CPCA: A program for principal component analysis with external information on subjects and variables

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A program is described for principal component analysis with external information on subjects and variables. This method is called *constrained principal component analysis* (CPCA), in which regression analysis and principal component analysis are combined into a unified framework that allows a full exploration of data structures both within and outside known information on subjects and variables. Many existing methods are special cases of CPCA, and the program can be used for multivariate multiple regression, redundancy analysis, double redundancy analysis, dual scaling with external criteria, vector preference models, and GMANOVA (growth curve models).

Regression analysis and principal component analysis (PCA) typically proceed from opposite ends of the data analysis spectrum. Regression methods are usually considered explanatory techniques in which the variability in a set of dependent variables is partitioned into two orthogonal components: variability accounted for by known external information (e.g., one or more predictor variables), and variability that is independent of external information (i.e., error variability). PCA, on the other hand, is an exploratory technique used to investigate structures in data when no external information is available. In this paper, we describe a program for constrained principal component analysis (CPCA), in which features of regression analysis and PCA are combined into a unified framework that captures advantages of both.

Takane and Shibayama (1991) have described in detail the rationale and methodology underlying CPCA. Briefly, the analysis can be conceptualized as proceeding in two distinct stages: (1) an external (regression) analysis in which a data set is partitioned into predictable and error variation, and (2) an internal analysis in which PCA is applied to the results of the external analysis. The external analysis decomposes an N subject $\times n$ variable matrix of dependent variables \mathbb{Z} , into variation predictable from external information on subjects and variables and variation unrelated to the external information. Assuming the information on subjects (e.g., age, gender, IQ, treatment

matrix for pair comparisons, characteristics such as order, familiarity, and complexity of stimuli presented in a memory task, etc.) into an $n \times q$ ($\ll n$) matrix, H, the model can be written as

$$Z = GMH' + BH' + GC + E,$$
 (1)

where \mathbf{M} $(p \times q)$, \mathbf{B} $(N \times q)$, and \mathbf{C} $(p \times n)$ are matrices of coefficients to be estimated, and \mathbf{E} $(N \times n)$ is a matrix of error components. The four terms on the right side of the equation explain unique (orthogonal) portions of the variation in \mathbf{Z} . The first term pertains to variation that can be explained uniquely by information on subjects and variables combined, the second term by information on variables alone, and the third term by information on subjects alone. The fourth term includes variation in \mathbf{Z} that is independent of information on subjects and variables.

At this stage, the analysis is related to multivariate multiple regression. In fact, if only the main data Z, and information on subjects, G, were available, then only the third and fourth terms in Equation 1 would be estimated. and the parameters in C would equal the regression coefficients obtained from a standard multivariate multiple regression analysis (assuming both G and Z contained continuous variables). Thus, the Stage 1 full model can be viewed as a doubly multivariate multiple regression analysis with external information not only on subjects but also on variables. Of course, the full model may not always be desired, or other submodels may be of substantive interest along with the full model. In such cases, it is possible to obtain Z = BH' + E (by simply omitting G from the analysis), or Z = GC + E (by omitting H from the analysis).

The second stage, the internal analysis, applies PCA to each of the decomposed submatrices from the Stage 1

condition, etc.) is coded into an $N \times p$ ($\ll N$) matrix, G,

and the information on variables (e.g., occasions, design

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Table 1 Summary of Analyses

Number	Data	External Analysis	Internal Analysis
1	Z	none	PCA of Z
2	G,Z	Regress Z on G to obtain _G Z'	PCA of _G Z'
3	G,Z	Regress Z on G to obtain _G Z'	PCA of $Z - {}_{G}Z'$
4	Z,H	Regress Z on H to obtain Z _u	PCA of Z' _H
5	Z,H	Regress Z on H to obtain Z' _H	PCA of $\mathbf{Z} - \mathbf{Z}'_{\mathbf{H}}$
6	G,Z,H	Regress Z on G and and H to obtain GZ'H	PCA of _G Z' _H
7	G,Z,H	Regress Z on H to obtain Z' _H and on G and H to obtain _G Z' _H	PCA of $\mathbf{Z_{H}}' - {}_{\mathbf{G}}\mathbf{Z'_{H}}$
8	G,Z,H	Regress Z on G to obtain _G Z' and on G and H to obtain _G Z' _H	PCA of $_{\mathbf{G}}\mathbf{Z}'{\mathbf{G}}\mathbf{Z}'_{\mathbf{H}}$
9	G,Z,H	Regress to obtain $_{G}Z^{\prime},Z_{H}^{\prime}$ and $_{G}Z_{H}^{\prime}$	PCA of $\mathbf{Z} - {}_{\mathbf{G}}\mathbf{Z}' - \mathbf{Z}'_{\mathbf{H}} - {}_{\mathbf{G}}\mathbf{Z}'_{\mathbf{H}}$

analysis. Alternatively, submatrices can be recombined for PCA if desired. For example, the first and second terms can be combined, amounting to PCA of that part of Z that is predictable overall from external information on variables (including that which is unique to H and that part which H shares with G). Similarly, the first and third terms can be combined, and the part of Z that is predictable from G, both uniquely and in common with H, can be component analyzed.

