

**TWINSPAN : a FORTRAN program for arranging multivariate data
in an ordered two-way table by classification of the individuals and
attributes / M.O. Hill**

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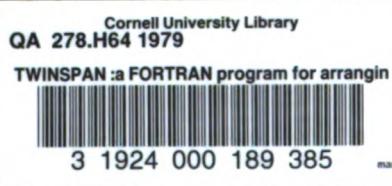
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TWINSPAN

A FORTRAN Program for Arranging Multivariate
Data in an Ordered Two-way Table by Classification
of the Individuals and Attributes

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Ithaca, New York 14850

July 1979

		11	1	
		21	9807631245	
9	SOL I SEM	211-3-----	0000	
5	PHRA COM	53415-----	0001	
3	IVA FRU	452113-----	0010	
11	SPAR PAT	3254-5-----	0011	
8	SCIR OLN	-1---53-----	01	
2	DIST SPI	-5414545-424	100	
1	ATRI PAT	231--3221411	101	
6	SAL I EUR	----22-23411	1100	
10	SPAR ALT	--142-135553	1101	
7	SAL I VIR	-----34	1110	
12	SUAE MAR	-----54	1110	
4	JUNC GER	-----151---	1111	
		000000111111		
		000001001111		
		00011 010011		
		00101 0101		
		01		

**TWINSPAN -- A FORTRAN program for arranging multivariate data
in an ordered two-way table by classification of
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July 1979

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PREFACE

TWINSPAN is program CEP-41 in the Cornell Ecology Programs series, written by Mark O. Hill. TWINSPAN is a FORTRAN program for two-way indicator species analysis, which is an improvement upon the original indicator species analysis (Journal of Ecology 63:597-613) in that species are classified as well as samples. This two-way classification makes possible a tabular matrix arrangement which approximates the result of Braun-Blanquet paperwork. TWINSPAN also constructs a key to the sample classification by identifying one to several species which are particularly diagnostic of each division in the classification.

TWINSPAN is a polythetic divisive method of classification; details of the algorithm are presented in this documentation. Although developed and discussed in a phytosociological context where the data matrix is one of species abundances in samples, the method is applicable to a wide range of data matrices expressing attributes of individuals (for example, vocabulary of persons, market preferences of customers, and types of artifact in archeological sites). TWINSPAN contains about 2300 cards.

The computer time required by TWINSPAN has a linear dependence on the amount of data, so large problems can be tackled readily. This is in marked contrast to most classification methods with requirements rising with the second or higher power of the number of samples. Furthermore, TWINSPAN does not store zero values of the data matrix (which usually predominate in large data matrices), and it does not produce or store a secondary matrix of sample similarities. Hence TWINSPAN is also very efficient with computer memory.

Copies of this documentation and magnetic tape copies of the program are available at cost. A catalog to the Cornell Ecology Programs series is available upon request. TWINSPAN is in the public domain and may be copied. TWINSPAN is offered with no warranty, express or implied, and with no legal liability or responsibility.

In many cases it is useful to parallel TWINSPAN classification with an ordination. In some cases if the data set is very large preliminary

clustering of essentially replicate samples may reduce the data set to manageable proportions, facilitating study of subsequent computer outputs. Other programs in the Cornell Ecology Programs series may be useful for these functions.

TWINSPAN requires that the input data matrix be supplied in a particular format which simplifies program code and permits fast data input. This format is essentially the condensed format in sample order of ORDIFLEX (an earlier program in the Cornell Ecology Programs series). However, in some cases this format may not be convenient for the initial keypunching of a data set. In order to allow greater flexibility in data matrix format, TWINSPAN is supplied with a utility program, CONDENSE, which reads data matrices in any format accepted by ORDIFLEX, and copies them into the restricted format used by TWINSPAN. CONDENSE is supplied without additional charge. This same format is also used by DECORANA, a Cornell Ecology Program by Mark O. Hill for detrended correspondence analysis and reciprocal averaging.

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July 1979

RRATA

Bugs in TWINSPLAN and DECORANA

To date, no bugs of note have appeared in DECORANA. Several bugs have appeared in TWINSPLAN, most of which will only appear when it is attempted to run large problems, where there are problems of dimensioning. Those that are known to me at present are as follows.

