Correspondence Analysis & Related Methods

Michael Greenacre

SESSION 13: Diagnostics, contributions in weighted PCA and Correspondence Analysis

Inertia contributions in weighted PCA

 PCA is a method of data visualization which represents the true positions of points in a map which comes closest to all the points, closest in sense of weighted least-squares.



The inertia (weighted variance) explained in the map applies to all the points: if we say 83% of the inertia is explained in the map, 71% on the first dimension and 12% on the second, this is a figure calculated for all row (or column) points together.

Inertia contributions in weighted PCA

- This type of "inertia-explained-by-axes" calculation can be made for individual points.
- These more detailed results are aids to interpretation in the form of numerical diagnostics, called contributions.
- Especially when there is not a high percentage of inertia explained by the map, these contributions will help us to identify points which are represented inaccurately.
- The inertias and their percentages tell us how much of the variance in the table is explained by the principal axes. The contributions do the same, but for each point individually, and help us to see:
 - (a) which points are being explained better than others;
 - (b) which points are contributing to the solution more than others.

Geometry of inertia contributions



Total inertia of the cloud of points = $\sum_{i} m_{i} d_{i}^{2} = \sum_{i} m_{i} \sum_{k} f_{ik}^{2} = \sum_{k} \lambda_{k}$ Inertia of *i*-th point = $m_{i} d_{i}^{2} = m_{i} \sum_{k} f_{ik}^{2}$ Inertia contribution of *i*-th point to *k*-th axis = $m_{i} f_{ik}^{2}$

Decomposition of inertia



Inertia contributions for CA of "author"

col	name	mass	qlt	inr	k=:	L co	r ctr	k=2	2 CO1	ctr	
1	a	80	162	10	2	1	0	-19	161	8	1
2	b	16	365	18	86	338	15	-24	27	2	Ĺ
3	c	23	831	60	185	691	102	-83	140	43	İ
4	d	46	920	89	-169	788	170	-69	132	59	İ
5	e	127	357	34	8	12	1	-42	345	60	ĺ
6	f	19	529	28	112	456	32	-45	72	10	İ
7	g	20	344	26	-89	325	21	21	19	2	İ
8	h	65	735	83	-131	721	146	-18	14	6	İ
9	i	70	465	28	23	74	5	54	392	55	İ
10	j	1	28	7	40	9	0	56	18	1	İ
11	k	9	724	43	-241	661	70	75	64	14	ĺ
12	1	43	555	33	89	548	44	-10	7	1	Ĺ
13	m	26	436	35	62	153	13	85	284	50	Ĺ
14	n	69	166	21	-18	54	3	-25	112	12	Ĺ
15	0	77	205	32	-9	12	1	39	193	31	Ĺ
16	р	15	515	51	141	317	39	-112	198	51	ĺ
17	q	1	416	12	357	376	11	-116	40	2	Ĺ
18	r	52	374	35	52	215	18	-45	159	28	Ĺ
19	s	61	413	49	75	374	45	25	40	10	Ĺ
20	t	93	90	13	-9	30	1	12	59	4	Ĺ
21	u	30	283	23	14	14	1	62	268	31	Ĺ
22	v	10	550	37	200	548	50	11	2	0	Ĺ
23	w	26	888	75	-219	883	161	-17	6	2	Ĺ
24	x	1	418	22	292	237	13	256	182	21	Ĺ
25	У	22	899	106	0	0	0	286	899	485	ĺ
26	z	1	576	30 İ	596	511	37	-213	65	10	ĺ

Inertía contributions



<u>Summary:</u> Contributions to inertia

- Each principal inertia can decomposed into parts due to each point, either row points or column points. These contributions explain how each principal axis has been constructed (hence the influence of each point in defining the dimension).
- The inertia of a point is similarly decomposed over all the axes, thanks to using Euclidean-type distance and Pythagoras' theorem. Each component on an axis can be expressed relative to the point inertia and this is the same as the squared cosine (i.e., squared correlation) between the point and the axis. These values can be added over axes and tell you how well the point is represented in the solution space.