In all, nine analyses are possible allowing a full exploration of main data structures both inside and outside the known external information (see Table 1, and the Example Analyses section). The nine analyses include: (1) PCA of Z unconstrained by G or H. This is a straightforward principal component analysis of Z. (2) PCA of Z constrained by G and (3) PCA of Z independent of G. Analysis 2 is a component analysis of the part of Z that is predictable overall from subject information, whereas Analysis 3 component analyzes the part of Z from which subject information has been partialed. In these analyses, external information on variables is ignored, which amounts to combining the first and third terms from the full model presented above (obviously, these would be the analyses of choice if no stimulus information were available). Notice, however, that Analysis 3 is not an analysis of E from the full model. The error component will differ depending on whether H is included or ignored. (4) PCA of Z constrained by H, and (5) PCA of Z independent of H. These analyses are equivalent to Analyses 2 and 3 above, except that now information on variables is incorporated and subject constraints are ignored (or are unavailable). (6-9) PCA of the four terms in the full model.

RELATIONS TO OTHER METHODS

CPCA is a very general method for finding a reduced rank representation of the relationship among sets of

variables, and it subsumes several existing methods as special cases. For example, if only G and Z were available and both contained continuous variables, CPCA would reduce to what has been called principal components of instrumental variables (Rao, 1964), reduced rank regression (Anderson, 1951), and redundancy analysis (Van den Wollenberg, 1977). The unique aspect of CPCA is that it extends redundancy analysis to incorporate external information on variables and on subjects, suggesting as a possible synonym for CPCA the term double redundancy analysis (or if only Z and H are available, we refer to the analysis as redundancy analysis of structured variables). Additionally, unlike standard redundancy analysis, CPCA focuses not only on structuring the predictable parts of Z but also on structuring

Table 2 Analysis of Main Data Matrix (Z)

	A	nalysis of Main	Data Matrix (Z)
	LYSIS 1: Ana		_	
		external analysis	•	
	484.000 %S			
SS &	%SS for the	internal analysis	•	
SS=	356.148 %SS	(per term SS) =	73.584 %SS (per t	total SS)= 73.584
I	Dimensionwis	e SS & %SS	1	2
		SS=	253.493	102.654
	%SS (per	term SS)	52.375	21.210
	%SS (per	total SS)	52.375	21.210
Matr	ix A of Comp	onent Loadings	•	
ı	.761	464		
2	.763	.357		
3	592	.573		
4	903	.335		
5	.810	.321		
6	678	487		
7	574	549		
8	.883	.246		
9	450	619		
10	.638	378		
11	774	.566		

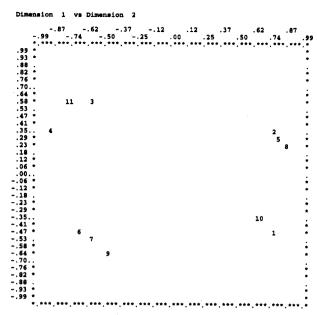


Figure 1. Plot of loading matrix for Analysis 1.

residuals in **Z** after variation accounted for by external information has been removed.

Other special cases of CPCA include growth curve models (GMANOVA; see Khatri, 1966; Rao, 1965) and two-way CANDELINC (Carroll, Pruzansky, & Kruskal, 1980), which both analyze only the first term in Decomposition 1. If **Z** is a matrix of pairwise preference data, **G** is a vector of ones, and **H** is a design matrix for paired comparison, then three vector models for pairwise preference data (e.g., De Soete & Carroll, 1983; Heiser & de Leeuw,1981; Takane and Shibayama,1988) are all special cases of CPCA that differ according to which term in Decomposition 1 is analyzed. (See Takane & Shibayama, 1991, for details and for a discussion of how CPCA relates to yet other methods. Example applications are also included.)

A commonly used method for investigating the reduced rank relationship between two sets of variables (e.g., Z and G) is canonical correlation. It obtains linear combinations (components) of each set of variables under the constraint that corresponding pairs of components are maximally correlated. The problem with canonical correlation is that components that maximally correlate might account for only a small portion of the variation in their respective sets. As a result, a component in one set (e.g., G) might correlate highly with a component in a second set (e.g., Z) but might explain only a small portion of the overall variation in Set Z (and vice versa). This overall portion of variation in one set that is explained by a component of another set is called redundancy, and it is this quantity that CPCA maximizes(for a full description of the difference between canonical correlation and redundancy analysis, see Van den Wollenberg, 1977). CPCA also differs from canonical correlation by accommodating information on subjects and on variables. A method that does incorporate both subject and variable constraints is GENFOLD2 (DeSarbo & Rao, 1984). However, GENFOLD2 does constrained unfolding analysis by fitting a distance model using an iterative gradient procedure, whereas CPCA does constrained principal component analysis by fitting a bilinear model.

THE CPCA PROGRAM

Program Description

CPCA is written in Fortran for the IBM PC XT, AT, PS/2, or compatibles under MS-DOS/PC-DOS 3.0 or higher. A math coprocessor is supported. The program can analyze problems with up to 1,000 subjects and 40 variables in each of Z, G, and H. At this point in development, the program does not accommodate missing data. At least 550 K of available memory is required. The program is very efficient. For example, using a 486DX2/66

Table 3
Analysis of Z Regressed on G (the Subjects Information Matrix)

ANALYSIS 2: Analysis of P(G)Z ● SS & %SS for the external analysis ● SS= 346.747 %SS= 71.642

Matr	ix C transpose	ed (in Model Z	= GC + E) •	
1	.422	-1.267	1.202	357
2	1.484	399	152	933
3	608	1.145	652	.115
4	984	1.293	895	.586
5	1.466	538	014	914
6	712	157	498	1.367
7	667	125	306	1.098
8	1.355	547	.294	-1.102
9	600	258	039	.898
10	068	731	1.478	679
11	- 735	1 451	- 903	177

