Principal Components & Multiple Correspondence Analyses with resampling approaches for stability assessments

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RRI RTC

May 01, 2019

Overview

► Something here

Overview

- Something here
- ► Where to find everything



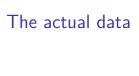
Slide A

 $\mathsf{Slide}\ \mathsf{A}$

Slide B

Slide B

Taxonomy



PCA

Stuff

PCA

How & When to use it

CA

Stuff

CA

How & When to use it

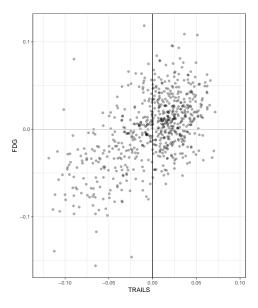
Today

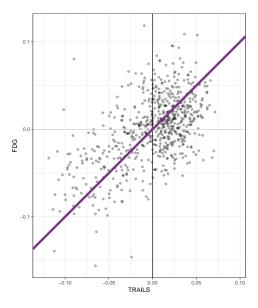
Some alternatives

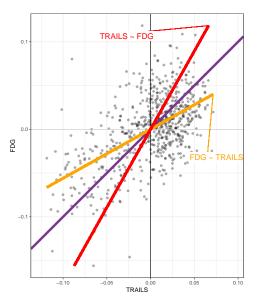
Principal Components Analysis

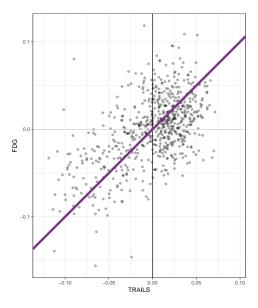


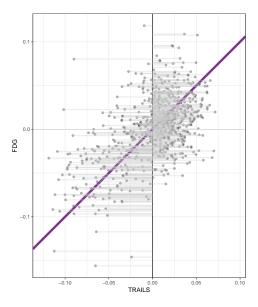
TRAILS

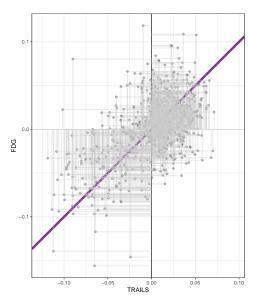


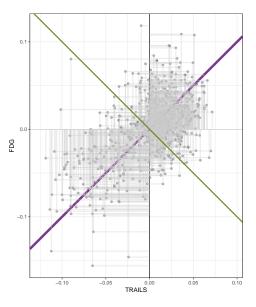




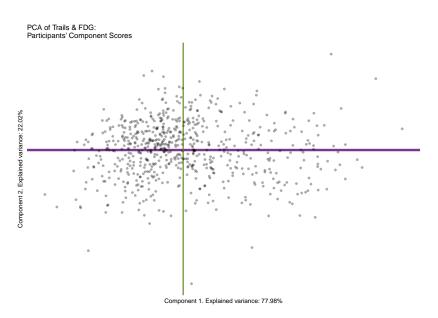




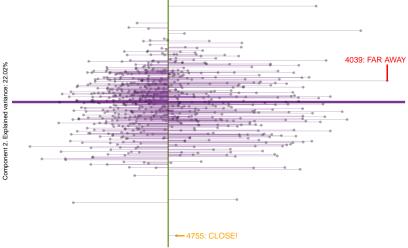




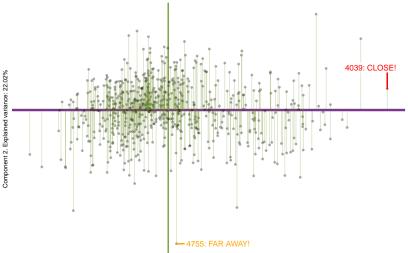
The SVD



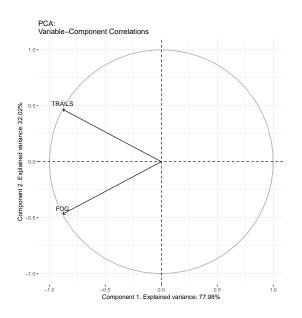
PCA of Trails & FDG: Participants' Component Scores

