.38	1182704.57	7481	20797	-11.5
5031	80.2	1.14	1.52	908133.86	5040	19032	-8.21
5037	67.3	0.98	1.21	1161499.61	5831	21428	-12.8
5040	75.9	1.24	1.01	943160.57	7994	16634	0.94
5047	68.8	1.7	1.48	1070406.07	7920	22043	-4.9
5054	74	1.12	1.43	1138040.06	6580	20836	-7.63
5058	61.8	0.97	1.54	1195549.29	7318	22757	-9.18
5063	71.5	0.92	1.61	817421.23	5364	12542	-15.03

A new plot

- ▶ Scree (Cattell)

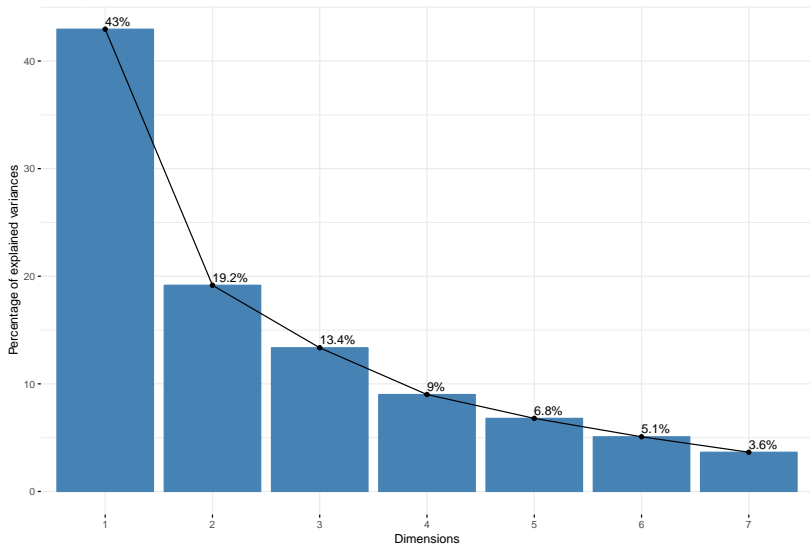
A new plot

- ▶ Scree (Cattell)
- ▶ Junk at the bottom of a slope

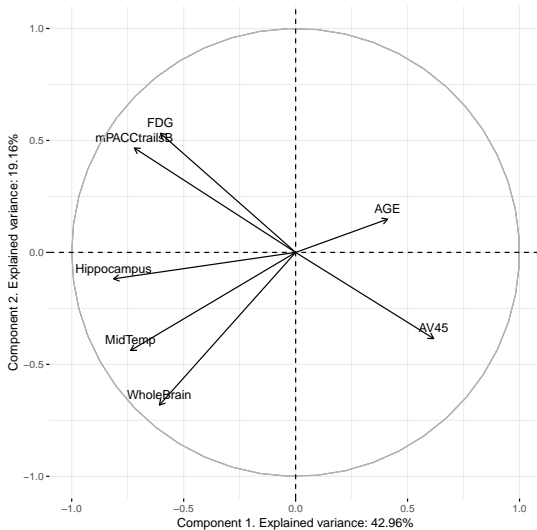
A new plot

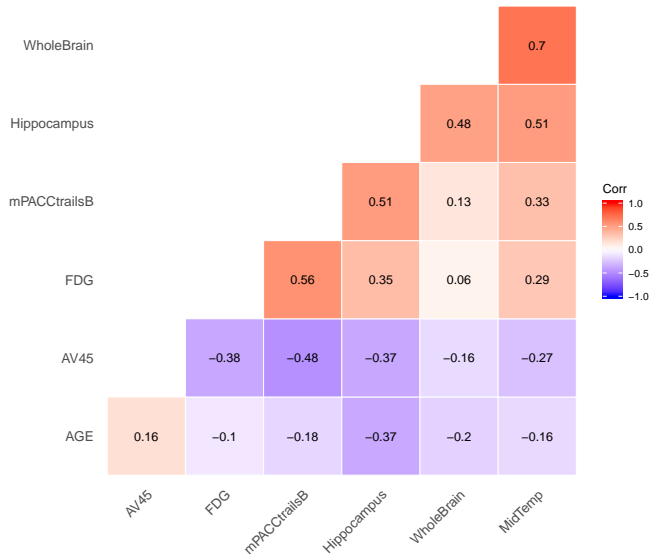
- ▶ Scree (Cattell)
- ▶ Junk at the bottom of a slope
- ▶ Shows us explained variance (%) per component

Scree plot

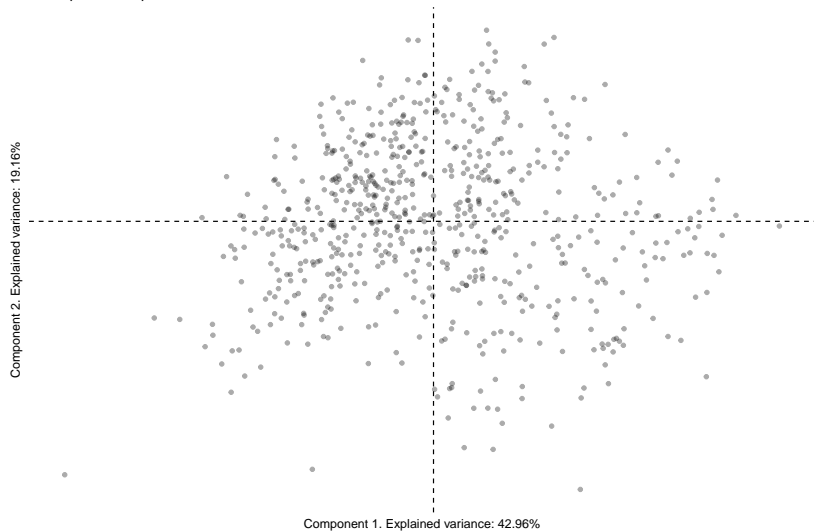


PCA:
Variable-Component Correlations



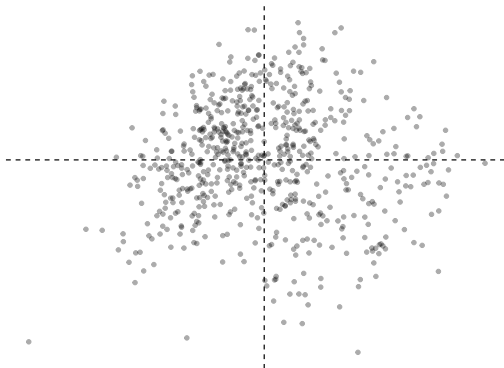


PCA:
Participants Component Scores

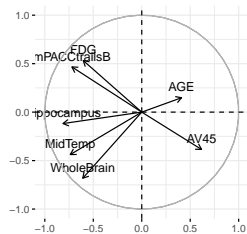


PCA:
Trails & FDG

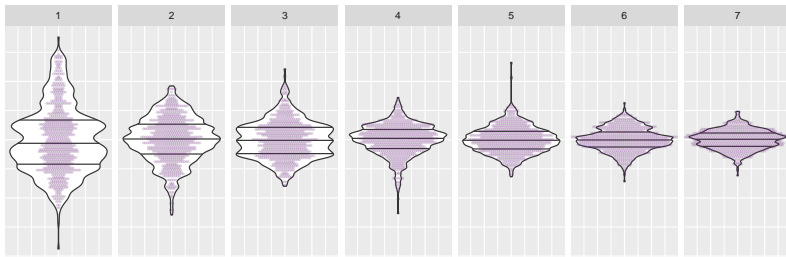
Component 2. Explained variance: 19.16%



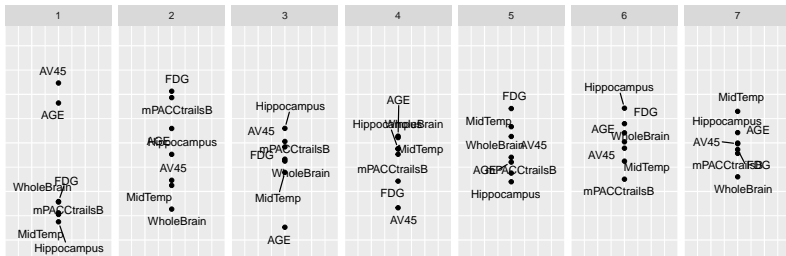
Component 1. Explained variance: 42.96%



COMPONENT_SCORES



CORRELATIONS



Correspondence analyses

CA

- ▶ Like PCA in many ways

CA

- ▶ Like PCA in many ways
- ▶ Slightly different interpretations

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- ▶ So much cooler

CA

- ▶ Like PCA in many ways
- ▶ Slightly different interpretations
- ▶ So much cooler
 - ▶ Handles all types of data

Illustrative data

	DX	PTRACCAT
5023	CN	Asian
5026	MCI	White
5027	Dementia	White
5028	Dementia	White
5031	MCI	White
5037	Dementia	Black
5040	CN	Black
5047	MCI	Black
5054	Dementia	White
5058	Dementia	Asian
5063	Dementia	White

Disjunctive data

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0

Disjunctive data

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0

- ▶ Row sums are total number of *original* variables

Disjunctive data

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g. DX) is total number of rows

Disjunctive data

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g. DX) is total number of rows
- ▶ Sum of the table is rows \times columns

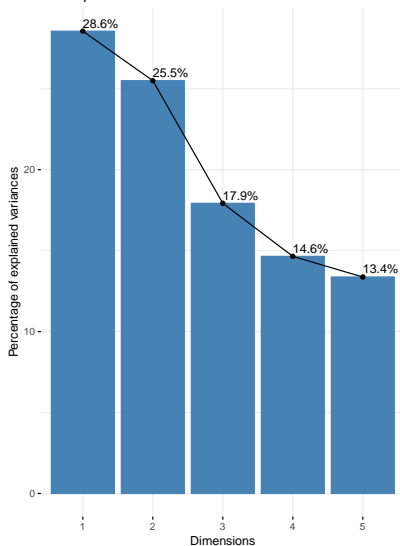
A bad idea: PCA

- ▶ “coding categorical variables with the indicator matrix of dummy variables and considering them as Gaussian, for instance, is almost a crime.”

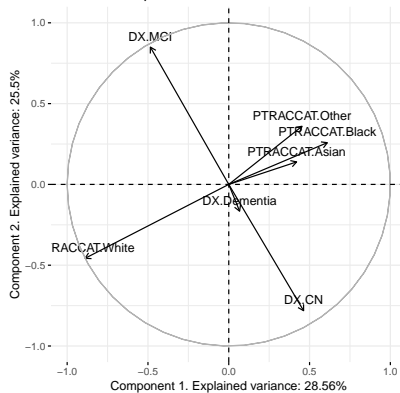
A bad idea: PCA

- ▶ “coding categorical variables with the indicator matrix of dummy variables and considering them as Gaussian, for instance, is almost a crime.”
 - ▶ “Jan de Leeuw and the French School of Data Analysis”
(Husson, Josse, Saporta)

Scree plot



PCA:
Variable-Component Correlations



Why is that a bad idea?