SS & %SS for the internal analysis 10

SS= 319.763 %SS (per term SS)= 92.218 %SS (per total SS)= 66.067

Dimensionwise SS &	k %SS	1	2
SS	S=	243.72 7	76.036
%SS (per term S	SS)	70.29 0	21.929
%SS (per total S	SS)	50 .357	15.710

Matrix A of Component Loadings 6

1	.764	500
2	.741	.456
3	619	.344
4	933	.257
5	.792	.373
6	613	394
7	502	360
8	.870	.313
9	346	40
10	.621	440
11	796	453

Matrix of correlations between G and F

1	.686	.480
2	567	.667
3	.458	591
4	576	555

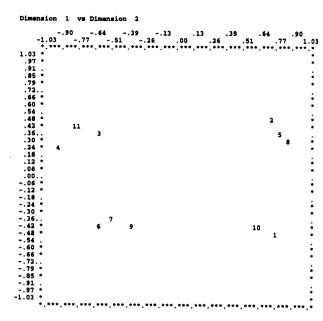


Figure 2. Plot of loading matrix for Analysis 2.

computer, an analysis of 200 subjects with 11 variables in **Z**, 4 variables in **G**, and 3 variables in **H** required 24 sec of CPU time to perform all nine analyses.

Input

Input to the program consists of job parameter information on number of subjects, number of variables in \mathbb{Z} , number of variables in \mathbb{G} , number of variables in \mathbb{H} , data transformations (raw data, column centered, or standardized), number of components, input/output options, and input data format. This information along with the data for N subjects (both \mathbb{Z} and \mathbb{G} data), a design matrix for \mathbb{H} , and optional target matrices for procrustes rotations are all placed in an input file using any standard word-processing program or editor. The program is run by typing cpca-t followed by an input filename and an optional output filename. The syntax is

cpca-t infile.ext outfile.ext

where infile.ext is the file containing job parameter information and the data, and outfile.ext is an optional file that receives the results of the analysis.

Output

For Analysis 1, the program output includes total sums of squares (SS) for the main data, SS and percent sums of squares (%SS) for each of the components retained, component loadings, and component scores. Loadings and scores can be plotted for each pair of components, and, to aid interpretation, optional varimax, promax, and procrustes rotations are available. When rotations are requested, SS and %SS of squares for rotated components are printed, and, in the cases of promax rotation, correlations among components are given.

For all analyses in which the main data are constrained by external information (e.g., Analyses 2, 4, 6, 7, and 8 above), the program prints out (1) total SS for the main data, (2) SS and %SS accounted for by the external information, (3) matrices of parameter estimates (i.e., matrix M, B, or C), (4) SS and %SS for each of the components retained in the internal analysis, (5) loadings of the main variables on components constrained by external information, and scores on those components (optional plots are available), (6) weights and loadings of the external variables on the externally defined components (e.g., these would be redundancy variate weights and loadings for the case in which only G and Z are analyzed), and (7) all of the above for optional varimax, promax, and procrustes rotations, as well as correlations among factors for promax rotated components.

For all analyses of error terms (e.g., Analyses 3, 5, and 9), the program printout includes total SS for the main data, SS and %SS for the part of the main data that is independent of the external information, SS and %SS for each of the components retained in the internal analysis, and component loadings and scores. Again, optional plots and rotations are available.

Availability

A copy of the program and the program user's manual may be obtained from the first author or can be downloaded from http://castle.uvic.ca/psyc/files.html

EXAMPLE CPCA ANALYSES

This example analyzes scores on four groups, each containing 11 psychiatric patients (G1 = manic-depressive [depressed]; G2 = manic-depressive [manic]; G3 = simple schizophrenia; G4 = paranoid schizophrenia) on 11 psychopathological items from the Brief Psychiatric Rat-

Table 4

Analysis of Residuals After Regressing Z on C

	Analysis	of Residuals A	fter R egress ing	Z on G
		alysis of Q(G)Z external analysis		
	137.253 %S		•	
			_	
22.0	7033 for the	internal analysis		
33-	13.001 7033	(per term 35) = 3	3.832 %SS (per t	total SS)= 15.266
	Dimensionwis	e SS & %SS	1	2
		SS=	45.523	28.364
	%SS (per	term SS)	33.167	20.665
	%SS (per	total SS)	9.406	5.860
Matr	ix A of Comp	onent Loadings (
1	.761	464		
1	.139	026		
2	.035	099		
2 3 4	.432	.403		
4	.054	063		
5	.119	.117		
6	317	.140		
7	572	354		
8	.067	059		
9	594	.551		
10	.089	.006		
11	.132	2.035		

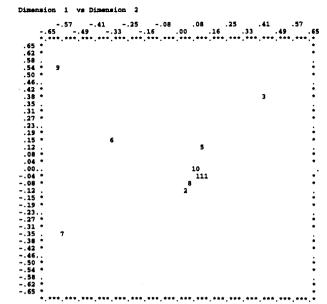


Figure 3. Plot of loading matrix for Analysis 3.