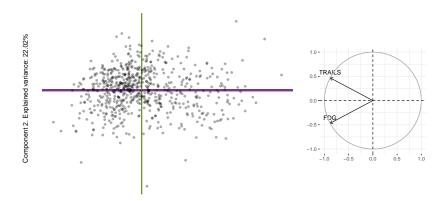


Component 1. Explained variance: 77.98%



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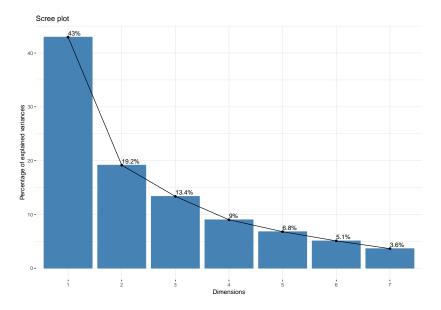


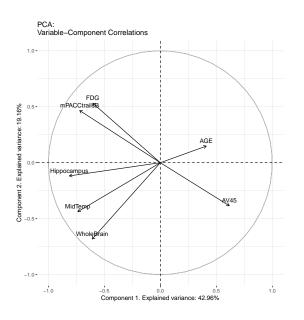
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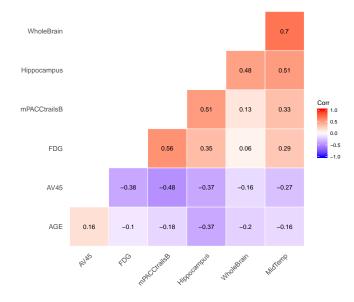
Component 1. Explained variance: 77.98%

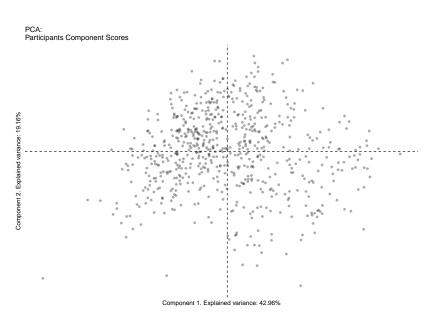
Scaling up

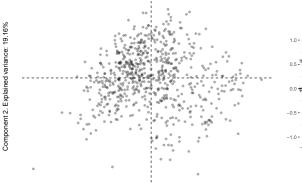
here: show the table and perhaps the code? maybe the code per slide for easy slides?

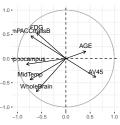






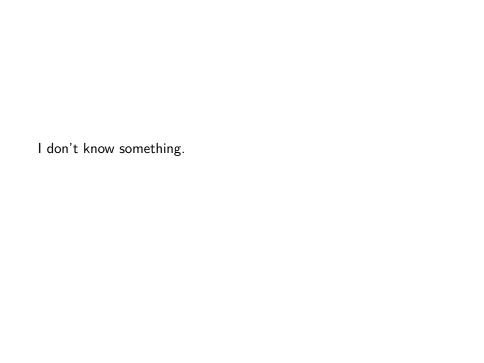






Component 1. Explained variance: 42.96%





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

	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

	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

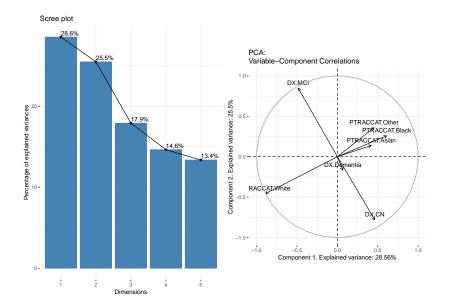
	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
5023	0	1	0	0	0	0	1
5026	1	0	0	1	0	0	0
5027	0	0	1	1	0	0	0
5028	0	0	1	1	0	0	0
5031	1	0	0	1	0	0	0
5037	0	0	1	0	0	1	0
5040	0	1	0	0	0	1	0
5047	1	0	0	0	0	1	0
5054	0	0	1	1	0	0	0
5058	0	0	1	0	0	0	1
5063	0	0	1	1	0	0	0

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

	DX.MCI	DX.CN	DX.Dementia	${\sf PTRACCAT.White}$	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 × columns

A bad idea: PCA



Why is that a bad idea?