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
<i>DX.MCI</i>	1	-0.815	-0.363	0.045	0.032	-0.043	-0.072
<i>DX.CN</i>	-0.815	1	-0.243	-0.047	0	0.067	0.003
<i>DX.Dementia</i>	-0.363	-0.243	1	0	-0.053	-0.035	0.116
<i>PTRACCAT.White</i>	0.045	-0.047	0	1	-0.562	-0.657	-0.45
<i>PTRACCAT.Other</i>	0.032	0	-0.053	-0.562	1	-0.031	-0.021
<i>PTRACCAT.Black</i>	-0.043	0.067	-0.035	-0.657	-0.031	1	-0.025
<i>PTRACCAT.Asian</i>	-0.072	0.003	0.116	-0.45	-0.021	-0.025	1

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
<i>DX.MCI</i>	365	0	0	341	11	10	3
<i>DX.CN</i>	0	235	0	213	6	12	4
<i>DX.Dementia</i>	0	0	65	60	0	1	4
<i>PTRACCAT.White</i>	341	213	60	614	0	0	0
<i>PTRACCAT.Other</i>	11	6	0	0	17	0	0
<i>PTRACCAT.Black</i>	10	12	1	0	0	23	0
<i>PTRACCAT.Asian</i>	3	4	4	0	0	0	11

A better idea

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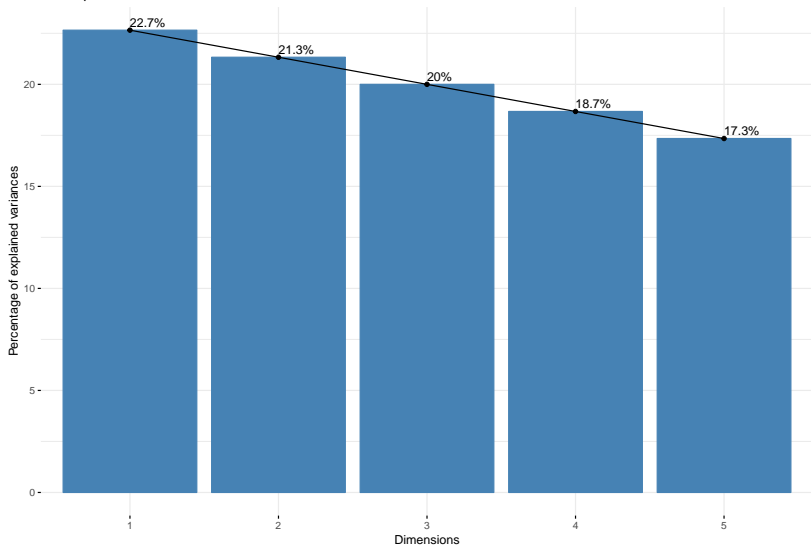
A better idea

- ▶ Correspondence analysis (CA)
 - ▶ Think of it as a χ^2 PCA
- ▶ Designed to handle things that look like counts
 - ▶ That includes categories
 - ▶ And some other things
- ▶ Row and column component scores exist on same scale

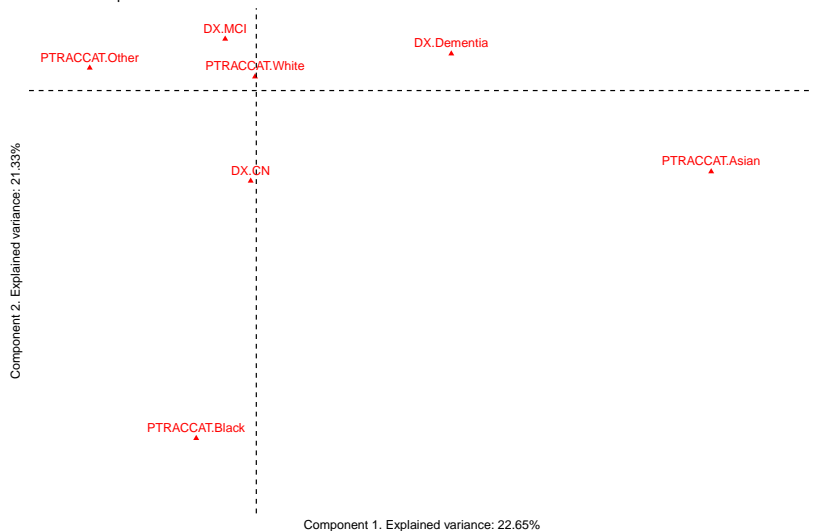
A better idea

- ▶ Correspondence analysis (CA)
 - ▶ Think of it as a χ^2 PCA
- ▶ Designed to handle things that look like counts
 - ▶ That includes categories
 - ▶ And some other things
- ▶ Row and column component scores exist on same scale
 - ▶ CA is a *bivariate* technique

Scree plot

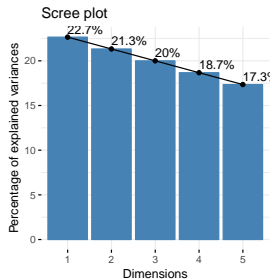


CA:
Variable Component Scores



	DX	PTRACCAT
5023	CN	Asian
5026	MCI	White
5027	Dementia	White
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5047	MCI	Black
5054	Dementia	White
5058	Dementia	Asian
5063	Dementia	White

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0



Multiple correspondence analysis

- ▶ An extension of CA

Multiple correspondence analysis

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- ▶ Accommodates multiple categorical variables (CA only does 2)

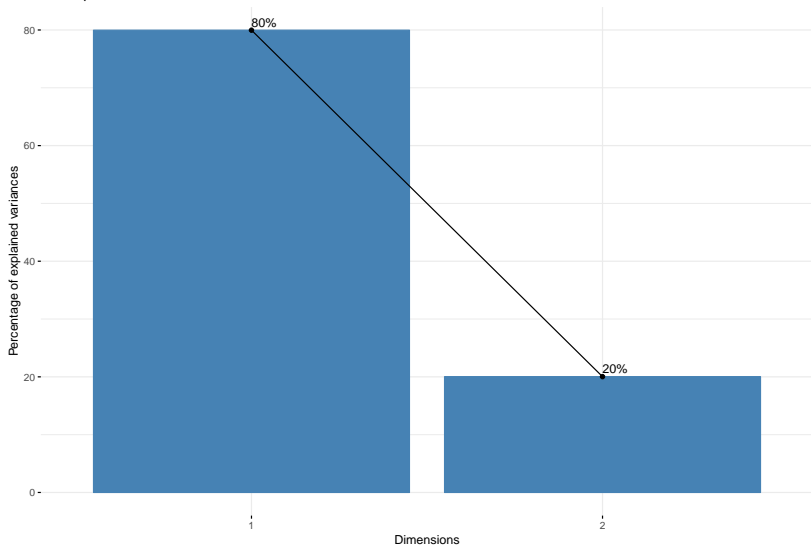
Multiple correspondence analysis

- ▶ An extension of CA
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- ▶ Corrects the dimensionality

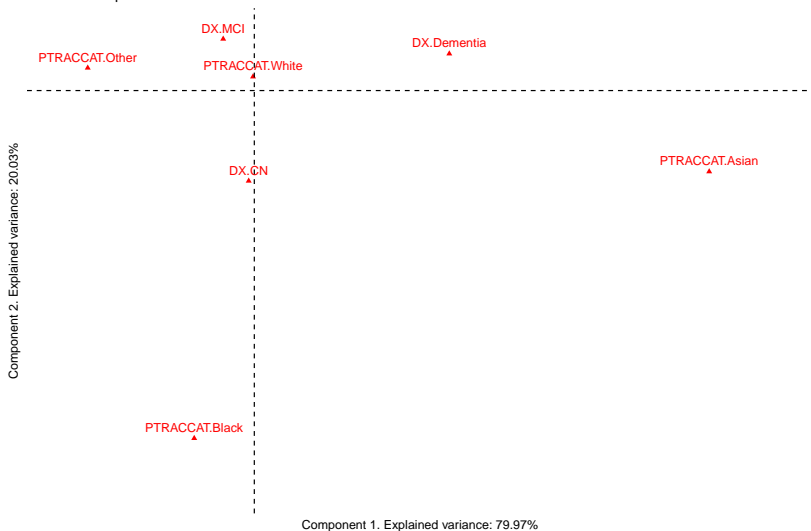
Multiple correspondence analysis

- ▶ An extension of CA
- ▶ Accommodates multiple categorical variables (CA only does 2)
- ▶ Corrects the dimensionality
- ▶ Has nearly magical properties (we'll see later)

Scree plot

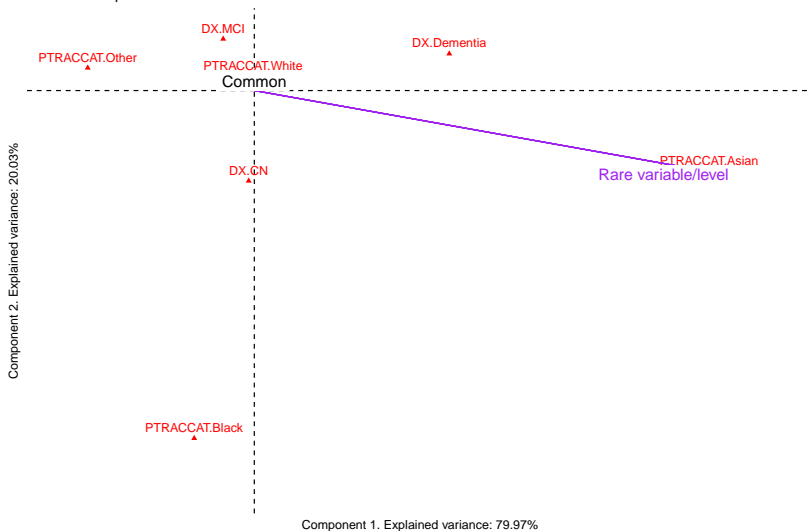


MCA:
Variable Component Scores

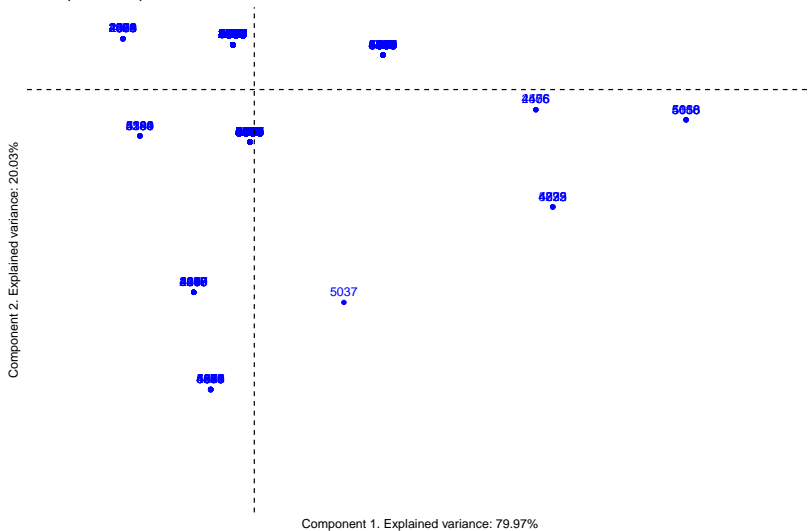


New interpretations

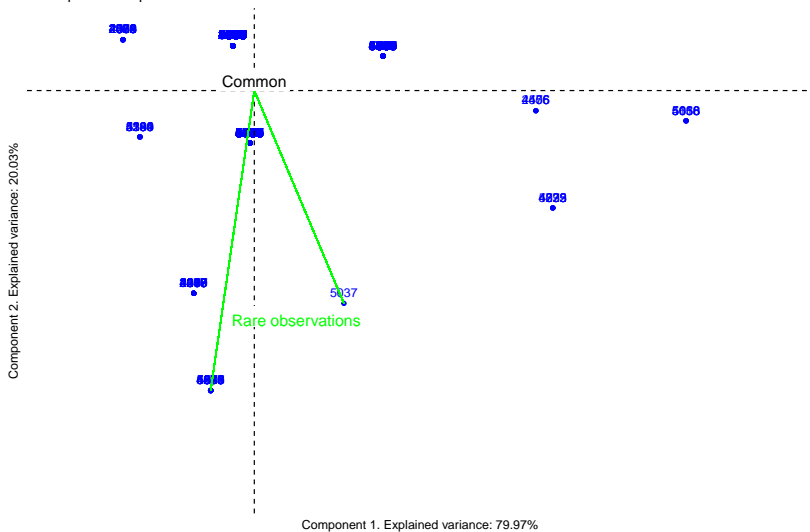
MCA:
Variable Component Scores



MCA:
Participants Component Scores



MCA:
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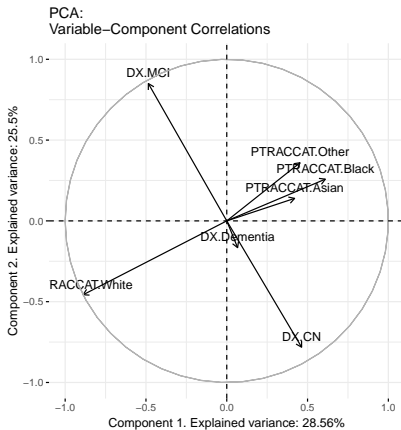
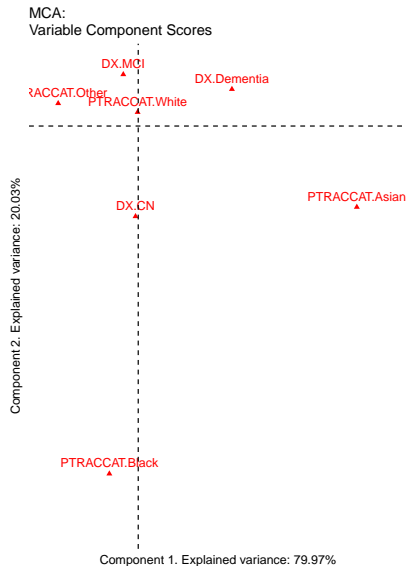


Why does it look like that?

DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
1	0	0	1	0	0	0
1	0	0	0	1	0	0
1	0	0	0	0	1	0
0	1	0	0	1	0	0
1	0	0	0	0	0	1
0	1	0	1	0	0	0
0	1	0	0	0	1	0
0	0	1	1	0	0	0
0	1	0	0	0	0	1
0	0	1	0	0	0	1
0	0	1	0	0	1	0

These are *all* the possible combinations from all 665

Compare the results



Compare the results

	PCA Comp. 1	PCA Comp. 2	PCA Comp. 3	PCA Comp. 4	PCA Comp. 5
MCA Comp. 1	0.17	-0.25	0.92	0.06	-0.26
MCA Comp. 2	-0.78	0.36	0.28	-0.42	0.03

- ▶ CA & MCA produce identical results, except MCA:

Compare the results

	PCA Comp. 1	PCA Comp. 2	PCA Comp. 3	PCA Comp. 4	PCA Comp. 5
MCA Comp. 1	0.17	-0.25	0.92	0.06	-0.26
MCA Comp. 2	-0.78	0.36	0.28	-0.42	0.03

- ▶ CA & MCA produce identical results, except MCA:
 - ▶ Drops components

Compare the results

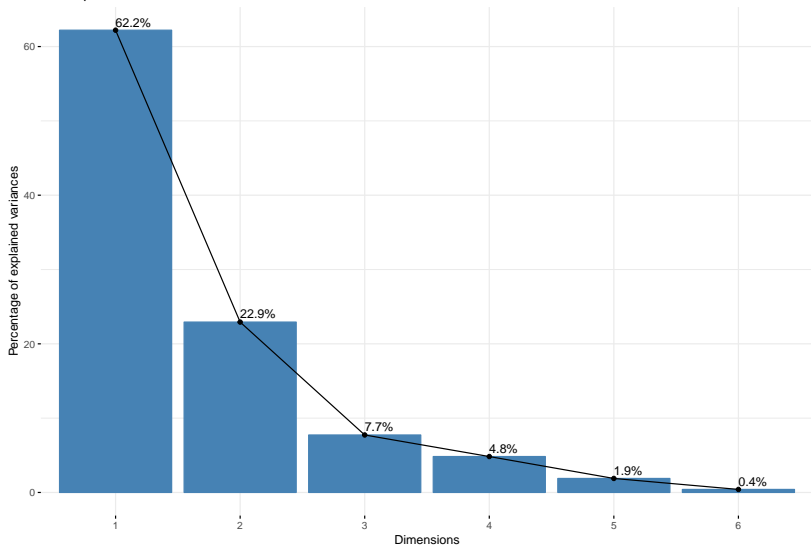
	PCA Comp. 1	PCA Comp. 2	PCA Comp. 3	PCA Comp. 4	PCA Comp. 5
MCA Comp. 1	0.17	-0.25	0.92	0.06	-0.26
MCA Comp. 2	-0.78	0.36	0.28	-0.42	0.03

- ▶ CA & MCA produce identical results, except MCA:
 - ▶ Drops components
 - ▶ Corrects explained variance

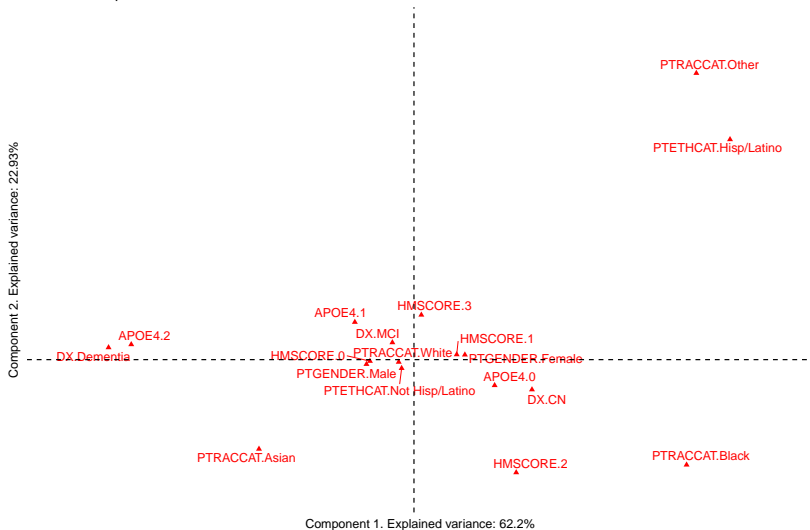
Scaling up

	DX	PTGENDER	PTETHCAT	PTRACCAT	APOE4	HMSCORE
5023	CN	Female	Not Hisp/Latino	Asian	0	0
5026	MCI	Female	Not Hisp/Latino	White	1	1
5027	Dementia	Male	Not Hisp/Latino	White	0	1
5028	Dementia	Male	Not Hisp/Latino	White	2	1
5031	MCI	Female	Hisp/Latino	White	0	1
5037	Dementia	Male	Not Hisp/Latino	Black	1	1
5040	CN	Female	Not Hisp/Latino	Black	0	1
5047	MCI	Female	Not Hisp/Latino	Black	2	1
5054	Dementia	Female	Not Hisp/Latino	White	1	0
5058	Dementia	Male	Not Hisp/Latino	Asian	0	0
5063	Dementia	Female	Not Hisp/Latino	White	1	1

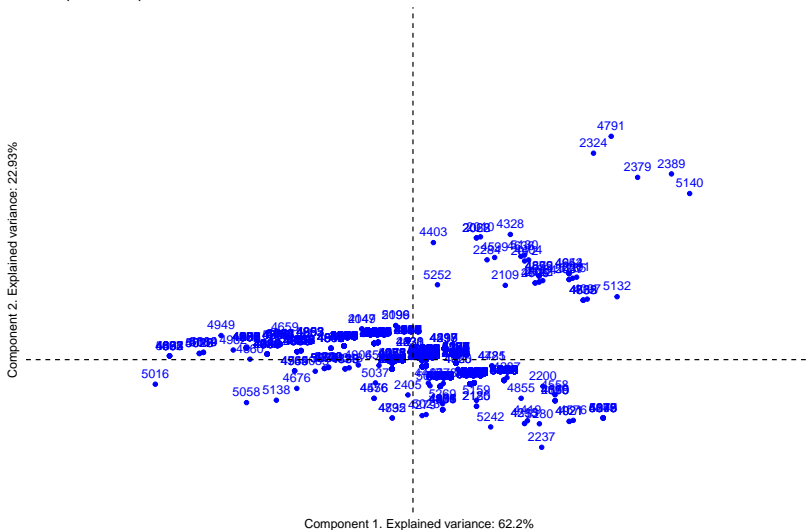
Scree plot



MCA:
Variable Component Scores

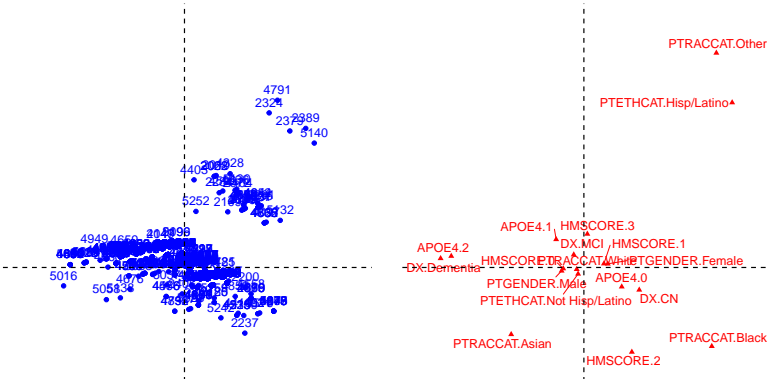


MCA:
Participants Component Scores



MCA

Component 2. Explained variance: 22.93%



Component 1. Explained variance: 62.2%

A very important detour

	PTGENDER	PTETHCAT
5023	Female	Not Hisp/Latino
5026	Female	Not Hisp/Latino
5027	Male	Not Hisp/Latino
5028	Male	Not Hisp/Latino
5031	Female	Hisp/Latino
5037	Male	Not Hisp/Latino
5040	Female	Not Hisp/Latino
5047	Female	Not Hisp/Latino
5054	Female	Not Hisp/Latino
5058	Male	Not Hisp/Latino
5063	Female	Not Hisp/Latino

Two variables with strictly two levels (i.e., binary data)

A very important detour

	PTGENDER.Male	PTGENDER.Female	PTETHCAT.Not Hisp/Latino	PTETHCAT.Hisp/Latino
5023	0	1	1	0
5026	0	1	1	0
5027	1	0	1	0
5028	1	0	1	0
5031	0	1	0	1
5037	1	0	1	0
5040	0	1	1	0
5047	0	1	1	0
5054	0	1	1	0
5058	1	0	1	0
5063	0	1	1	0

Disjunctive coding of two variables with strictly two levels (i.e., binary data) into four columns

A very important detour

	PTGENDER	PTETHCAT
5023	Female	Not Hisp/Latino
5026	Female	Not Hisp/Latino
5027	Male	Not Hisp/Latino
5028	Male	Not Hisp/Latino
5031	Female	Hisp/Latino
5037	Male	Not Hisp/Latino
5040	Female	Not Hisp/Latino
5047	Female	Not Hisp/Latino
5054	Female	Not Hisp/Latino
5058	Male	Not Hisp/Latino
5063	Female	Not Hisp/Latino

Two variables with strictly two levels (i.e., binary data)

A very important detour

	PTGENDER	PTETHCAT
5023	1	0
5026	1	0
5027	0	0
5028	0	0
5031	1	1
5037	0	0
5040	1	0
5047	1	0
5054	1	0
5058	0	0
5063	1	0

Binary coding of two variables with strictly two levels (i.e., binary data) in two columns

A very important detour

	PTGENDER	PTETHCAT
5023	0	1
5026	0	1
5027	1	1
5028	1	1
5031	0	0
5037	1	1
5040	0	1
5047	0	1
5054	0	1
5058	1	1
5063	0	1

Alternate but equivalent binary coding of two variables with strictly two levels (i.e., binary data) in two columns

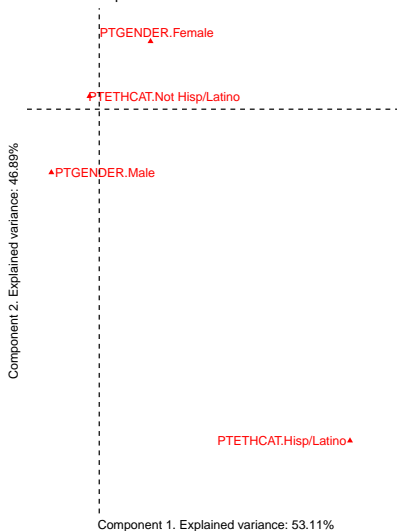
Always a bad idea?