ing Scale. The 11 items are (1) emotional withdrawal. (2) guilt feelings, (3) tension, (4) grandiosity, (5) depressive mood, (6) suspiciousness, (7) hallucinatory behavior, (8) motor retardation, (9) unusual thought content, (10) blunted affect, and (11) excitement. For purposes of the analyses presented below, Z is a 44 × 11 matrix containing scores on the 11 items for all 44 patients, G is a 44×4 matrix of dummy coded variables indicating the four patient-type groups, and H is an 11×4 matrix of dummy coded variables indicating item-type classifications. For this example, we classified together items that might be expected to be symptomatic of depression (guilt, depressive mood, and motor retardation), manic state (tension, grandiosity, and excitement), simple schizophrenia (emotional withdrawal and blunted affect), and paranoid schizophrenia (suspiciousness, hallucinatory behavior, and unusual thought content). In what follows, we show the results of each of the nine analyses provided by the CPCA program, along with explanatory comments. Note that optional output, such as printed factor scores and graphs of factor scores, and the results of various rotations are excluded due to space constraints.

Table 2 and Figure 1 show the output for Analysis 1. The highlighted numbers below correspond to those shown in Table 2.

- Analysis 1 is a straightforward principal components analysis of **Z**.
- The SS and %SS are given for the external (or regression) analysis. In Analysis 1, there are no external constraints on Z; therefore, in this case, the values of 484 and 100% reflect the total variability in Z.
- Summary results are given for the internal (or PCA) analysis. The SS term reflects the amount of variability accounted for by the number of components retained. In

Table 5
Analysis of Z Regressed on H (the Variables Information Matrix)
ANALYSIS 4: Analysis of ZP(H)

SS & %SS for the external analysis

■

SS = 400.371 %SS = 82.721					
Matr	ix B in Model	Z = BH (transp	oose) + E 🛮		
1	038	1.629	464	883	
2	.638	. 1.465	-1.028	-1.341	
3	279	1.163	-1.039	.252	
4	714	1.722	574	231	
5	123	1.350	-1.109	046	
6	.273	1.683	−.74 1	909	
7	026	1.381	332	523	
8	.178	1.663	9 19	-1.174	
9	.604	1.502	838	445	
10	.705	.996	56 5	759	
11	.730	1.227	922	-1.198	
12	-1.036	480	1.171	.336	
13	-1.006	−.538	1.093	.807	
14	897	552	1.486	176	
15	822	332	1.767	206	
16	780	424	1.273	-1.228	
17	-1.174	412	1.257	529	
18	-1.006	343	1. 36 3	897	
19	897	379	1.773	704	
20	-1.006	671	.915	.966	
21	-1.187	-1.093	.884	.268	
22	-1.174	214	1.279	617	
23	1.584	292	558	588	
24	.990	.607	910	363	
25	1.259	349	-1.180	.032	
26	1.888	350	767	619	
27	.865	.145	-1.071	.442	
28	1.424	063	-1.139	301	
29	1.317	.487	072	-1.414	
30	1.670	.081	956	028	
31	1.024	305	920	.605	
32	1.482	.445	834	730	
33	1.233	.064	539	129	
34	285	995	111	1.552	
35	945	809	.443	1.070	
36	621	-1.066	.099	1.020	
37	678	459	304	1.620	
38	396	971	.658	.810	
39	519	-1.071	.007	1.330	
40	860	942	.902	.655	
41	480	-1.022	.389	.737	
42	234	-1.369	.497	1.084	
43	956	780	.219	1.342	
44	.277	-1.331	.420	1.111	
SS & %SS for the internal analysis •					

SS & %SS for the internal analysis \bullet SS = 350.107 %SS (per term SS)= 87.446 %SS (per total SS)= 72.336

Dimensionwise SS & %SS	1	2
SS=	249.789	100.318
%SS (per term SS)	62.389	25.056
%SS (per total SS)	51.609	20.727

Matrix A of Component Loadings

-.426

.696

2	.819	.313
3	760	.488
4	760	.488
5	.819	.313
6	566	550
7	566	550
8	.819	.313
9	566	550
0	.696	426
1	760	.488

Matrix of correlations between H and A

.708	.434
638	.668
.464	432
- 473	- 728

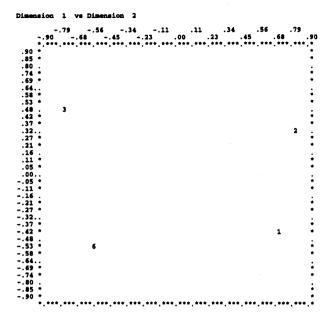


Figure 4. Plot of loading matrix for Analysis 4.

this example, the amount of variability accounted for by the first two principal components was 356.148. The %SS (per term SS) reexpresses this variability into a percentage of the variance in Z that is related to the external constraints. In contrast, the %SS (per total SS) reflects the percentage of total variability accounted for by the number of components retained. In Analysis 1, there are no external constraints; therefore, these two values are the same. The example results show that the first two principal components account for 73.584% of the total variation in Z. The dimensionwise SS and %SS give the same information, but for each component considered separately. For example, the current analysis indicates that the amount of variability accounted for by the first component is 253.493, which accounts for 52.375% of the total variance in **Z**.

• Matrix A contains the (unrotated) loadings of the variables in Z on the first two principal components, and the loadings are plotted in Figure 1 (if more than two components are retained, pairwise plots are produced). The plot indicates that the relations among the items are fairly clear: Items characteristic of depression (Items 2, 5 and 8) are located in the third quadrant, those characteristic of manic state (Items 3, 4, and 11) are in the fourth quadrant, simple schizophrenic symptoms (Items 1 and 10) are in the second quadrant, and those related to paranoid schizophrenia (Items 6,7, and 9) are in the first quadrant.