	DX.MCI	DX.CN	DX.Dementia	PTRACCAT.White	PTRACCAT.Other	PTRACCAT.Black	PTRACCAT.Asian
DX.MCI	1	-0.815	-0.363	0.045	0.032	-0.043	-0.072
DX.CN	-0.815	1	-0.243	-0.047	0	0.067	0.003
DX.Dementia	-0.363	-0.243	1	0	-0.053	-0.035	0.116
PTRACCAT.White	0.045	-0.047	0	1	-0.562	-0.657	-0.45
PTRACCAT. Other	0.032	0	-0.053	-0.562	1	-0.031	-0.021
PTRACCAT.Black	-0.043	0.067	-0.035	-0.657	-0.031	1	-0.025
PTRACCAT.Asian	-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
DX.MCI	365	0	0	341	11	10	3
DX.CN	0	235	0	213	6	12	4
DX.Dementia	0	0	65	60	0	1	4
PTRACCAT.White	341	213	60	614	0	0	0
PTRACCAT.Other	11	6	0	0	17	0	0
PTRACCAT.Black	10	12	1	0	0	23	0
PTRACCAT.Asian	3	4	4	0	0	0	11

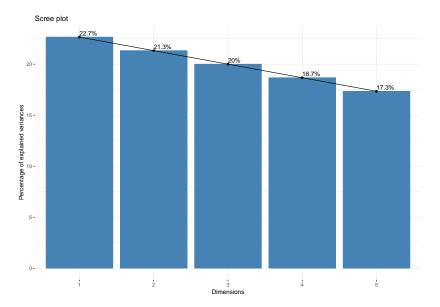
► Correspondence analysis (CA)

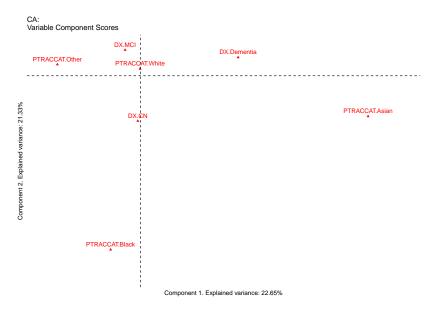
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 - Think of it as a χ^2 PCA

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- ▶ Row and column component scores exist on same scale

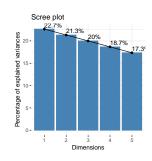
- Correspondence analysis (CA)
 - ► Think of it as a χ^2 PCA
- Deals with categories, counts (amongst others)
- Row and column component scores exist on same scale
 - CA is a bivariate technique