- ▶ MCA on the disjunctive coded data

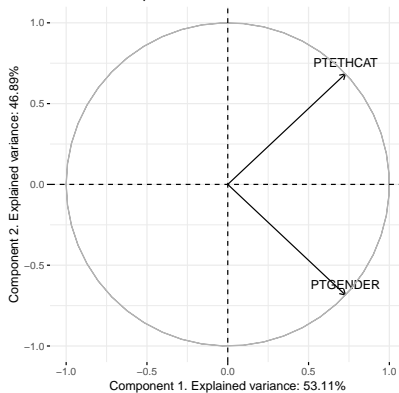
Always a bad idea?

- ▶ MCA on the disjunctive coded data
- ▶ PCA on the binary coded data

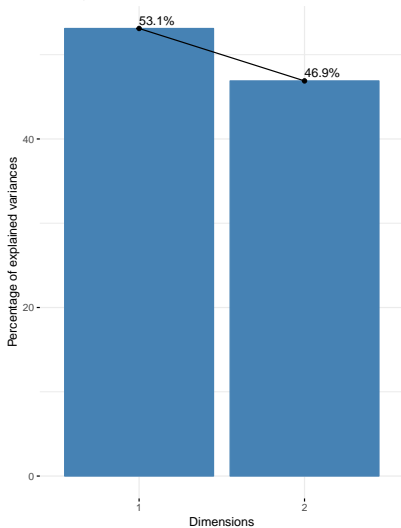
MCA:
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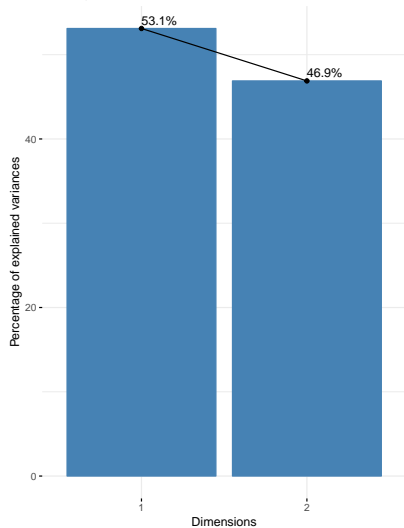
PCA:
Variable-Component Correlations



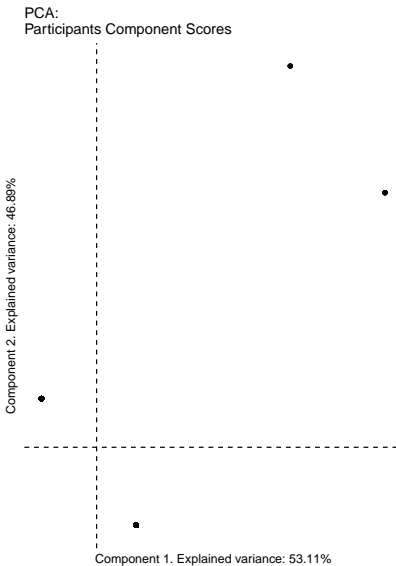
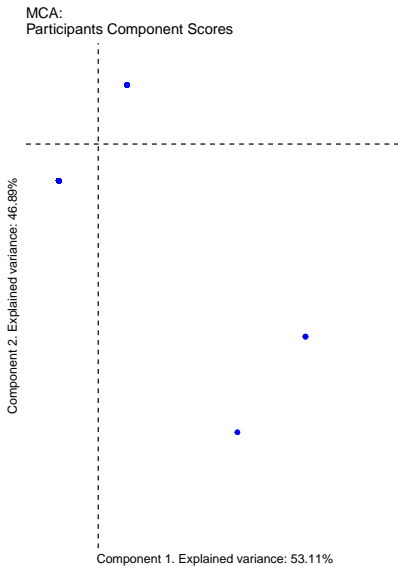
MCA:
Scree plot



PCA:
Scree plot



Oh, weird!



Component 2 is "flipped"
We will revisit this

	PCA Comp. 1	PCA Comp. 2
MCA Comp. 1	1	0
MCA Comp. 2	0	-1

Oh, double weird!

Let's get weird

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

Let's get weird

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

► $\phi = 0.06$

Let's get weird

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

- ▶ $\phi = 0.06$
- ▶ Deep connections between χ^2 , Normal, binomial (and others)

Let's get weird

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

- ▶ $\phi = 0.06$
- ▶ Deep connections between χ^2 , Normal, binomial (and others)
- ▶ We can expand the idea of “binary” or “binomial”

An old friend

	mPACCtrailsB	FDG
5023	1.12	0.13
5026	0.46	-1.31
5027	-2.77	-1.48
5028	-1.59	-0.97
5031	-0.92	-0.87
5037	-1.86	-2.00
5040	0.94	-0.21
5047	-0.25	3.05
5054	-0.80	-1.05
5058	-1.12	-2.13
5063	-2.31	-2.49

We perform(ed) PCA on these data

Escofier's Geometric Magic

- ▶ One of the “fuzzy” or “bipolar” coding schemes

Escofier's Geometric Magic

- ▶ One of the “fuzzy” or “bipolar” coding schemes
- ▶ Take each Z-scored continuous variable

Escofier's Geometric Magic

- ▶ One of the “fuzzy” or “bipolar” coding schemes
- ▶ Take each Z-scored continuous variable
- ▶ Duplicate it as $\left[\frac{1-Z}{2} \frac{1+Z}{2} \right]$

Escofier's Geometric Magic

	mPACCtrailsB-	mPACCtrailsB+	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	1.57	-0.57
5063	1.66	-0.66	1.74	-0.74

Escofier's Geometric Magic

	mPACCtrailsB-	mPACCtrailsB+	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	1.57	-0.57
5063	1.66	-0.66	1.74	-0.74

- Row sums are total number of *original* variables

Escofier's Geometric Magic

	mPACCtrailsB-	mPACCtrailsB+	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	1.57	-0.57
5063	1.66	-0.66	1.74	-0.74

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g., FDG) is total number of rows

Escofier's Geometric Magic

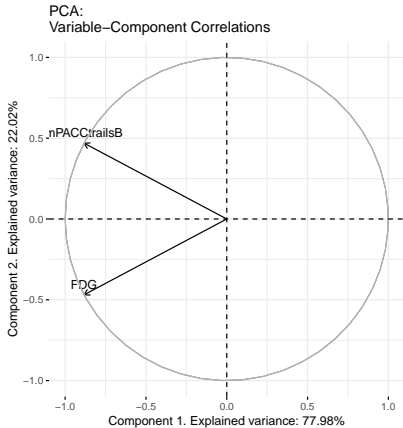
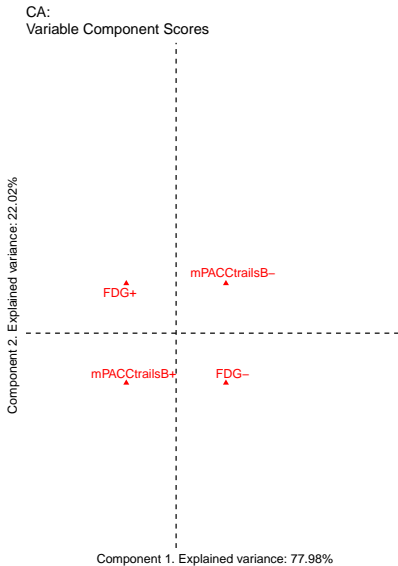
	mPACCtrailsB-	mPACCtrailsB+	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	1.57	-0.57
5063	1.66	-0.66	1.74	-0.74

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g., FDG) is total number of rows
- ▶ Sum of the table is rows \times columns

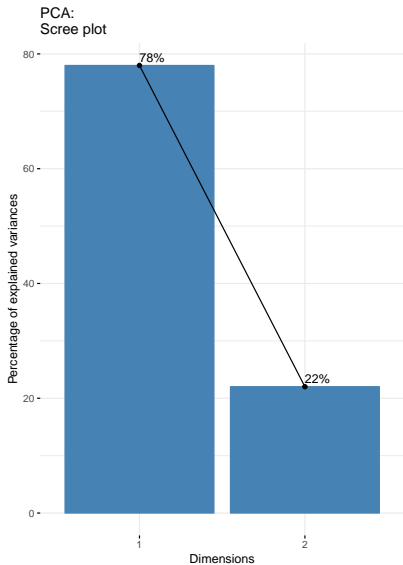
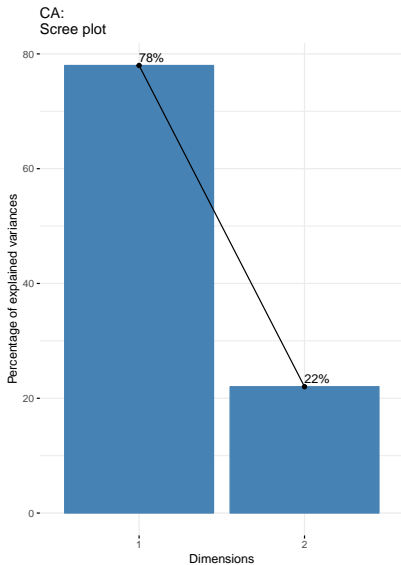
Escofier's Geometric Magic

	mPACCtrailsB-	mPACCtrailsB+	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	1.57	-0.57
5063	1.66	-0.66	1.74	-0.74

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g., FDG) is total number of rows
- ▶ Sum of the table is rows \times columns
- ▶ *These behave like disjunctive data!*

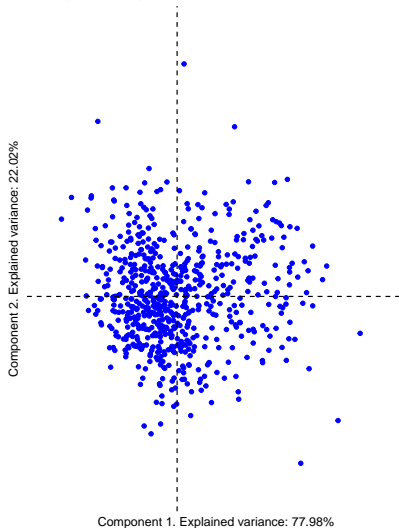


Oh, interesting!
Take note: each variable has two "poles"

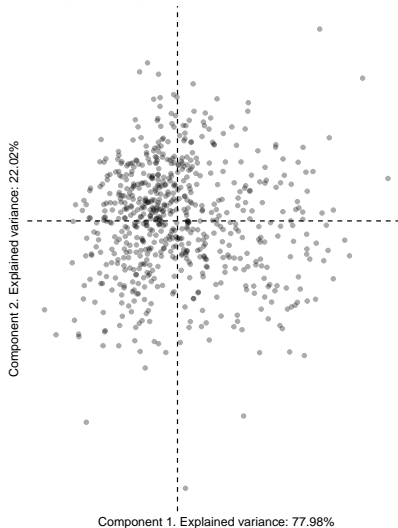


Oh, weird!

CA:
Participants Component Scores

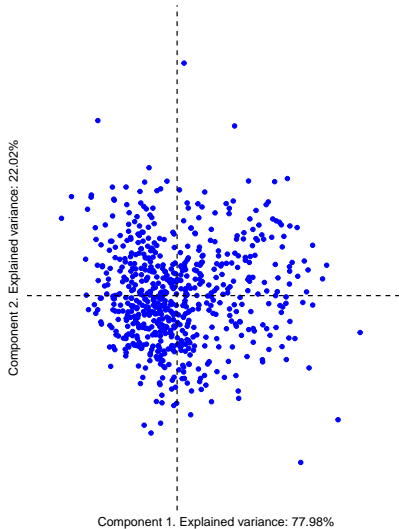


PCA:
Participants Component Scores

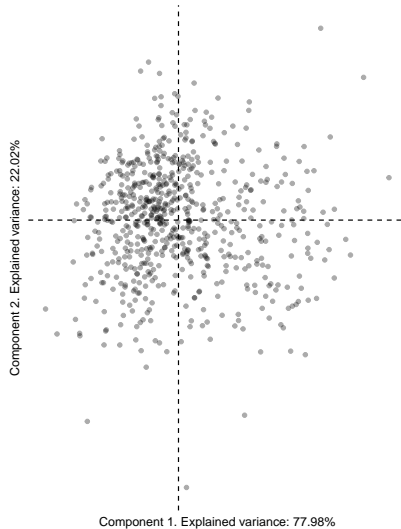


Oh, double weird!