The results for Analysis 2 are shown in Table 3 and Figure 2. The highlighted numbers below correspond to those shown in Table 3.

• Analysis 2 is an analysis of the part of **Z** that is predictable from **G**—that is, it is an analysis of $_{\mathbf{G}}\mathbf{Z}'$, where $_{\mathbf{G}}\mathbf{Z}' = \mathbf{GC}$, and is equivalent to redundancy analysis. In

the current example, it is an analysis of between-groups variation.

● The values in this section show that the betweengroups variation is 346.747, which accounts for 71.642% of the total variation. This total percent of variation in Z that can be accounted for by G is called the total redundancy.

Matrix C is a matrix of parameter estimates for predicting Z from G. In general, this matrix contains the regression coefficients that would be produced via a multivariate multiple regression of Z on G. In the current example, G is a matrix of dummy variables coding group membership, so the values in matrix C are the means for each group on each of the 11 variables.

This section indicates that when PCA is applied to the between-groups variation, $_{G}Z'$, the variation accounted for by a two-component solution is 319.76, which represents 92.218% of the between-groups variation and 66.067% of the total variation. In the language of redundancy analysis, we would say that the first two predictor (redundancy) variates account for 66.067% of the overall variance in Z, which represents 92.218% of the total redundancy. The dimensionwise SS and %SS give corresponding values separately for the first and second components. Thus, the first two redundancy variates account for 50.357% and 15.710%, respectively, of the total variance in Z.

Matrix A shows the loading matrix that arises when PCA is applied to $_{\rm G}{\bf Z}'$. These loadings are plotted in Figure 2 and are strikingly similar to the ones presented earlier for the unconstrained solution. Overall, these results suggest that most of the variation in ${\bf Z}$ (and the factor structure arising out of it), is due to differences among patient groups. In redundancy analysis, Matrix A would be referred to as the matrix of cross-loading of the criterion variables (${\bf Z}$) on the predictor (redundancy) variates.

Table 6
Analysis of Residuals After Regressing Z on H

6SS for the 6 3.629 %SS= 6SS for the i	external analysis = 17.279 nternal analysis	₽	total SS)= 9.911
mensionwis	e SS & %SS	1	2
%SS (per	SS= term SS)	28.680 34.294 5.926	19.288 23.064 3.985
A of Compo	onent Loadings C	•	
.016 .119 372 .218 124 001 .454 .005 453 016	226 057 017 068 083 443 .219 .141 .224 .226		
	%SS for the e 3.629 %SS= 6SS for the i 7.968 %SS (mensionwise %SS (per the second of t	6SS for the external analysis of 3.629 %SS = 17.279 6SS for the internal analysis of 7.968 %SS (per term SS) = 5 mensionwise SS & %SS	6SS for the internal analysis ● 7.968 %SS (per term SS) = 57.358 %SS (per term SS) = 28.680 SS = 28.680 %SS (per term SS) 34.294 %SS (per total SS) 5.926 A of Component Loadings ● .016226 .119057372017 .218068124083001443 .454 .219 .005 .141453 .224016 .226

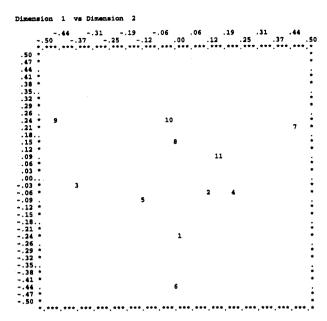


Figure 5. Plot of loading matrix for Analysis 5.

♠ The matrix of correlations between G and F is a loading matrix that reflects the structure of the predictor (redundancy) variates. The sign and size of the loadings in the current example indicate that the difference between Groups 1 and 3 (depressed and simple schizophrenic patients) and Groups 2 and 4 (manic and paranoid schizophrenics) accounts for most of the variability in Z (50.357% as seen above). The second redundancy variate, accounting for 15.71% of the variance in Z contrasts depressed and manic patients with the two groups of schizophrenics.

The results for Analysis 3 are shown in Table 4 and Figure 3. The highlighted numbers below correspond to those in Table 4.

- $lack \Phi$ Analysis 3 is an analysis of the part of $\bf Z$ that is not predictable from $\bf G$ —that is, it is an analysis of $({\bf Z}-{}_{\bf G}{\bf Z}')$, where ${}_{\bf G}{\bf Z}'={\bf G}{\bf C}$. In the current example, it is an analysis of within-groups variation.
- The values in this section show that the withingroups variation is 137.253, which represents 28.358% of the total variation.
- This section indicates that when PCA is applied to the within-groups variation (**Z** − **GZ**'), the variance accounted for by a two-component solution is 73.887, which represents 53.832% of the within-groups variation but only 15.266% of the total variation. The dimensionwise SS and %SS give corresponding values separately for the first and second within-groups components.
- The loading matrix given in this section and illustrated in Figure 3 shows that the within-groups structure differs substantially from the total structure.

The results for Analysis 4 are shown in Table 5 and Figure 4. The highlighted numbers below correspond to those shown in Table 5.