	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

	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

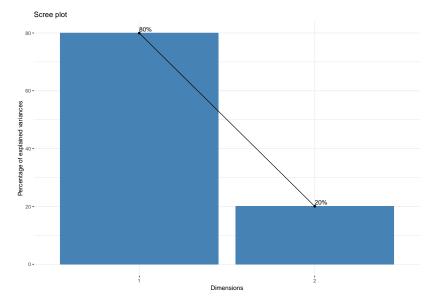


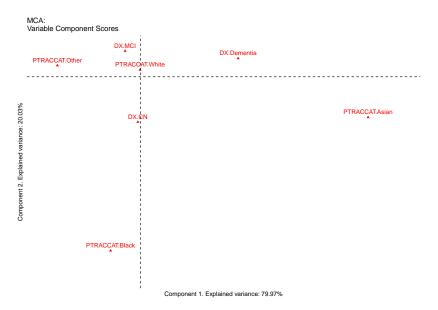
► An extension of CA

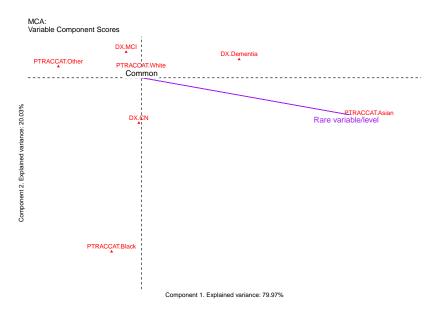
- An extension of CA
- ► Accomodates multiple categorical variables (CA only does 2)

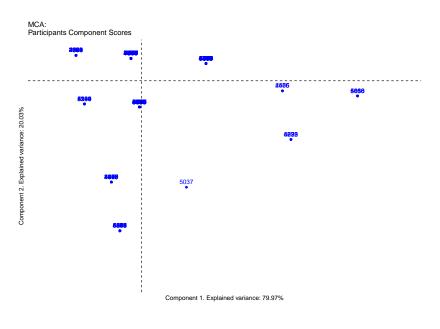
- An extension of CA
- ► Accomodates multiple categorical variables (CA only does 2)
- Corrects the dimensionality

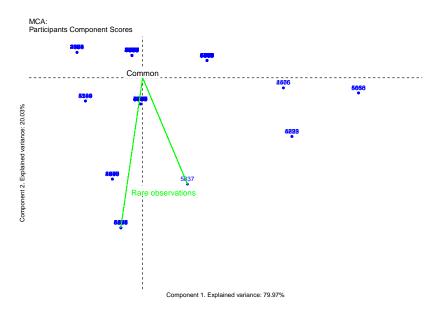
- An extension of CA
- Accomodates multiple categorical variables (CA only does 2)
- Corrects the dimensionality
- ► Has nearly magical properties (we'll see later)











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

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

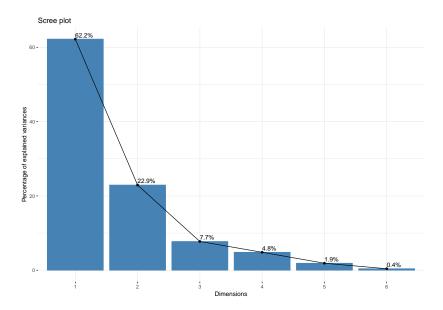
Compare the results

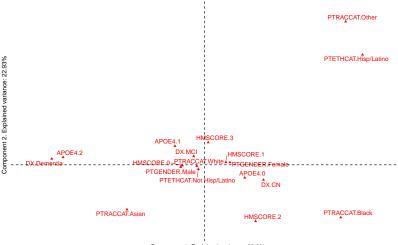
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- ► CA & MCA produce identical results, except MCA:
 - Drops components
 - Corrects explained variance

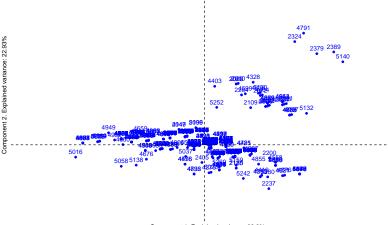
Scaling up

SHow the data here a bit

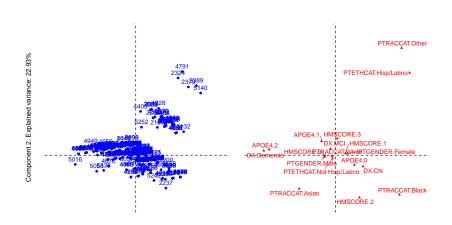




MCA: Participants Component Scores



Component 1. Explained variance: 62.2%



Component 1. Explained variance: 62.2%

	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)

	PTGENDER.Male	PTGENDER.Female	${\sf PTETHCAT.Not\ Hisp/Latino}$	${\sf PTETHCAT.Hisp}/{\sf Latino}$
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

	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)