CA:
Participants Component Scores



PCA:
Participants Component Scores



Flips: They don't matter.

	PCA Comp. 1	PCA Comp. 2
CA Comp. 1	1	0
CA Comp. 2	0	-1

Flips: They don't matter.

Escofier's Geometric Trick

- ▶ Apply PCA to continuous data or

Escofier's Geometric Trick

- ▶ Apply PCA to continuous data or
- ▶ Apply CA to “Escofier transformed” data

Thermometer

- ▶ For ordinal data

Thermometer

- ▶ For ordinal data
- ▶ Another “fuzzy” or “bipolar” coding

Thermometer

- ▶ For ordinal data
- ▶ Another “fuzzy” or “bipolar” coding
- ▶ More Escofier Geometric Magic

Thermometer

- ▶ For ordinal data
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 - ▶ Subtract the maximum (minimum is now 0)

Thermometer

- ▶ For ordinal data
- ▶ Another “fuzzy” or “bipolar” coding
- ▶ More Escofier Geometric Magic
 - ▶ Subtract the maximum (minimum is now 0)
 - ▶ $\left[\frac{\max(x) - x}{\max} \quad \frac{x - \min(x)}{\max} \right]$

Thermometer

- ▶ For ordinal data
- ▶ Another “fuzzy” or “bipolar” coding
- ▶ More Escofier Geometric Magic
 - ▶ Subtract the maximum (minimum is now 0)
 - ▶ $\left[\frac{\max(x) - x}{\max} \quad \frac{x - \min(x)}{\max} \right]$
- ▶ Apply CA

More Geometric Magic

	PTEDUCAT	CDRSB	ADAS13	MOCA
5023	18	0.0	6	30
5026	18	1.5	8	24
5027	18	4.0	27	19
5028	16	3.5	20	19
5031	14	2.0	16	20
5037	16	5.0	35	17
5040	18	0.0	8	20
5047	16	1.0	17	24
5054	18	3.5	22	21
5058	20	3.0	17	21
5063	14	2.5	38	16

More Geometric Magic

	PTEDUCAT+	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	0.91	0.09	0.76	0.24	0.07	0.93
5040	0.75	0.25	0.00	1.00	0.17	0.83	0.29	0.71
5047	0.50	0.50	0.18	0.82	0.37	0.63	0.57	0.43
5054	0.75	0.25	0.64	0.36	0.48	0.52	0.36	0.64
5058	1.00	0.00	0.55	0.45	0.37	0.63	0.36	0.64
5063	0.25	0.75	0.45	0.55	0.83	0.17	0.00	1.00

More Geometric Magic

	PTEDUCAT+	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	0.91	0.09	0.76	0.24	0.07	0.93
5040	0.75	0.25	0.00	1.00	0.17	0.83	0.29	0.71
5047	0.50	0.50	0.18	0.82	0.37	0.63	0.57	0.43
5054	0.75	0.25	0.64	0.36	0.48	0.52	0.36	0.64
5058	1.00	0.00	0.55	0.45	0.37	0.63	0.36	0.64
5063	0.25	0.75	0.45	0.55	0.83	0.17	0.00	1.00

- ▶ Row sums are total number of *original* variables

More Geometric Magic

	PTEDUCAT+	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	0.91	0.09	0.76	0.24	0.07	0.93
5040	0.75	0.25	0.00	1.00	0.17	0.83	0.29	0.71
5047	0.50	0.50	0.18	0.82	0.37	0.63	0.57	0.43
5054	0.75	0.25	0.64	0.36	0.48	0.52	0.36	0.64
5058	1.00	0.00	0.55	0.45	0.37	0.63	0.36	0.64
5063	0.25	0.75	0.45	0.55	0.83	0.17	0.00	1.00

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g. EDU) is total number of rows

More Geometric Magic

	PTEDUCAT+	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	0.91	0.09	0.76	0.24	0.07	0.93
5040	0.75	0.25	0.00	1.00	0.17	0.83	0.29	0.71
5047	0.50	0.50	0.18	0.82	0.37	0.63	0.57	0.43
5054	0.75	0.25	0.64	0.36	0.48	0.52	0.36	0.64
5058	1.00	0.00	0.55	0.45	0.37	0.63	0.36	0.64
5063	0.25	0.75	0.45	0.55	0.83	0.17	0.00	1.00

- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g. EDU) is total number of rows
- ▶ Sum of the table is rows \times columns

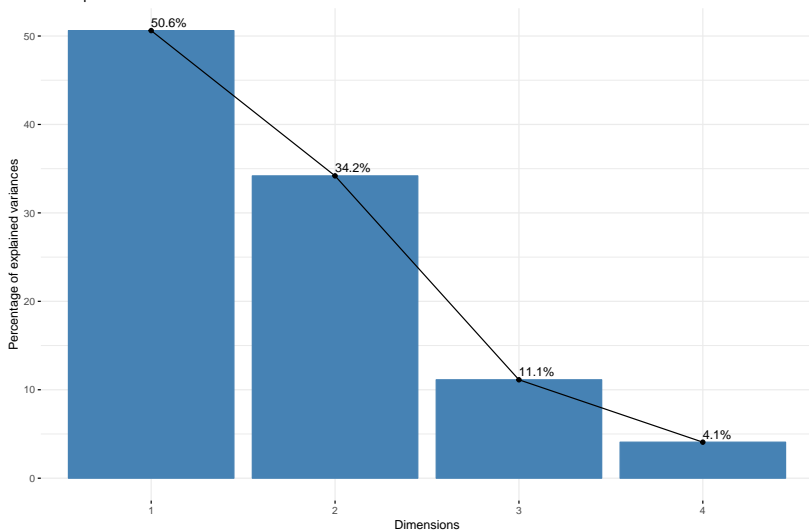
More Geometric Magic

	PTEDUCAT+	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	0.91	0.09	0.76	0.24	0.07	0.93
5040	0.75	0.25	0.00	1.00	0.17	0.83	0.29	0.71
5047	0.50	0.50	0.18	0.82	0.37	0.63	0.57	0.43
5054	0.75	0.25	0.64	0.36	0.48	0.52	0.36	0.64
5058	1.00	0.00	0.55	0.45	0.37	0.63	0.36	0.64
5063	0.25	0.75	0.45	0.55	0.83	0.17	0.00	1.00

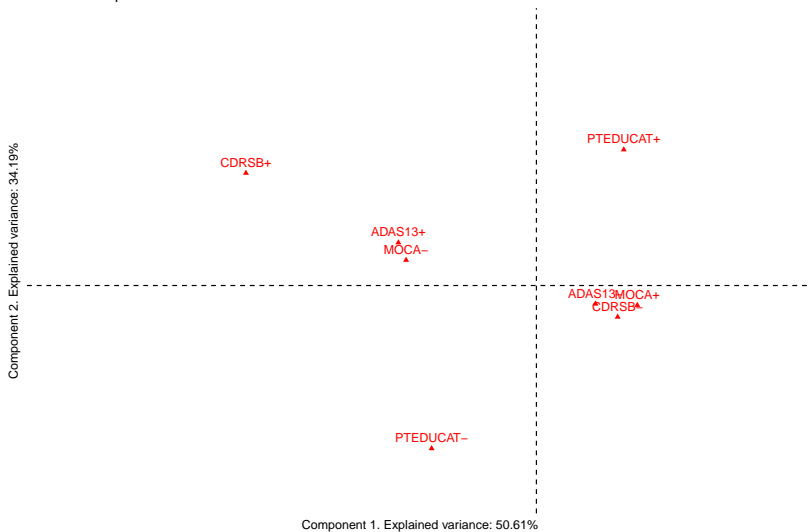
- ▶ Row sums are total number of *original* variables
- ▶ Sum within a variable (e.g. EDU) is total number of rows
- ▶ Sum of the table is rows \times columns
- ▶ *These behave like disjunctive data!*

Let's take a look

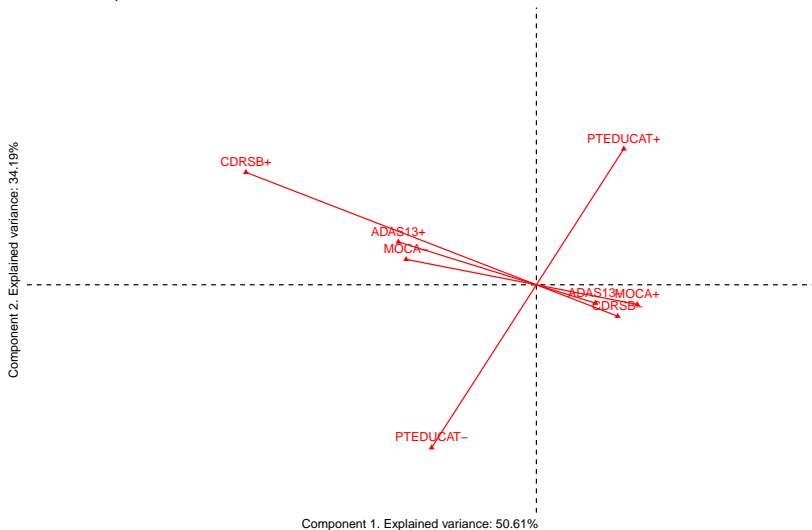
CA:
Scree plot



CA:
Variable Component Scores

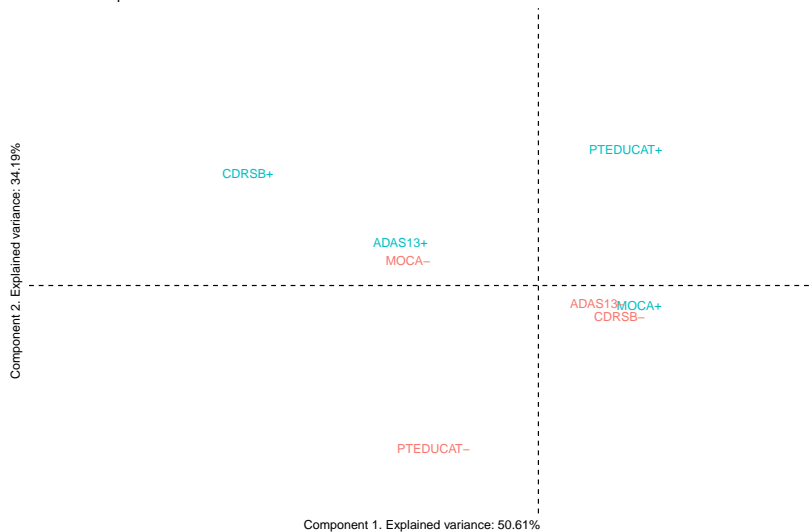


CA:
Variable Component Scores

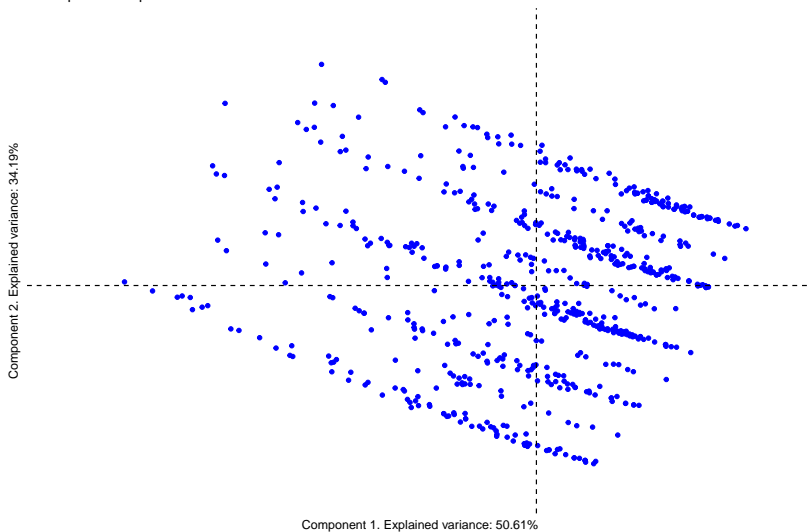


Special properties: Bipolar coding passes through 0.