- $oldsymbol{\Phi}$ Analysis 4 is an analysis of the part of Z that is predictable from H—that is, it is an analysis of Z'_H , where $Z'_H = BH'$. We refer to this analysis as redundancy analysis of structured variables. In the current example, it is an analysis of between-item-type variation, in which items are classified into types as described above.
- The values in this section show that the betweenitem-type variation is 400.371, which represents 82.721% of the total variation in Z. Again, this percent represents redundancy in Z, but with respect to item types rather than patient types.
- **oldsymbol{\Theta}** Matrix $oldsymbol{B}$ is a matrix of parameter estimates for predicting $oldsymbol{Z}$ from $oldsymbol{H}$. In general, this matrix contains the regression coefficients that would be produced if $oldsymbol{Z}$ were transposed to produce a p (variables) \times N (subjects) matrix, which was then subjected to a multivariate multiple regression on $oldsymbol{H}$. In the current example, $oldsymbol{H}$ is a matrix of dummy variables coding item types; the values in matrix $oldsymbol{B}$ are the means for each subject on each of the four types of items. For example, items classified as characteristic of depression include guilt, depressive mood, and motor

Table 7

Analysis of Z Regressed on G and H

Analysis of Z Regressed on G and H							
ANALYSIS 6: Analysis of P(G)ZP(H) ● SS & %SS for the external analysis ● SS = 335.453 %SS = 69.309							
Matri	ix M in the ful	ll model ❸					
1	.177	1.435	776	660			
2	999	494	1.296	180			
3	1.340	.043	813	281			
4	518	983	.293	1.121			
SS & %SS for the internal analysis ● SS = 313.775 %SS (per term SS) = 93.538 %SS (per total SS) = 64.830							
I	Dimensionwis	e SS & %SS	1	2			
		SS=	239.164	74.611			
	%SS (per	,	71.296	22.242			
	%SS (per	total SS)	49.414	15.416			
Matr	ix A of Compe	onent Loadings	•				
1	.691	472					
2	.802	.381					
3	783	.351					
4	783	.351					
5	.802	.381					
6	486	385					
7	486	385					
8	.802	.381					
9	486	385					
10	.691	472					
11	783	.351					
Matrix of correlations between H and A •							
1	.701	.580					
2	681	.534					
3	.464	578					
4	4422614						
Matrix of correlations between G and F ●							
1	.689	.482					
2	569	.665					
3	.454	596					

-.574

-.551

77553311 .11 .33 .55 .7788664422 .00 .22 .44 .66 .8188 *.83 *88 *66 *11 .33 .55 .7762 .5762 .5762 .5762 .5762 .5762 .5763 .57	Dimens	ion 1	vs Dir	ension	2							
.88 * .83 * .787877 * .67 * .67 * .6257 * .52 * .4742 * .36 * 1136 * .31 * .30 * .30 * .31 * .30 * .31 * .32 * .33 * .34 * .35 * .35 * .36 * .31 * .37 * .38 * .		7	!	is	33	11	.11	.3	3	. 55	.77	
88 * 83 * 83 * 84 * 84 * 84 * 84 * 84 *	-	. 88	66	44	22		0	. 22	. 44		6	. 86
.83 *		. * * * . * '	**.***.	***.***	••••	•.•••.					,	٠.:
.78												
.73 * .67 * .67 * .6257 * .52 * .4742 * .36 * 1136 * .3130 * .30 * .31 * .40 * .31 * .41 * .42 * .43 * .44 * .44 * .44 * .44 * .44 * .44 * .4748 * .4748 * .48 * .48 * .49 * .40 * .40 * .40 * .41 * .42 * .44 * .44 * .44 * .44 * .4748 * .48 * .48 * .49 * .40 * .40 * .40 * .41 * .42 * .44 * .44 * .4748 * .48 * .48 * .49 * .40 * .40 * .40 * .40 * .41 * .41 * .42 * .43 * .44 * .44 * .44 * .45 * .4748 * .48 * .48 * .49 * .49 * .40 * .												
.62												*
.57 *												•
52 * 47 .												:
47												
42 *												
31												•
26 *		11									8	•
.21 *												:
16												:
.10 *												
.00051011162126313631364247525257626773788383 -	.10 *											•
05 *10 *101121 *25 *3136 * 942 *4752 *57 *6267 *73 *7883 *												•
10 *1521 *22 *3136 *3136 *42 *4752 *52 *57 *6267 *73 *7883 *												:
16												
21 *												
31	21 *											•
36 * 9 *												•
42 *												:
- 47				•								*
52 *										:	10	
- 62 - 67 * - 73 * - 78 . - 78 .												•
67 *73 *78												•
73 *78						,						:
78 . 83 *	73 *											
	78											
88 *												
	88 *											

Figure 6. Plot of loading matrix for Analysis 6.

retardation; the parameters in **B** associated with this classification would equal the subjects' means on these three items.

- This section indicates that when PCA is applied to the between-item-type variation, the variance accounted for by a two-component solution is 350.107, which represents 87.446% of the between-item-type variation and 72.336% of the total variation (i.e., the item-type redundancy is 72.336). The dimensionwise SS and %SS indicate that the first two item-type redundancy variates account for 51.609% and 20.727%, respectively, of the total variance in Z.
- Matrix A shows the loading matrix that arises when PCA is applied to the Z'_H obtained from $Z'_H = BH'$. These loadings are plotted in Figure 4. These results show that dummy coding the items into types has the effect of constraining the loadings for items within a type to be equal. Moreover, the figure clearly indicates that these constraints capture the essence of the structure of Z (only one item from each item type appears in the figure because same-type items overlap).
- ♠ The matrix of correlations between H and A indicates that the first item-type redundancy variate is defined by the difference between Item Types 1 and 3 (depressed and simple schizophrenic) and Item Types 2 and 4 (manic and paranoid schizophrenic). The second item-type redundancy variate contrasts the combined manic and depressive item types with item types characteristic of schizophrenia (simple and paranoid item types combined).