R implementation of CA (repeat)

# read in data into data-frame data_set					
# the next 14 commands are all you need to compute CA results					
data.P <- data_set/sum(data_set)					
data.c <- apply(data.P,2,sum)					
data.Dr <- diag(data.r) data.Dc <- diag(data.c)					
<pre>data.Drmh <- diag(1/sqrt(data.r)) data.Dcmh <- diag(1/sqrt(data.c))</pre>					
<pre>data.P <- as.matrix(data.P) data.S <- data.Drmh %*% (data.P-data.r%o%data.c) %*% data.Dcmh data.svd <- svd(data.S)</pre>					
<pre>data.rsc <- data.Drmh%*%data.svd\$u data.csc <- data.Dcmh%*%data.svd\$v data.rpc <- data.rsc%*%diag(data.svd\$d) data.cpc <- data.csc%*%diag(data.svd\$d)</pre>					
# the symmetric map					
<pre>plot(data.rpc[,1],data.rpc[,2],type="n",pty="s") text(data.rpc[,1],data.rpc[,2],label=rownames(data))</pre>					
# now do it in one shot using ca package (first install from CRAN)					
library(ca)					

plot(ca(data_set))

Computation of contributions

compute matrix of contributions for rows and inertias data.rcon <- data.rpc^2 * data.r</pre> apply(data.rcon, 1, sum) # compute contributions and squared correlations data.rctr <- t(t(data.rcon) / apply(data.rcon, 2, sum))</pre> data.rcor <- data.rcon / apply(data.rcon, 1, sum)</pre> # compute qualities in 2-d solution apply(data.rcor[,1:2], 1, sum) # compute matrix of contributions for columns and inertias data.ccon <- data.cpc^2 * data.c apply(data.ccon, 1, sum) # compute contributions and squared correlations data.cctr <- t(t(data.ccon) / apply(data.ccon, 2, sum))</pre> data.ccor <- data.ccon / apply(data.ccon, 1, sum)</pre> # compute qualities in 2-d solution apply(data.ccor[,1:2], 1, sum)

Correspondence Analysis & Related Methods

Michael Greenacre

SESSION 14: 1. CORRESPONDENCE ANALYSIS & CLUSTER ANALYSIS

2. CORRESPONDENCE ANALYSIS & BIPLOT

 Correspondence analysis (CA) is a method of data visualization that reveals continuous structures (the dimensions)



• But in our search for structure in the table we can also consider clustering the rows and columns, to reveal discrete structures (the clusters, or classes):



A símple example

 988 students, males and females classified each according to their parents having been or not to university, cross-tabulated with their choice of studies at high school

F_no	94	43	197	61
F_uni	28	17	103	37
M_no	65	19	51	132
M_uni	17	9	30	85

MΔ

LS PS

NS

Inertia = 0.1848 Chi-square = 182.6

Which two rows can we merge so that inertia

(or chi-square) is reduced the least?

F_no and F_uni: reduces inertia by 0.0070

- F_no: female, parents no university F_uni: female, parents university M_no: male, parents no university M_uni: male, parents university
- NS: non-science MA: mathematics LS: life sciences PS: physical sciences

A símple example

 988 students, males and females classified each according to their parents having been or not to university, cross-tabulated with their choice of studies at high school

	NS	MA	LS	PS	
{F_no, F_uni}	122	60	300	98	
M_no	65	19	51	132	
M_uni	17	9	30	85	
					·

Inertia = 0.1778

Which two rows can we merge so that inertia is reduced the *least*?

M_no and M_uni: reduces inertia by 0.0104

- F_no: female, parents no university F_uni: female, parents university M_no: male, parents no university M_uni: male, parents university
- NS: non-science MA: mathematics LS: life sciences PS: physical sciences

A simple example

988 students, males and females classified each according to their • parents having been or not to university, cross-tabulated with their choice of studies at high school

	113	MA	L3	F3
{F_no, F_uni}	122	60	300	98
{M_no, M_uni}	83	28	81	217

....

NC

Inertia = 0.1674

Which two rows can we merge so that inertia (or chi-square) is reduced the least?