CA:
Variable Component Scores

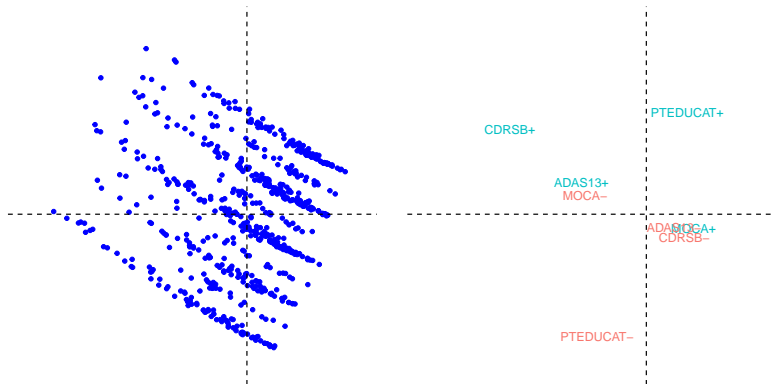


CA:
Participants Component Scores



CA

Component 2. Explained variance: 34.19%



Component 1. Explained variance: 50.61%

Thermometer vs. Disjunctive

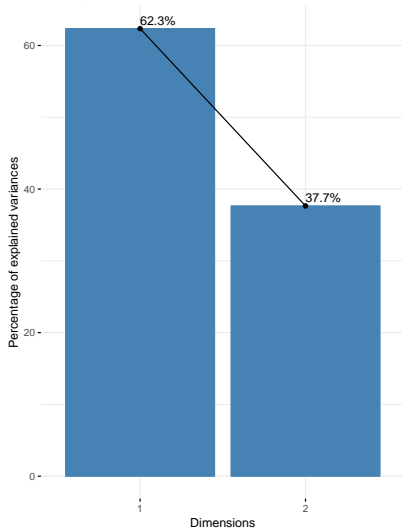
- ▶ Sometimes data could be either

Thermometer vs. Disjunctive

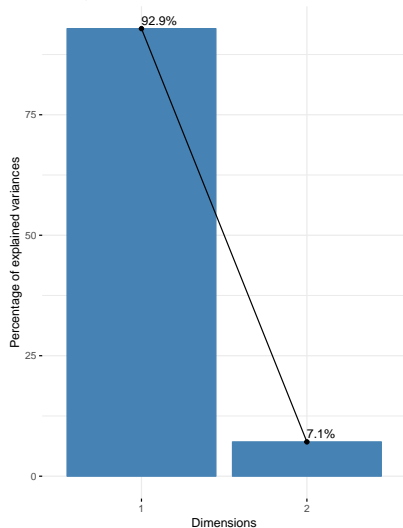
- ▶ Sometimes data could be either
- ▶ Let's analyze it both ways

	APOE4	HMSCORE
5023	0	0
5026	1	1
5027	0	1
5028	2	1
5031	0	1
5037	1	1
5040	0	1
5047	2	1
5054	1	0
5058	0	0
5063	1	1

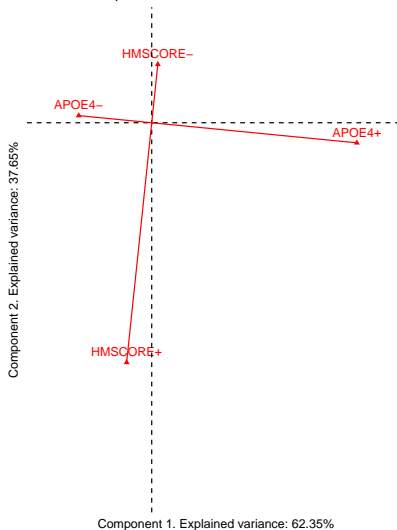
CA (thermometer):
Scree plot



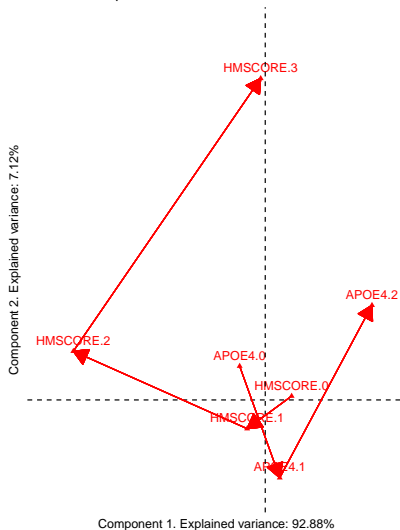
MCA (disjunctive):
Scree plot



CA (thermometer):
Variable Component Scores



MCA (disjunctive):
Variable Component Scores



Thermometer vs. Disjunctive

- ▶ For a small (reasonable) number of levels: disjunctive

Thermometer vs. Disjunctive

- ▶ For a small (reasonable) number of levels: disjunctive
- ▶ Otherwise: thermometer

Thermometer vs. Disjunctive

- ▶ For a small (reasonable) number of levels: disjunctive
- ▶ Otherwise: thermometer
- ▶ Interpretation:

Thermometer vs. Disjunctive

- ▶ For a small (reasonable) number of levels: disjunctive
- ▶ Otherwise: thermometer
- ▶ Interpretation:
 - ▶ Thermometer is “easier”

Thermometer vs. Disjunctive

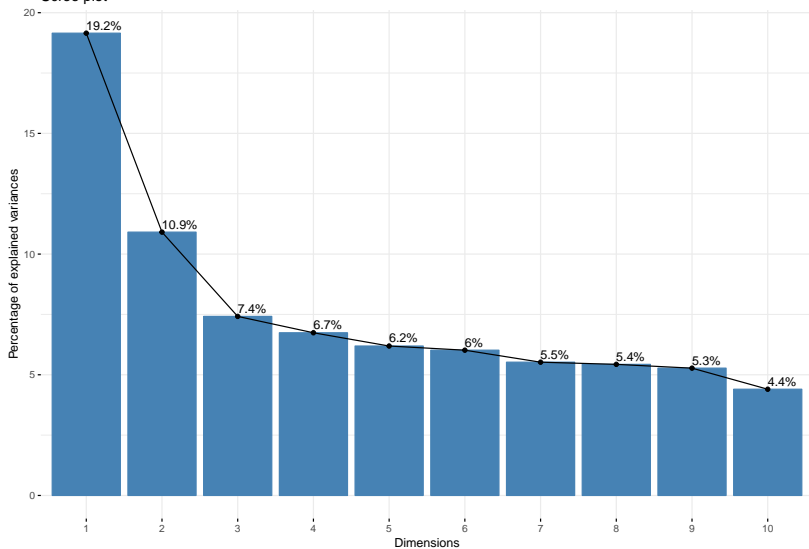
- ▶ For a small (reasonable) number of levels: disjunctive
- ▶ Otherwise: thermometer
- ▶ Interpretation:
 - ▶ Thermometer is “easier”
 - ▶ Disjunctive is more informative

All of the data

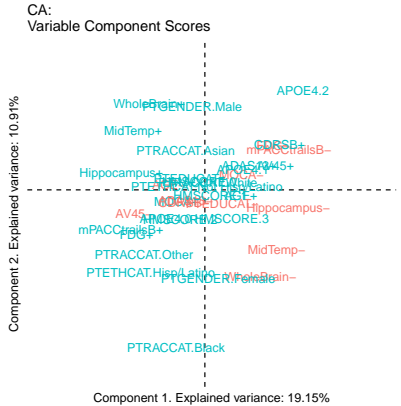
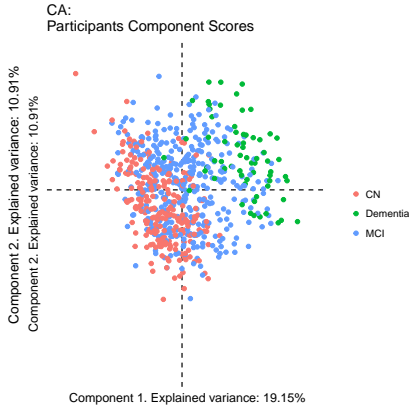
All of the data

	DX	AGE	PTGENDER	PTEDUCAT	PTETHCAT	PTRACCAT	APOE4	FDG	AV45	CDRSB	ADAS13	MOCA	WholeBrain	Hippocampus	MidTemp	mPACtrailsB	HMSCORE
5023	CN	63.9	Female	18	Not Hisp/Latino	Asian	0	1.29	1.03	0.0	6	30	1057351.0	7904	21306	1.81	0
5026	MCI	70.5	Female	18	Not Hisp/Latino	White	1	1.08	1.44	1.5	8	24	1023057.3	8051	16501	-1.45	1
5027	Dementia	75.5	Male	18	Not Hisp/Latino	White	0	1.06	1.44	4.0	27	19	986723.7	6534	17437	-17.27	1
5028	Dementia	61.9	Male	16	Not Hisp/Latino	White	2	1.13	1.38	3.5	20	19	1182704.6	7481	20797	-11.50	1
5031	MCI	80.2	Female	14	Hisp/Latino	White	0	1.14	1.52	2.0	16	20	908133.9	5040	19032	-8.21	1
5037	Dementia	67.3	Male	16	Not Hisp/Latino	Black	1	0.98	1.21	5.0	35	17	1161499.6	5831	21428	-12.80	1
5040	CN	75.9	Female	18	Not Hisp/Latino	Black	0	1.24	1.01	0.0	8	20	943160.6	7994	16634	0.94	1
5047	MCI	68.8	Female	16	Not Hisp/Latino	Black	2	1.70	1.48	1.0	17	24	1070406.1	7920	22043	-4.90	1
5054	Dementia	74.0	Female	18	Not Hisp/Latino	White	1	1.12	1.43	3.5	22	21	1138040.1	6580	20836	-7.63	0
5058	Dementia	61.8	Male	20	Not Hisp/Latino	Asian	0	0.97	1.54	3.0	17	21	1195549.3	7318	22757	-9.18	0
5063	Dementia	71.5	Female	14	Not Hisp/Latino	White	1	0.92	1.61	2.5	38	16	817421.2	5364	12542	-15.03	1

Scree plot



CA:
Everything!



Component 1. Explained variance: 19.15%

Resampling

Background

- ▶ Generally the Gifi or Benzecri principles

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- ▶ Pause!

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My beliefs

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 - ▶ Provides critical diagnostics

Definitions

- ▶ Permutation: break relationships in the data

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- ▶ Split-half: mutually exclusive sets

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- ▶ Permutation: break relationships in the data
- ▶ Split-half: mutually exclusive sets
- ▶ Bootstrap: resample with reselection

Base data

	VARIABLE 1	VARIABLE 2	VARIABLE 3
OBS. 1	a	b	c
OBS. 2	d	e	f
...
OBS. N-1	u	v	w
OBS. N	x	y	z

Tiny illustrative data

Permutation

	VARIABLE 1	VARIABLE 2	VARIABLE 3
OBS. 1	x	e	c
OBS. 2	u	b	f
...
OBS. N-1	a	v	z
OBS. N	d	y	w

Tiny permuted illustrative data

Split-half

	VARIABLE 1	VARIABLE 2	VARIABLE 3
OBS. 1	a	b	c
OBS. 3	g	h	i
...

	VARIABLE 1	VARIABLE 2	VARIABLE 3
OBS. 42	α	π	ω
OBS. 2	d	e	f
...