The results of Analysis 5 are shown in Table 6 and Figure 5. The highlighted numbers below correspond to those shown in Table 6.

O Analysis 5 is an analysis of the part of **Z** that is not predictable from **H**—that is, it is an analysis of $(\mathbf{Z} - \mathbf{Z}'_{\mathbf{H}})$,

Table 8

Analysis of the Part of Z That Is Predictable From H

But Unrelated to G

But Unrelated to G						
		ysis of Q(G)ZP				
SS & %SS for the external analysis SS = 64.917 %SS = 13.413						
	x B in the full		•••	202		
1	215	.194	.311	223 - 691		
2	.461 456	.031 272	253 263	681 .912		
4	43 6 891	.287	203 .202	.428		
5	300	085	3 34	.614		
6	.096	.249	.035	249		
7	203	053	.443	.136		
8	.001	.229	144	514		
9	.427	.067	062	.214		
10	.528	438	.211	099 539		
11 12	.553	208 .014	146 126	538 .516		
13	038 008	044	20 4	.987		
14	.102	057	.189	.004		
15	.177	.162	.471	026		
16	.219	.071	023	-1.048		
17	175	.083	040	349		
18	008	.152	.067	717		
19	.102	.115	.477	524		
20	008	177	381	1.146		
21 22	188 175	599 .280	413 018	.448 437		
23	.244	334	01 6 .256	307		
24	349	.565	0 9 6	082		
25	081	392	366	.313		
26	.548	393	.046	337		
27	474	.102	258	.723		
28	.084	106	326	019		
29	022	.445	.741	-1.133		
30	.330	.038	143	.254		
31	316	348	107	.887		
32 33	.143 106	.403 .021	020 .274	449 .152		
34	.233	011	404	.431		
35	427	.174	.150	051		
36	103	083	194	101		
37	160	.525	597	.499		
38	.122	.012	.365	311		
39	001	088	285	.209		
40 41	342 .038	.041 039	.610 .097	466 384		
42	.284	386	.205	037		
43	438	.203	074	.221		
44	.795	348	.127	010		
\$ 22		nternal analysis				
				r total SS)= 11.098		
		-	1	2		
ı	Dimensionwise					
	0/65 (no-	SS=	41.157 63.400	12.558 19.344		
	%SS (per %SS (per	total SS)	8.504	19.344 2.595		
N/	· -			2.373		
	•	nent Loadings	y			
1 2	.120	270				
3	.081 .187	.205 .066				
4	.187	.066				
5	.081	.205				
6	511	.015				
7	511	.015				
8	.081	.205				
9	511	.015				
10	.120	270				
П	.187	.066				
Matrix of correlations between H and A @						
1	.266	.680				
2	.492	.144				
3	.268	888				

-.991

-.055

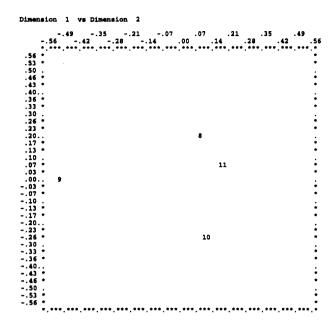


Figure 7. Plot of loading matrix for Analysis 7.

where $Z'_{H} = BH'$. In the current example, it is an analysis of within-item-type variation.

- The values in this section show that the variation within item types is 83.629, which is only 17.279% of the total variation in **Z**.
- $lacktriangled{\Theta}$ This section indicates that when PCA is applied to the within-item-type variation ($\mathbf{Z}-\mathbf{Z}_{\mathbf{H}}'$), the variance accounted for by a two-component solution is 47.968, which represents 57.358% of the within-item-type variation, but only 9.911% of the total variation. The dimensionwise SS and %SS give corresponding values separately for the first and second within-item-type components.
- The loading matrix given in this section and illustrated in Figure 5 shows that the within-item-type structure differs substantially from the total structure.

The results for Analysis 6 are shown in Table 7 and Figure 6. The highlighted numbers below correspond to those shown in Table 7.

- $oldsymbol{\Phi}$ Analysis 6 analyzes the part of Z that is predictable from both G and H. It is an analysis of ${}_{G}Z'_{H}$, where ${}_{G}Z'_{H} = GMH'$. We refer to this analysis as double redundancy analysis. In the current example, it is an analysis of between-groups variation in between-item-type variation (i.e., it evaluates whether the item-type profiles vary over groups).
- The values in this section show that combined, G and H account for 69.309% of the total variation in Z. This percent represents redundancy in Z as a function of patient-type variation in item types.
- ♠ Matrix M is a matrix of parameter estimates for predicting Z from G and H. In the current example, because both G and H are matrices of dummy variables, the values

in matrix M are the means for each of the four patient types on each of the four item types.