/ Lines 479,480 should read MM,MMAX, not M,MMAX. At present, if the allowed number of samples is exceeded, a bad error message is obtained. Changing MM to M puts the error message right.

/ Line 987 should read
IP(IP.LE.NMAX) INDPOT(IP)=JJJ
The point here is that it should not be possible to write outside the bounds of the array INDPOT. An error message will appear shortly, with an indication of the required dimensioning (cf. lines 1022-1026).

/ Line 552 should read
CALL PARA(MMIN,2,10000,5)
The point here is that groups smaller than 2 should not be attempted to be divided. The program gets into difficulties if it tries to do this.

/ Line 1246 should read
1004 FORMAT((1X,7(A4,1X,A4,I1,'(,A1,)',3X)))
i.e. the FORMAT statement should have an extra pair of brackets round the outside of it. If this is not put in, and if more than 7 indicators are asked for, the first letter of the 8th indicator is regarded as a control character and omitted from the print-out.

/ Lines 1076-1083 Common blocks are in wrong order; this leads to diagnostic messages being put out by some compilers, but has never to my knowledge been regarded as a fatal error.

/ Line 388 Numerous users have had trouble because the number read by the program as NITEM (-8 in diagram on p. 18) was wrongly positioned in the data. The awkward position of this number is a relic from days when the data were routinely submitted on cards. Users may think it worth their while to test that the value of NITEM lies between suitable limits (e.g. 4 to 20); this would eliminate almost all cases of wrongly submitted data. If users do want to test the value of NITEM, then they should insert a test statement immediately following line 388.

/ Fig. 1 (on page 2). The lines indicating the dichotomy are omitted. Do not be confused by this.

Restrictions not indicated in existing write-up are as follows

NMAX must exceed $(2^{**}(\text{LEVMAX}+1)-1)*3$ (-381 in default case). This will not cause trouble unless it is desired to use a large number for LEVMAX, i.e. maximum level of divisions - cf. p. 13.

Note: These four changes are already made on magnetic tape copies of TWINSPLAN made since

June 1981

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Bugs in TWINSPAN, continued

Dimensions of TOT, TOTJ must exceed
 $(2^{**}(\text{LEVMAX}+1)-1)$ (-127 in default case).
The dimensions supplied on line 35, of 511 ($-2^{**}9-1$)
allow 8 levels of division.

In addition to the restrictions given on p.33, an additional restriction is that the data when held in non-pseudospecies form must not occupy more than half the data matrix. If they do occupy more than this, the species classification may not be possible. This can be tested for following line 103 of the program, by demanding that
 2^*MDAT.LT.NDAT

If (as is usually the case) it is not wanted to run the program unless the species classification is possible, then it may be sensible to put in an instruction to terminate the program following lines 116, 1016 if the condition is not satisfied, and following line 103 (addition).

GENERAL COMMENT

None of the above matters will cause any trouble with a standard analysis of a normal-sized problem. The bugs are pretty unimportant in general. On the other hand, the restrictions can be serious when trying to analyse big problems. Provided that the level of the divisions is not altered, the main difficulty is in ensuring that there is enough room in the main data matrix. This amounts to ensuring that

- (i) the data when in pseudospecies form do not have length greater than $\frac{1}{2}(\text{NDAT})$
- (ii) the data when in non-pseudospecies form do not have length greater than $\frac{1}{2}(\text{NDAT})$.

Provided that these two conditions can be assured, there will probably not be too much difficulty in running large problems.

It is of course unfortunate that the matter of dimensioning is not made clearer by the program. This is one aspect of the fact that the program would be better if it could be simplified. However, I do not see myself having time to make a simplification for some years, and users must content themselves with its present form for the time being.

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26 March 1980

7 April 1980

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DIVISION OF BIOLOGICAL SCIENCES
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SECTION OF ECOLOGY & SYSTEMATICS

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- For your information
 For appropriate action
 Please comment and return
 For your files or disposition
 Please answer direct
 Need not return

To: TWINSPAN Users:

Mark Hill has supplied this list of bugs and restrictions for TWINSPAN, and we are circulating it to TWINSPAN users. Should problems or questions arise, it would ordinarily be best to write to Dr. Hill directly.

From Hugh G. Gauch, Jr.
Hugh G. Gauch, Jr.