Tiny split half illustrative data

Bootstrap

	VARIABLE 1	VARIABLE 2	VARIABLE 3
OBS. 42	α	π	ω
OBS. 42	α	π	ω
...
OBS. 1	a	b	c
OBS. N-1	u	v	w

Tiny bootstrap illustrative data

Uses in PCA & CA

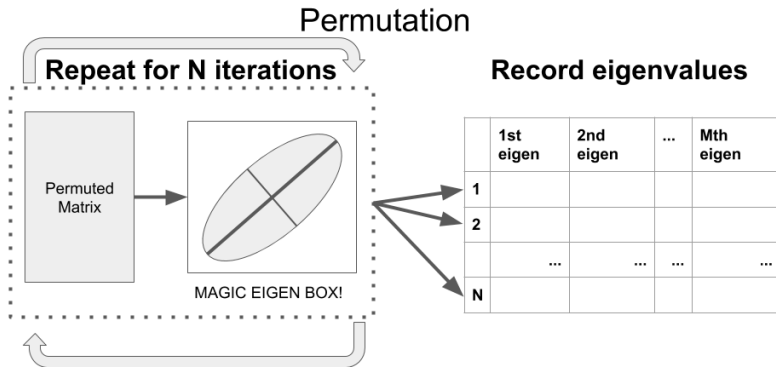
- ▶ Permutation: Effect size tests of components

Uses in PCA & CA

- ▶ Permutation: Effect size tests of components
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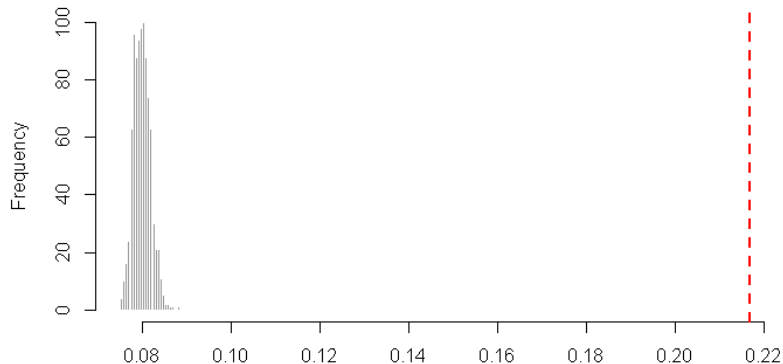
Uses in PCA & CA

- ▶ Permutation: Effect size tests of components
- ▶ Split-half: Replication of components
- ▶ Bootstrap: Stability of variables



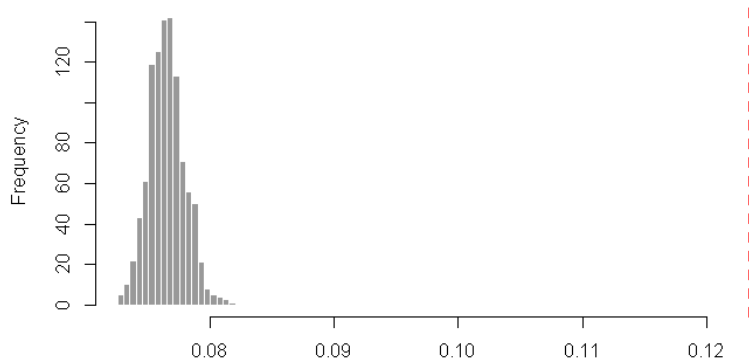
Permutation diagram

First Component Permutation Distribution



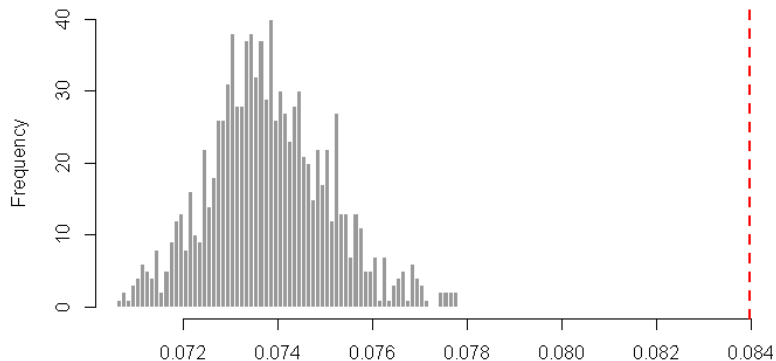
Permutation: First component

Second Component Permutation Distribution



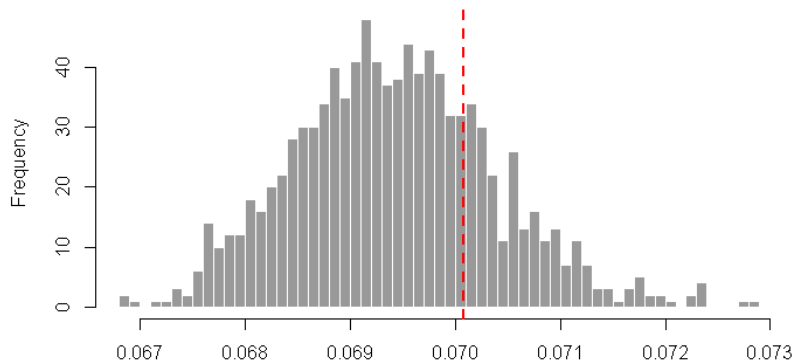
Permutation: Second component

Third Component Permutation Distribution



Permutation: Third component

Fifth Component Permutation Distribution



Permutation: Fifth component

Iterations: 1000

Comp.1	Comp.2	Comp.3	Comp.4	Comp.5	Comp.6	Comp.7
0.001	0.001	0.001	0.001	0.236	0.255	1

Comp.8	Comp.9	Comp.10	Comp.11	Comp.12	Comp.13	Comp.14
0.999	0.998	1	1	1	1	1

Comp.15	Comp.16	Comp.17	Comp.18	Comp.19	Comp.20	Comp.21
1	1	1	1	1	1	1

Permutation: p-values

Another detour

- ▶ p-values should (inversely) follow the scree

Another detour

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- ▶ Diagnostic tests:

Another detour

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Another detour

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- ▶ Diagnostic tests:
 - ▶ Large or erratic jumps
 - ▶ First or first few $ps \geq .5$

Conclusions

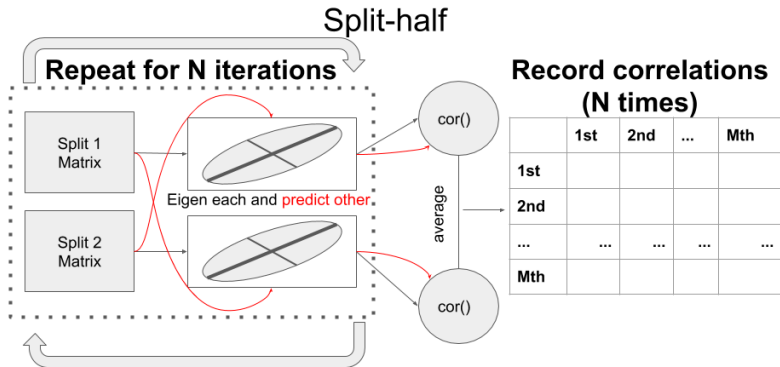
- ▶ First few components have larger than expected effect sizes

Conclusions

- ▶ First few components have larger than expected effect sizes
 - ▶ More variance than null

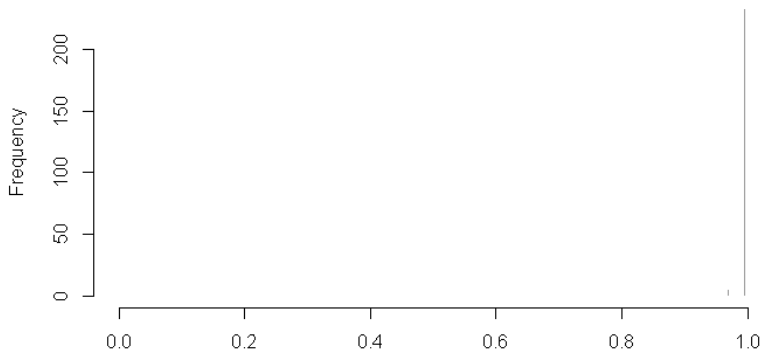
Conclusions

- ▶ First few components have larger than expected effect sizes
 - ▶ More variance than null
- ▶ We do not know if these generalize



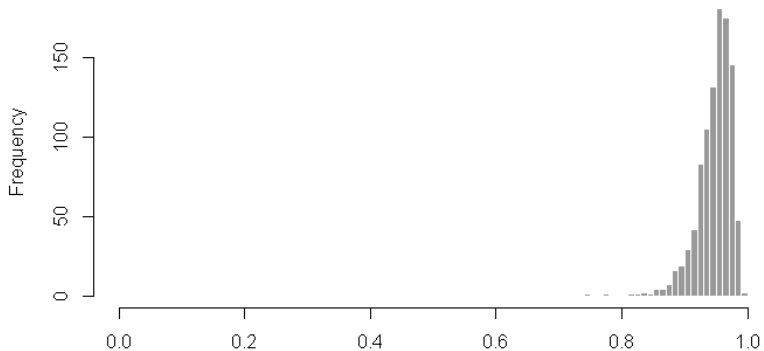
Split half diagram

First Component Split Correlations Distribution



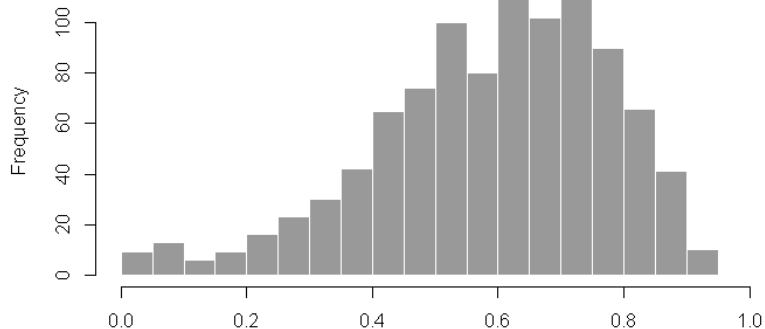
Split-half correlations: First component

Second Component Split Correlations Distribution

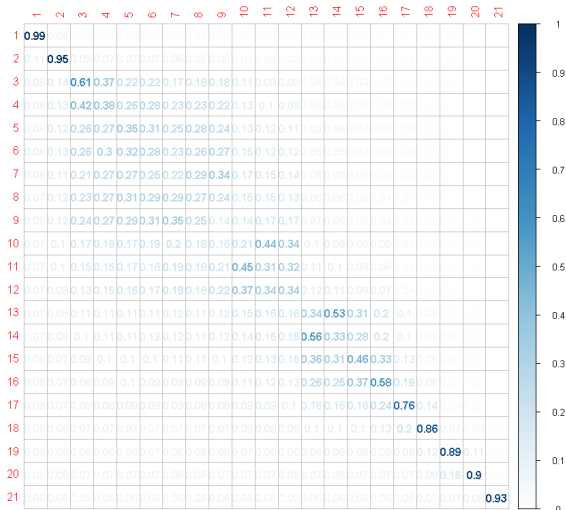


Split-half correlations: Second component

Third Component Split Correlations Distribution



Split-half correlations: Third component



Split-half correlations: All components

Conclusions

- ▶ We do *sort of* know if these generalize

Conclusions

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Conclusions

- ▶ We do *sort of* know if these generalize
- ▶ First two really do
- ▶ Next few: Maybe
- ▶ Key observation:
 - ▶ Components *flip* order!
 - ▶ We need to question the meaning of order of components in our data

Bootstrap goes last

- ▶ You need to know the number of components to interpret

Bootstrap goes last

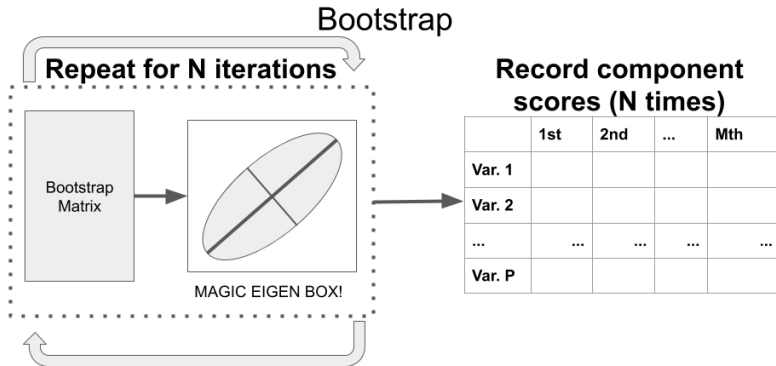
- ▶ You need to know the number of components to interpret
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Bootstrap goes last