Only two rows left to merge and this reduces inertia by 0.1674

- **F_no**: female, parents no university **F_uni**: female, parents university M_no: male, parents no university M_uni: male, parents university
- NS: non-science MA: mathematics LS: life sciences PS: physical sciences

A simple example

988 students, males and females classified each according to their parents having been or not to university, cross-tabulated with their choice of studies



From Greenacre(1993:118), the critical point for the chi-square is 13.11, that is for the inertia: 13.11/988 = **0.0133**. This gives multiple comparison test for differences between rows.



Ward clustering

- The type of clustering performed by this procedure of "minimizing the reduction of inertia at each step" is called Ward clustering (see our earlier classes on cluster analysis)
- Ward clustering is a hierarchical clustering analysis which needs:

(a) description vectors of objects to be clustered

(b) weights for each object

If you prefer to have a "distance" criterion for clustering, this is it:

Masses of clusters
$$G_1$$
 and G_2

 $d(G_1, G_2) = \frac{r_1 r_2}{r_1 + r_2} \| prof_1 - prof_2 \|_c^2$

Chi-square distance Profiles of clusters

 G_1 and G_2

- We want to perform Ward clustering on the profiles, with weights equal to the masses.
- Since Ward clustering calculates Euclidean distances between vectors, we would need to prepare the profiles so that the Euclidean distances will be chi-squared: that is, we have to divide the profile elements by the square roots of their average (expected) values.
- But we need to weight the points: use XLSTAT or Fionn Murtagh's R code: http://astro.u-strasbg.fr/~fmurtagh/mda-sw/correspondances

Bíplot

Correspondence analysis is based on the SVD of

$$\mathbf{D}_r^{-1/2}(\mathbf{P} - \mathbf{r}\mathbf{c}^{\mathsf{T}})\mathbf{D}_c^{-1/2} = \mathbf{U}\mathbf{D}_{\alpha}\mathbf{V}^{\mathsf{T}}$$

 $\begin{array}{ll} \mbox{Principal} & \mbox{\bf F} = \mbox{\bf D}_r^{-1/2} \mbox{\bf U} \mbox{\bf D}_\alpha & \mbox{Standard} & \mbox{\bf \Phi} = \mbox{\bf D}_r^{-1/2} \mbox{\bf U} \\ \mbox{coordinates} & \mbox{\bf G} = \mbox{\bf D}_c^{-1/2} \mbox{\bf V} \mbox{\bf D}_\alpha & \mbox{coordinates} & \mbox{\bf \Gamma} = \mbox{\bf D}_c^{-1/2} \mbox{\bf V} \end{array}$

We want the right hand side in the form of scalar products between the coordinate matrices

$$\mathbf{D}_r^{-1}(\mathbf{P} - \mathbf{r}\mathbf{c}^{\mathsf{T}})\mathbf{D}_c^{-1} = \mathbf{D}_r^{-1/2}\mathbf{U}\mathbf{D}_{\alpha}(\mathbf{D}_c^{-1/2}\mathbf{V})^{\mathsf{T}} = \mathbf{F}\mathbf{\Gamma}^{\mathsf{T}}$$

 $\left(\frac{p_{ij}}{r_i} - c_j\right) / c_j \approx (f_{i1}\gamma_{j1} + f_{i2}\gamma_{j2})$

In full space: $\frac{p_{ij} - r_i c_j}{r_i c_j} = f_{i1} \gamma_{j1} + f_{i2} \gamma_{j2} + \cdots$

In reduced space: $\frac{p_{ij} - r_i c_j}{r_i c_j} \approx f_{i1} \gamma_{j1} + f_{i2} \gamma_{j2}$

scalar product between row profile and column vertex

row profile element average profile element

Biplot variations by Gabriel & Greenacre



• Gabriel's modification:

 $\frac{p_{ij}}{r_i}$

deviation of profile from average

Greenacre's modification (the standard biplot):



standardized deviation of profile from average

• (Relative) column contribution:

 $c_{i}g_{ik}^{2}/\lambda_{k} = c_{i}(\lambda_{k}^{1/2}\gamma_{ik})^{2}/\lambda_{k} = c_{i}\gamma_{ik}^{2}$

scalar product between row profile and column vertex

vertices (standard coordinates) "shrunk by their respective masses

vertices shrunk by square roots of their respective masses; squares of these rescaled column coordinates are exactly the (relative) contributions of the column to the respective dimension