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1. INTRODUCTION

1.1 TWINSPAN and indicator species analysis

TWINSPAN (TWO-way INdicator SPECIES ANALYSIS) is a computer program in FORTRAN designed primarily for ecologists and phytosociologists who have collected data on the occurrence of a set of species in a set of samples. The samples may be stands, relevés, stomach contents, island faunas, or whatever is appropriate to the study. TWINSPAN is a development of a method already published under the name "indicator species analysis" (Hill, Bunce & Shaw, 1975). However, it is much more flexible than previous programs for indicator species analysis, and involves many new features.

The most significant new feature is that the program first constructs a classification of the samples, and then uses this classification to obtain a classification of the species according to their ecological preferences. The two classifications are then used together to obtain an ordered two-way table that expresses the species' synecological relations as succinctly as possible.

1.2 Ordered two-way tables

Consider the table in Fig. 1. This is an ordered two-way table, and is the sort of structure that TWINSPAN aims to exhibit. (In fact it was obtained by TWINSPAN, but with non-standard parameter settings.)

The table has been ordered to exhibit the relation between the species and the samples as clearly as possible. The top five species are more abundant on the left side of the primary division than on the right side. The bottom five species are more abundant on the right side of the primary division than on the left side. The middle two species are somewhat indifferent, occurring widely on both sides.

For the most part, the species form a fairly well defined sequence, from those that prefer the left side of the primary dichotomy to those that prefer the right side of the dichotomy. However, one species "4 JUNC GER" appears out of sequence, at the end of the list. This is

SEE ERRATA

			11 1	
			219807631245	
9	SOL I	SEM	211-3-----	0000
5	PHRA	COM	53415-----	0001
3	IVA	FRU	452113-----	0010
11	SPAR	PAT	3254-5-----	0011
8	SCIR	OLN	-1---53-----	01
2	DIST	SPI	-5414545-424	100
1	ATRI	PAT	231--3221411	101
6	SAL I	EUR	----22-23411	1100
10	SPAR	ALT	--142-135553	1101
7	SAL I	VIR	-----34	1110
12	SUA E	MAR	-----54	1110
4	JUNC	GER	-----151---	1111
			000000111111	
			000001001111	
			00011 010011	
			00101 0101	
			01	

Fig. 1 Two-way table ordered and classified by TWINSPLAN (data of L. F. M. Fresco discussed in Gauch, 1977, pp. 70-74). Species names are shown at the left; sample numbers along the top. The classifications of species and samples are indicated along the right and bottom margins (notation is explained in Section 3.3 and need not concern us here). The main dichotomy for the species is indicated by a horizontal line, and the main dichotomy for the samples is indicated by a vertical line. Values indicate a scale of abundance, with absence of a species represented by the symbol "-".

not a mistake of the program, but is a result of the fact that this species behaves differently from the others, occurring with high abundance in a sample (number 6) where the commoner species are rather poorly represented.

The ordered table, therefore, is not simply an ordination. If it were, then the odd species "4 JUNC GER" would be put in an intermediate position. Rather, the aim of the arrangement is to throw the salient features of the data into sharp relief, by grouping like species with like, and like samples with like. In the example, the odd species "4 JUNC GER" is unlike any other (though it obviously has some relation to "8 SCIR OLN") and is therefore placed in an anomalous position.

1.3 Differential species

A differential species is one with clear ecological preferences, so that its presence can be used to identify particular environmental conditions. In Fig. 1 the species "3 IVA FRU" is a fine differential species. Its presence exactly demarcates the group of samples to the left of the primary division, and is strongly indicative of conditions in which the other three of the top four species are likely to be found. On the other hand the species "2 DIST SPI" is a poor differential species, not showing any marked affinity for a particular group of samples.

1.4 Strategy of analysis in TWINSPAN

TWINSPAN is designed to construct ordered two-way tables, and the method of doing so is by identification of differential species. In this respect it closely resembles the "hand" method of classification outlined by Mueller-Dombois & Ellenberg (1974, Chapter 9). It differs, however, in its treatment of the species. In the method outlined by Mueller-Dombois & Ellenberg, the species are classified at the same time as the samples. In TWINSPAN, on the other hand, the samples are classified first, and the species are classified second, using the classification of the samples as a basis.

The basic structure of TWINSPAN is as follows.