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- ▶ Now you can interpret variables *per* component
 - ▶ Find the ones that are stable



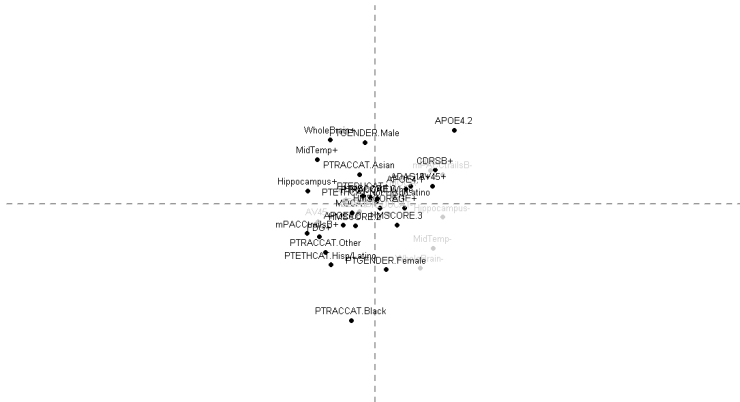
Bootstrap diagram

[illegible]

Component 1. Explained variance: 19.15%

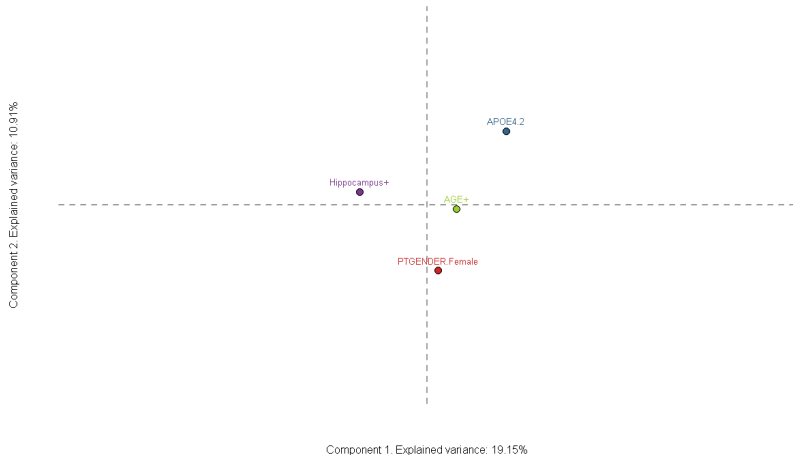
Bootstrap ratios

Component 2. Explained variance: 10.91%



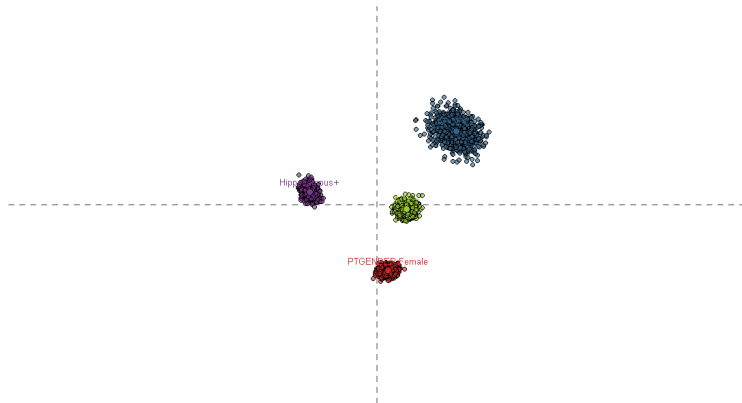
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Bootstrap ratios

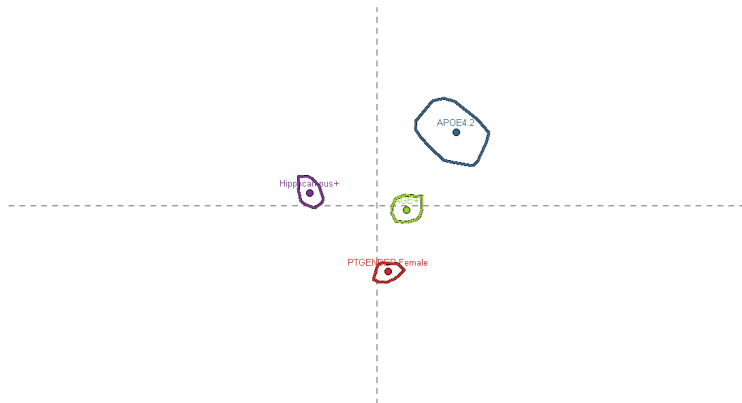
Component 2, Explained variance: 10.91%



Component 1, Explained variance: 19.15%

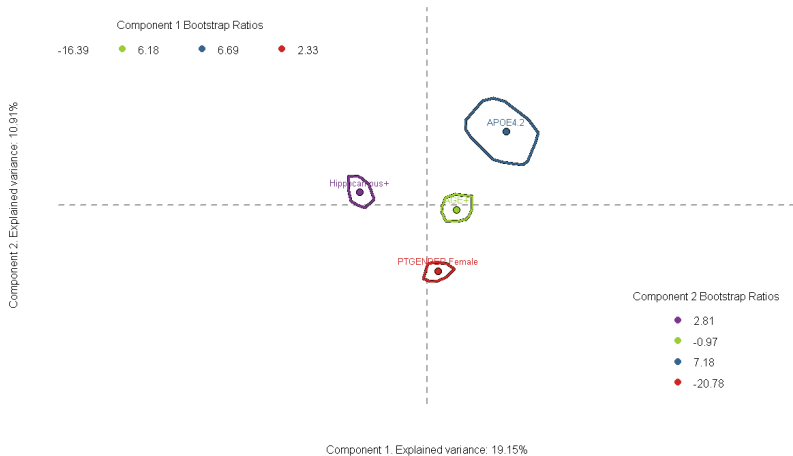
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Bootstrap ratios

Conclusions

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- ▶ APOE4 contributes to both
- ▶ The others generally contribute to one or the other

Final stretch

Things not discussed

- ▶ Corrections

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- ▶ Alternate preprocessing

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Rotation

- ▶ Two compelling examples rotation

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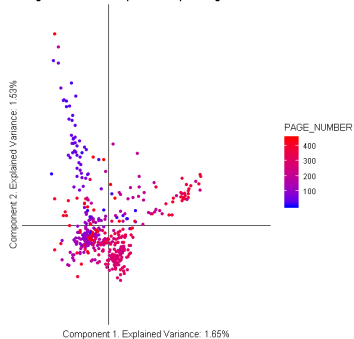
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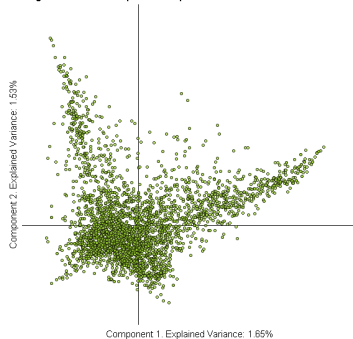
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 - ▶ see http://github.com/fahd09/neurosynth_semantic_map

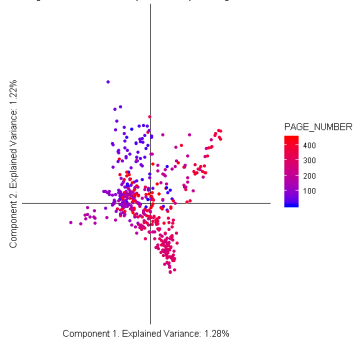
Pages x Words. Component Map of Pages.



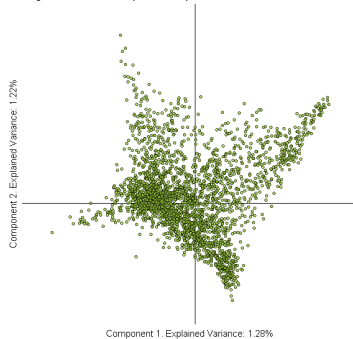
Pages x Words. Component Map of Words.



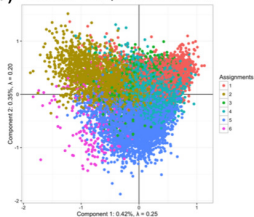
Pages x Lemmas. Component Map of Pages.



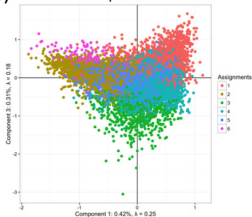
Pages x Lemmas Component Map of Lemmas.



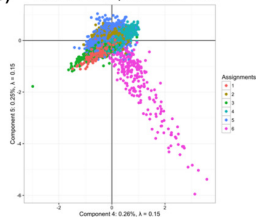
(a) Studies' Components: 1 vs. 2



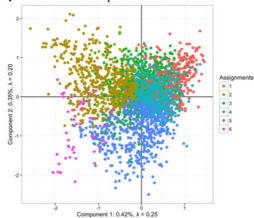
(c) Studies' Components: 1 vs. 3



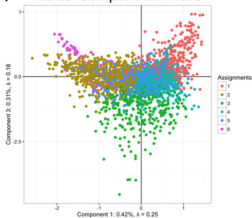
(e) Studies' Components: 4 vs. 5



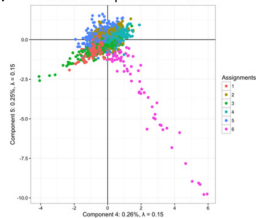
(b) Words' Components: 1 vs. 2



(d) Words' Component: 1 vs. 3



(f) Words' Components: 4 vs. 5



One table

- ▶ Independent Components Analysis

One table

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- ▶ Multidimensional scaling (MDS)

Two tables: Part 1

- ▶ Partial least squares (correlation)

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Two tables: Part 2

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 - ▶ Beaton, Saporta, Abdi (2019)
 - ▶ Mixed data, most two table techniques

More than two tables

- ▶ STATIS

More than two tables

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- ▶ Multiple factor analysis

Not as related

► tSNE

Not as related

- ▶ tSNE
- ▶ UMAP

Not as related

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- ▶ UMAP
- ▶ More akin to non-metric multidimensional scaling

Not as related

- ▶ tSNE
- ▶ UMAP
- ▶ More akin to non-metric multidimensional scaling
 - ▶ Not always a fair comparison

For all types of data

- ▶ Distances (MDS, DiSTATIS, CovSTATIS)

For all types of data

- ▶ Distances (MDS, DiSTATIS, CovSTATIS)
- ▶ Networks (CA)

For all types of data

- ▶ Distances (MDS, DiSTATIS, CovSTATIS)
- ▶ Networks (CA)
 - ▶ More magic!

(Some) References

See the reference sections of these

- ▶ Beaton, D., Saporta, G., Abdi, H., & Alzheimer's Disease Neuroimaging Initiative. (2019). A generalization of partial least squares regression and correspondence analysis for categorical and mixed data: An application with the ADNI data. bioRxiv, 598888.

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- ▶ Beaton, D., Saporta, G., Abdi, H., & Alzheimer's Disease Neuroimaging Initiative. (2019). A generalization of partial least squares regression and correspondence analysis for categorical and mixed data: An application with the ADNI data. bioRxiv, 598888.
- ▶ Beaton, D., Sunderland, K. M., Levine, B., Mandzia, J., Masellis, M., Swartz, R. H., ... & Strother, S. C. (2019). Generalization of the minimum covariance determinant algorithm for categorical and mixed data types. bioRxiv, 333005.

And these

- ▶ Abdi, H., Guillemot, V., Eslami, A., & Beaton, D. (2017). Canonical correlation analysis. Encyclopedia of Social Network Analysis and Mining, 1-16.

And these

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