	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

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

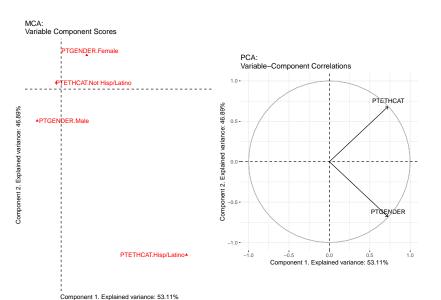
Alternate but equivalent binary coding of two variables with strictly two levels (i.e., binary data) in two columns

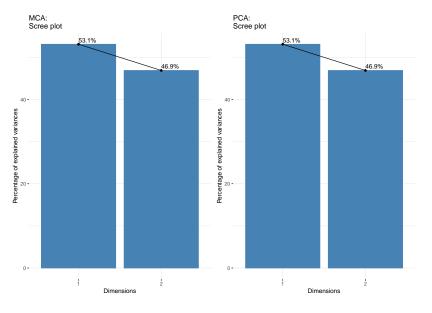
Always a bad idea?

► MCA on the disjunctive coded data

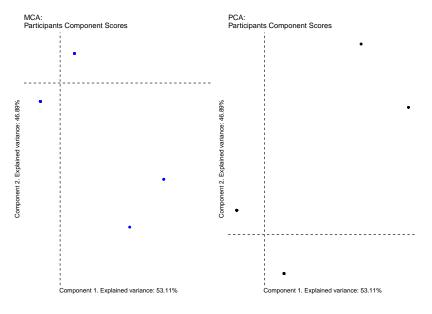
Always a bad idea?

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





Oh, weird!



Component 2 is "flipped"
We will revisit this

MCA Comp. 1 1 MCA Comp. 2 0 -			. e, t eep. =
MCA Comp. 2 0 -	MCA Comp. 1	1	0
•	MCA Comp. 2	0	-1

Oh, double weird!

PCA Comp. 1 PCA Comp. 2

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

 $[\]phi = 0.06$

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

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

	PTGENDER	PTETHCAT
PTGENDER	1.00	0.06
PTETHCAT	0.06	1.00

- $\phi = 0.06$
- ightharpoonup 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 PCA on these data

▶ One of the "fuzzy" or "bipolar" coding schemes

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- ► Take each Z-scored continuous variable

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

	${\sf mPACCtrailsB-}$	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

	${\sf mPACCtrailsB-}$	${\sf mPACCtrailsB} +$	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

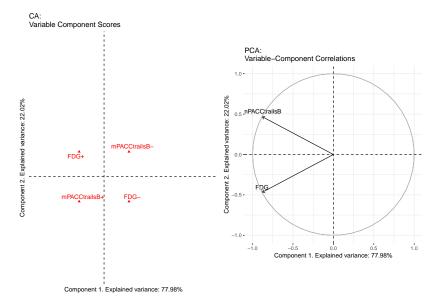
▶ Row sums are total number of *original* variables

	${\sf mPACCtrailsB-}$	${\sf mPACCtrailsB} +$	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

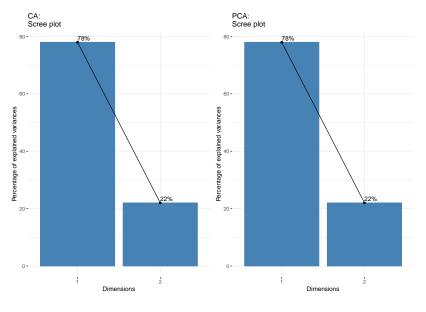
	${\sf mPACCtrailsB-}$	${\sf mPACCtrailsB} +$	FDG-	FDG+
5023	-0.06	1.06	0.43	0.57
5026	0.27	0.73	1.16	-0.16
5027	1.88	-0.88	1.24	-0.24
5028	1.30	-0.30	0.98	0.02
5031	0.96	0.04	0.93	0.07
5037	1.43	-0.43	1.50	-0.50
5040	0.03	0.97	0.60	0.40
5047	0.62	0.38	-1.03	2.03
5054	0.90	0.10	1.03	-0.03
5058	1.06	-0.06	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
- \triangleright Sum of the table is rows \times columns