1. Classify the samples in a divisive hierarchy, dividing them first into 2 subsets, then 4, 8, 16, etc.
2. Convert the sample classification into an ordering.
3. Using the groups of samples as a basis, construct attributes for the species. For example, in Fig. 1, species 9, 5, 3, 11, 8 would be described as "preferential to the left side of the major division," and would be registered as possessing the attribute "preferential to the left side of the major division."
4. Classify the species in the same way as the samples, but with the difference that whereas the species were treated as attributes of the samples, the species have attributes of the kind indicated above.
5. Convert the species classification also into an ordering.
6. Print out the resulting ordered two-way table.

1.5 Making a dichotomy

The basic activity in TWINSPAN is to make a dichotomy. Indeed, as indicated in Section 1.4, all that the program does, in effect, is to divide up the samples into groups by repeated dichotomization, and then to do the same for the species.

It can be argued that this is a bad way to organize a table because dichotomies do not arise naturally. Indeed, Mueller-Dombois & Ellenberg (1974) recommend dividing the species into three categories, those that are preferential to one side of the division, those that are preferential to the other, and those that are indifferent. It was noted above (Section 1.2) that the species in Fig. 1 can be put into three such categories, comprising the first five species (preferential to the left), the next two (indifferent), and the last five (preferential to the right). But more careful inspection will show that these three categories are nearly as arbitrary as the two indicated in the figure. Indifference and preferentiality are a matter of degree; a sharp natural distinction between the two is often not found. Indeed, the dichotomy of the species in Fig. 1 shows that the species that are preferential to the left form a more clearly defined category than the species that are preferential to

the right--a fact that would be missed if the species were initially divided into three categories of equal status.

Given that a table can be structured satisfactorily by successive dichotomization, how are dichotomies to be made? The method suggested by Mueller-Dombois & Ellenberg is very roughly as follows.

1. Identify a direction of variation in the data--i.e. find a group of differential species that tend to occur together in the samples studied.
2. Construct an ordination by assigning each of these species a score of +1 and adding together the scores of the species in the sample.
3. Divide the ordination at an appropriate point into distinguishable communities.
4. Divide the species into categories according as to whether they are preferential or not to these communities.

This description of the method of "hand" classification is deliberately oversimplified, but does nevertheless encapsulate some of its essential features.

Now consider how this method might be applied to the table in Fig. 1. Suppose, for example, that it has been observed that species 5, 9, and 11 tend to occur together. Then these species can be declared as differential species and the samples can be ordinated thus:

Samples 9, 11, 12 - score +3

Samples 8, 10 - score +2

Sample 7 - score +1

Remainder - score 0.

If the samples are now arranged in order of descending score, it will rapidly become apparent that species 3 is also a differential species of the same group. If this species is now also given a score of +1, then the scores would work out like this:

Samples 9, 11, 12 - score +4

Samples 8, 10 - score +3

Sample 7 - score +2

Remainder - score 0.

Now there is a discontinuity, suggesting that the samples that score 0 should be distinguished from the rest.

TWINSPAN makes its dichotomies in a broadly similar manner to that described above. There are many complications, but the basic idea is the same. The stages of making a dichotomy in TWINSPAN are as follows.

1. Identify a direction of variation in the data by ordinating the samples. This ordination is referred to below as the "primary" ordination, and is made by the method of reciprocal averaging (Hill, 1973).
2. Divide the ordination at its middle to get a crude dichotomy of the samples.
3. Identify differential species that are preferential to one side or the other of the crude dichotomy.
4. Construct an improved ordination (referred to below as the "refined" ordination), using the differential species as a basis.
5. Divide the refined ordination at an appropriate point to derive the desired dichotomy.

If this were all, TWINSPAN would be relatively easy to describe. However, yet a third ordination is also constructed, the "indicator" ordination. This is based on a small number of the most strongly differential species, and is designed to provide a simple criterion for re-identification of the groups. So a further stage must be added.

6. Construct a simplified ordination, the "indicator" ordination, based on a few of the most highly preferential species. See whether the dichotomy suggested by the refined ordination can be reproduced by a division of the indicator ordination.

To summarize, TWINSPAN makes its dichotomies by dividing ordinations in half. There are three ordinations involved:

1. The primary ordination (reciprocal averaging), which is divided to obtain an initial, crude dichotomy;
2. The refined ordination, which is derived from the primary ordination through the identification of differential species; and
3. The indicator ordination.