- This section indicates that when PCA is applied to the between-patient-type variation in item-type variation, the variance accounted for by a two-component solution is 313.775, which represents 93.538% of the explained variation and 64.83% of the total variation. The dimensionwise SS and %SS give corresponding values separately for each of the components.
- Matrix A shows the loading matrix that arises when PCA is applied to the $_{\mathbf{G}}\mathbf{Z}'_{\mathbf{H}}$ obtained from $_{\mathbf{G}}\mathbf{Z}'_{\mathbf{H}} = \mathbf{GMH}'$. These loadings are plotted in Figure 6. Again, these results show that the structure of the explained variation captures the essence of the total variation in \mathbf{Z} .
- ♠ and ♠ These sections show the matrix of correlations between H and A and between G and F, respectively. These matrices have have the same interpretation as redundancy variates, as described previously. In this case, they indicate that the first overall redundancy variate is defined by

Table 9

Analysis of the Part of Z That Is Predictable From G

But Unrelated to H

ANALYSIS 8: Analysis of P(G)ZQ(H) ●						
SS & %SS for the external analysis •						
full model						
8138	.161					
	.050					
1 .161	178					
3081	.294					
3 05 7	.069					
3217	.246					
5025	023					
2 . 252	119					
8 .242	223					
8 .138	161					
5 08 0	116					
SS & %SS for the internal analysis ● SS = 9.909 %SS (per term SS) = 87.750 %SS (per total SS) = 2.047						
SS 1	2					
5.96	2 3.947					
52.79						
1.23	2 .815					
dings 🌑						
1						
5						
4						
4						
2						
3						
Matrix of correlations between G and F •						
0						
5						
0						
	full model • 8					

.839

.075

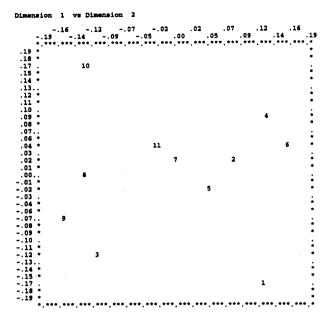


Figure 8. Plot of loading matrix for Analysis 8.

Table 10 Analysis of Residuals After Z Has Been Regressed on G and H

ANALYSIS 9: Analysis of O(G)ZO(H) 0 SS & %SS for the external analysis @ SS= 72.336 %SS= 14.946 SS & %SS for the internal analysis ® SS = 42.139 %SS (per term SS) = 58.255 %SS (per total SS) = 8.706 Dimensionwise SS & %SS 2 1 5.224 1.045 3.145

SS=	26.916	15
%SS (per term SS)	37.209	21
%SS (per total SS)	5.561	3
Matrix A of Component Loadings A		

Mati	rix A of Comp	onent Loadings 🛭	
1	.006	159	
2	.091	038	
3	329	010	
4	.171	039	
5	144	015	
6	098	429	
7	.498	.140	
8	.052	.053	
9	400	.289	
10	006	.159	
11	.159	.049	

the difference between patients who are depressed or simple schizophrenic and patients who are manic or paranoid schizophrenic in the item types hypothesized to be symptomatic of such patients. The second redundancy variate captures differences between schizophrenics overall and manic depressives overall in corresponding item

The results for Analysis 7 are shown in Table 8 and Figure 7. The highlighted numbers below correspond to those shown in Table 8.

• Analysis 7 is an analysis of the part of **Z** that is uniquely predictable from H, independent of G. It analyzes $(Z_H' - {}_GZ_H')$ and can be conceptualized as regressing Z on G, obtaining the residuals from that analysis, and then applying item-type constraints to those residuals. In the current example, it is an analysis of withinpatient-type variation in between-item-type variation.

2, **3**, **4**, **5**, and **6** These sections and Figure 7 report results identical in interpretation to similar results reported previously, with the proviso that they pertain to between-item-type variation after between-patient-type variation has been partialed out. Overall, the results indicate there is very little variation in Z that is uniquely attributable to H, and that the structure of this variation is quite different from the structure of the total variation. There is, however, some indication that, independent of group membership, items classified as characteristic of paranoid schizophrenics are responded to differently than other item types.

The results of Analysis 8 are shown in Table 9 and Figure 8. The highlighted numbers below correspond to those shown in Table 9.

• Analysis 8 is an analysis of the part of Z that is uniquely predictable from G, independent of H. It analyzes $(_{G}Z' - _{G}Z'_{H})$ and can be conceived as regressing Z (transposed) on H, obtaining the residuals from that analysis, and then applying subject constraints to those residuals. In the current example, it is an analysis of between-patienttype variation in within-item-type variation.

2, **3**, **4**, **5**, and **5** These sections and Figure 8 show results identical in interpretation to similar results reported previously, this time with the proviso that they pertain to between-patient-type variation after between-itemtype variation has been partialed out. The results indicate no independent effect of patient types.

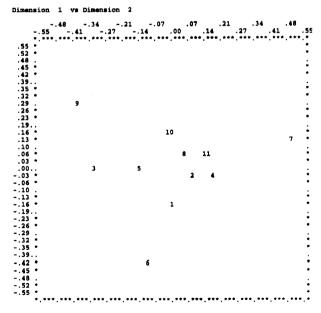


Figure 9. Plot of loading matrix for Analysis 9.

The results of Analysis 9 are shown in Table 10 and Figure 9. The highlighted numbers below correspond to those shown in Table 10.

- Analysis 9 is an analysis of the part of **Z** that is not predictable from **G** and **H**. In the current example, it is an analysis of within-patient-type variation in within-item-type variation.
- **②**, **③**, and **①** These sections and Figure 9 show that very little variation in **Z** remains that is independent of **G** and **H**, and that the structure of the overall residuals does not resemble the total structure at all.

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