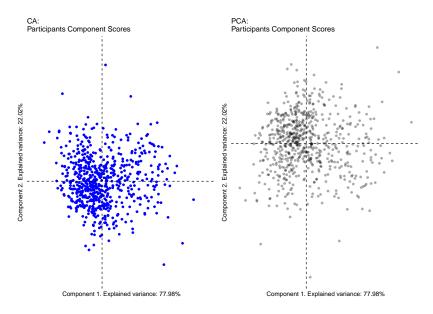


Oh, interesting!

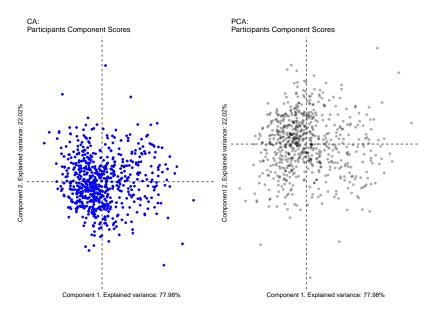
Take note: each variable has two "poles"



Oh, weird!



Oh, double weird!



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

► For ordinal data

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- ► Another "fuzzy" or "bipolar" coding

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Thermometer

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

	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

5023	ΓEDUCAT+ 0.75	PTEDUCAT-	CDRSB+	CDRSB-	ADAS13+	ADAS13-	MOCA+	MOCA-
5023	0.75	0.25						
3023		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

	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

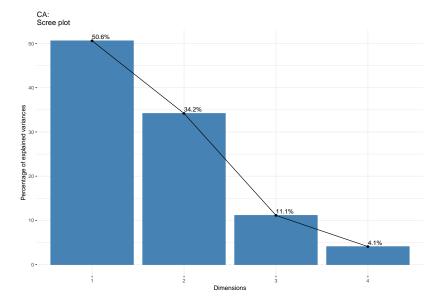
▶ Row sums are total number of *original* variables

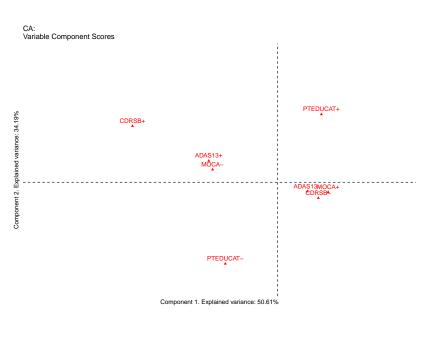
	PTEDUCAT +	PTEDUCAT-	CDRSB +	CDRSB-	ADAS13+	ADAS13-	MOCA +	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
5037	0.50	0.50	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

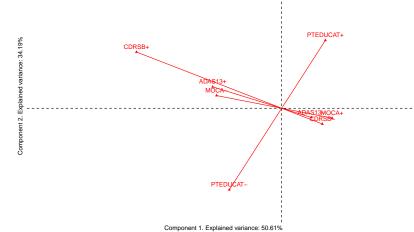
	PTEDUCAT +	PTEDUCAT-	CDRSB +	CDRSB-	ADAS13 +	ADAS13-	MOCA +	MOCA-
5023	0.75	0.25	0.00	1.00	0.13	0.87	1.00	0.00
5026	0.75	0.25	0.27	0.73	0.17	0.83	0.57	0.43
5027	0.75	0.25	0.73	0.27	0.59	0.41	0.21	0.79
5028	0.50	0.50	0.64	0.36	0.43	0.57	0.21	0.79
5031	0.25	0.75	0.36	0.64	0.35	0.65	0.29	0.71
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5040	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