With the exception of borderline cases (see Section 3.3), the refined

ordination is the one that is used to determine the dichotomy. The indicator ordination is essentially an appendage, put there for the convenience of users who want a succinct characterization of the dichotomy.

1.6 Warning about nomenclature

Unfortunately, the name "indicator species analysis" can cause much confusion. The method of division in TWINSPAN closely resembles the method published as "indicator species analysis" by Hill, Bunce & Shaw (1975). But, as already stressed, the indicator ordination is only an appendage, not the real basis of the method. "Differential species analysis" would possibly be a better name, but even this is not strictly accurate. Perhaps the best name would be "dichotomized ordination analysis," which would be a generic term to describe a wide variety of similar types of method.

Therefore, dear reader, please do not be confused by names. The basic method in TWINSPAN is the division of ordinations, and not the selection of indicator species.

1.7 Pseudospecies

The idea of a differential species is essentially qualitative, and to be effective with quantitative data must be replaced by a quantitative equivalent. This equivalent is the "pseudospecies" (Hill, Bunce & Shaw, 1975; Hill, 1977). The essential idea is that much of the quantitative information can be retained by expressing it on a relatively crude scale such as the Braun-Blanquet scale of cover-abundance (Mueller-Dombois & Ellenberg, 1974, p. 82; Westhoff & Maarel, 1973, p. 639). The levels of abundance that are used in TWINSPAN to define the crude scale are here termed "pseudospecies cut levels."

Consider, for the sake of example, the Braun-Blanquet scale. Ignoring the distinction between 1, +, and r, the scale is as follows:

1	0 - 4%	4	51 - 75%
2	5 - 25%	5	76 - 100% .
3	26 - 50%		

If quantitative data with cover expressed on a percentage scale are entered into TWINSPAN, then the Braun-Blanquet scale can be used by entering the pseudospecies cut levels:

0 5 26 51 76 .

Consider now a species, e.g. Poa annua, whose cover in one sample is 18% and in another is 36%. Although these values differ by a factor of 2, the samples have an important feature in common, namely that they share a moderate abundance of Poa annua. Given the Braun-Blanquet scale defined above, this fact can be expressed in terms of pseudospecies by saying that the samples contain the following pseudospecies:

Sample with Poa annua 18% - P. annua 1, P. annua 2;

Sample with Poa annua 36% - P. annua 1, P. annua 2, P. annua 3.

Hence, using the Braun-Blanquet scale, the samples are registered as having two pseudospecies in common, and one different. In spite of the rather large difference in cover, the samples are registered as having more in common than by way of difference.

The method of pseudospecies allows quantitative values to be used as differential "species" and as indicators. Thus, in TWINSPAN, instead of a differential species P. annua, it is possible to have a differential species P. annua 2, which--with the pseudospecies cut levels defined above--occurs when and only when P. annua has cover 5% or greater.

In practice, users of TWINSPAN who are mainly interested in a good tabular arrangement will hardly concern themselves with the technical detail of pseudospecies. For them it suffices to know that they are using a particular scale of abundance. However, there is one important practical point, namely that information on the occurrence of each pseudospecies is stored separately in the computer. Use of very numerous pseudospecies may therefore be undesirable with larger problems, as they take up space in the computer and use up additional computer time. The amount of space that they take up is printed in the output (see Section 3.2).

2. DATA INPUT AND OUTPUT

2.1 Input and output devices

Input for TWINSPAN is composed of two parts:

1. The data matrix itself, entered from input device 5; and
2. Parameters determining details of the analysis and output, entered from device 3.

If the operating system is interactive, then device 3 would ordinarily be a computer terminal.

Output from TWINSPAN is composed of two, or optionally three, parts:

1. Output to the lineprinter using the FORTRAN word PRINT;
2. Monitoring to output device 4, using the FORTRAN phrase WRITE(4,...); and
3. Optional output to device 7, using the FORTRAN phrase WRITE(7,...).

Output to devices 4 and 7 is intended to be construed as 1 line-control character and 79 other characters. No confusion will arise, however, if the control character is output explicitly. Indeed, in the case of output to device 7, the control character is always blank, so that the output will not even be unsightly.