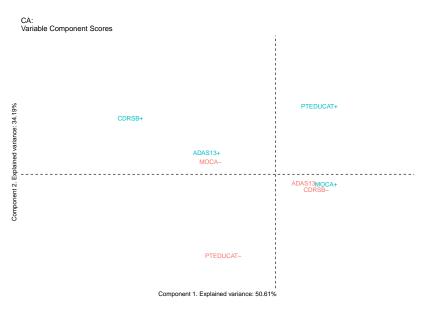
- ▶ Row sums are total number of *original* variables
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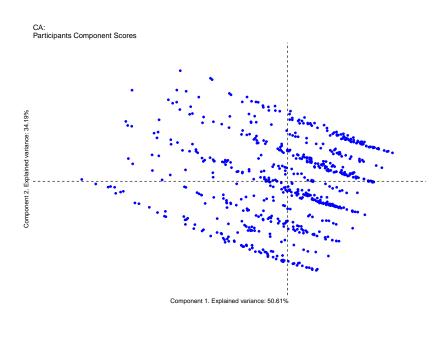


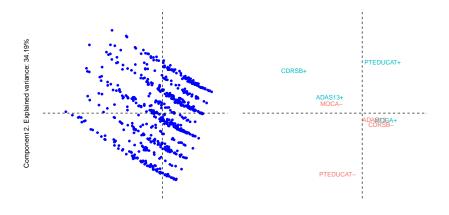










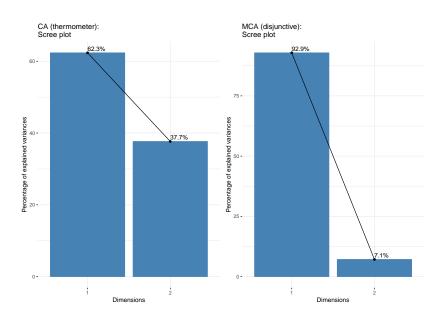


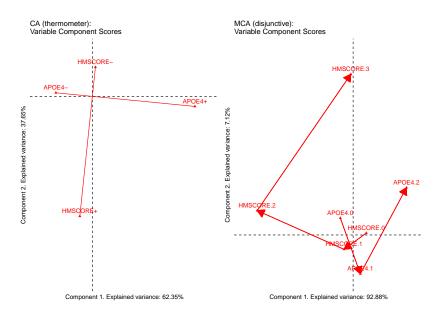
Component 1. Explained variance: 50.61%

Sometimes data could be either

- 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





► For a small (reasonable) number of levels: disjunctive

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- ► Otherwise: thermometer

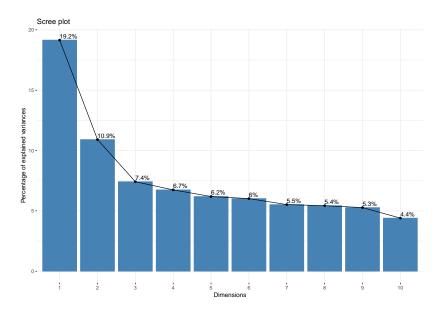
- ► For a small (reasonable) number of levels: disjunctive
- Otherwise: thermometer
- ► Interpretation:

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- Otherwise: thermometer
- Interpretation:
 - ► Thermometer is "easier"

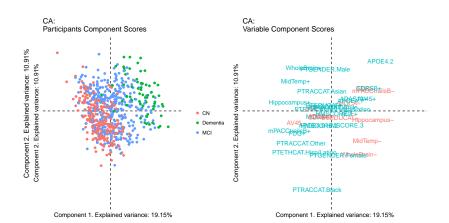
- For a small (reasonable) number of levels: disjunctive
- Otherwise: thermometer
- Interpretation:
 - ► Thermometer is "easier"
 - Disjunctive is more informative

All of the data

	DX	AGE	PTGENDER	PTEDUCAT	PTETHCAT	PTRACCAT	APOE4	FDG	AV45	CDRSB	ADAS13	MOCA	WholeBrain	Hippocampus	MidTemp	mPACCtrailsB	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



CA: Everything!



Component 1. Explained variance: 19.15%

Resampling

FUCK