Output to the printer uses 1 line-control character and a maximum of 124 other characters. Users with narrower lineprinters will need to modify the following FORMAT statements:

In subroutine QUIKIN: 1102, 1103, 1105, 1107;
In subroutine CLASS: 1010, 1004;
In subroutine REPORT: 1050;
In subroutine XLIST: 1000, 1001;
In subroutine YLIST: 1000;
In subroutine TABOUT: 1000, 2000.

Most of the changes are easy enough to make. In XLIST and YLIST it is necessary to change the values of NITEM (numbers of items printed out per line) as well as the FORMAT statements. In TABOUT, small economies of field-width can be achieved by omitting either the species names or the species numbers. Larger economies can be achieved by reducing the number of samples that can be included on a page of the table. At

present this number is 100, but it can be reduced by changing the value of the parameter NUMTAB (line 53 of source code).

The monitoring to device 4 is intended for users with an interactive operating system, and would ordinarily be sent to the console of a computer terminal. It is quite dispensable, and its main use is to allow checking of input parameters as they are entered. Furthermore, it is reproduced almost exactly in the output to the lineprinter, so that once the program has been run to completion, the output to device 4 serves no further purpose.

The output to device 7 is a copy of the sample and species classifications in machine-readable form. It is intended for users who wish to subject the data to further automatic data-processing (see Section 3.7 below). It is optional, and is not supplied unless requested by the user.

2.2 Entering the input parameters (standard analysis)

To get the standard analysis, 11 cards (or card images), each with a -1 on it, should be entered on device 3. All input parameters are read in free format. If the operating system is interactive, then device 3 would ordinarily be a computer terminal, and the user is then prompted by monitoring on device 4 (which would be the console of the terminal).

The standard analysis should suffice in many applications, but in others the user will want to vary the parameters. In particular, the pseudospecies cut levels provided by the standard analysis assume that either the data are quantitative with quantities given as percentages (percent cover, percent frequency, percent biomass, etc.), or the data are presence-and-absence data, in which case the quantities must all be 1. (It is permissible for the quantities all to be 0, but then there is a distinction between a species present with quantity 0--i.e. present but with unspecified quantity arbitrarily represented as 0--and species that are genuinely absent.)

To vary the parameters, values other than -1 must be supplied.

2.3 Omitting samples

The first action of the program is to read in the data matrix from device 5 (see Section 2.14 for the correct format). Then a message appears on the console (i.e. is written to device 4):

```
DO YOU WISH TO OMIT SOME SAMPLES?  
ENTER NUMBERS (NOT NAMES) OF ITEMS TO BE OMITTED  
ONE PER CARD, ENDING LIST WITH A -1.  
OTHER NEGATIVE NUMBERS DENOTE SEQUENCES. FOR EXAMPLE  
A 4 FOLLOWED BY A -8 OMITS ITEMS 4 THROUGH 8.
```

The standard response of -1 omits no samples. Typing

```
5  
-29  
35  
-1
```

omits samples 5-29, and 35. Omission of items is the one case where the monitoring on the terminal is not reproduced exactly in the output to the lineprinter. For ease of comprehension, the input considered above would be written

```
OMIT ITEMS      5      -29  
OMIT ITEM       35
```

in the output to the lineprinter.

2.4 Pseudospecies cut levels

Message on console:

```
ENTER NUMBER (NOT EXCEEDING 9) OF PSEUDOSPECIES CUT LEVELS  
OR TYPE -1 FOR DEFAULT CUT LEVELS, WHICH ARE 0 2 5 10 20.
```

If a value other than -1 is entered, then appropriate cut levels must be supplied. They must be supplied on the next card, and not on the same card as the number of cut levels. Correctly entered parameters would be (for example)

```
4  
0.0   0.5   1   5
```

The cut levels need not all be supplied on the same card; equally correct would be

```
4  
0.0   0.5  
1     5
```

A point to note about free-format input is that the program must know in advance how many values to expect. Hence the necessity of typing the number of cut levels first, on a separate card.

Levels should be chosen so as to reflect typical values of abundance, e.g. present, a little, a lot, more-or-less dominant. It is important, however, not to over-weight the effect of dominance by including many relatively high cut levels. With percentage data the default cut levels have proved very effective, and the user is recommended to adhere to them until he has some experience with the program.

2.5 Minimum group size for division

Message on console:

```
ENTER MINIMUM GROUP SIZE FOR DIVISION
TYPE -1 FOR DEFAULT VALUE (= 5); OTHERWISE TYPE VALUE REQUIRED,
WHICH MUST NOT LIE OUTSIDE LIMITS    0    10000
```

Groups smaller than this value are not divided further. Two points should be noted.

1. Groups much smaller than the specified size will often be formed, e.g. when the natural structure of the data is one large group and a few outliers.
2. The same minimum size for division is applied to the species classification as to the sample classification.

In general it is better to control the maximum level of divisions (Section 2.8) than to control the size of groups generated; though there certainly does come a size below which it is not worth dividing groups further.

2.6 Maximum number of indicators

Message on console:

```
ENTER MAXIMUM NUMBER OF INDICATORS PER DIVISION
TYPE -1 FOR DEFAULT VALUE (= 7); OTHERWISE TYPE VALUE REQUIRED,
WHICH MUST NOT LIE OUTSIDE LIMITS    0    15
```

This determines the maximum number of species that can be used in the indicator ordination (see Section 3.3 for a further explanation of what this involves). If no indicator ordination is desired, then this parameter can be set to zero. If a particular dichotomy can be specified precisely

by an indicator ordination based on a smaller number of indicators than the maximum, then the smaller number will be used.

2.7 Maximum number of species in final tabulations

Message on console:

```
ENTER MAXIMUM NUMBER OF SPECIES IN FINAL TABULATION  
TYPE -1 FOR DEFAULT VALUE (= 100); OTHERWISE TYPE VALUE REQUIRED,  
WHICH MUST NOT LIE OUTSIDE LIMITS 0 1000
```

In the two-way table at the end of the output, it is often inconvenient to clutter the diagram with rarer species. Therefore only the N commonest species are printed, where N is the value supplied. The default value is 100, which occupies about two pages of standard lineprinter output. If the value zero is supplied here, the final tabulation is omitted.

2.8 Maximum level of divisions

Message on console:

```
ENTER MAXIMUM LEVEL OF DIVISIONS  
TYPE -1 FOR DEFAULT VALUE (= 6); OTHERWISE TYPE VALUE REQUIRED,  
WHICH MUST NOT LIE OUTSIDE LIMITS 0 15
```

Even with large data sets there is little purpose in continuing to subdivide groups beyond a certain limit. Six levels of division are apt to produce about 64 groups. If there are larger numbers of groups, then interpretation can be difficult. Users will soon get a feel for how much division they require with any problem. Rarely will they want fewer than 4 levels, and almost never more than 7.

2.9 Diagrams of divisions

Message on console:

```
TYPE 1 IF DIAGRAMS OF DIVISIONS ARE WANTED.  
TYPE -1 FOR DEFAULT VALUE (= 0); OTHERWISE TYPE VALUE REQUIRED,  
WHICH MUST NOT LIE OUTSIDE LIMITS 0 1
```

In some applications it may be of interest to see how the indicator ordination relates to the refined ordination, and in particular to see whether the misclassifications are approximately borderline cases. If the value 1 is supplied, a scatter diagram relating the two ordinations

is printed out for each dichotomy (see Section 3.3 for what the diagrams mean).

2.10 Machine-readable copy of solution

Message on console:

TYPE 1 IF MACHINE-READABLE COPY OF SOLUTION TO BE WRITTEN TO DEVICE 7.
TYPE -1 FOR DEFAULT VALUE (= 0); OTHERWISE TYPE VALUE REQUIRED,
WHICH MUST NOT LIE OUTSIDE LIMITS 0 1

This refers to the optional output to device 7 mentioned in Section 2.1. Both the sample and species classifications are printed, together with the classes that items belong to in both binary and decimal notation. (See Section 3.3 for an explanation of the class numbering.)

2.11 Weights for levels of pseudospecies

Message on console:

ENTER WEIGHTS FOR LEVELS OF PSEUDOSPECIES.
FOR EXAMPLE WEIGHTS 1 2 2 2 SIGNIFY THAT PSEUDOSPECIES
CORRESPONDING TO 3 HIGHER CUT LEVELS ARE TO BE GIVEN TWICE
THE WEIGHT OF PSEUDOSPECIES AT THE LOWEST LEVEL.
TYPE -1 FOR DEFAULT VALUES (I.E. IF ALL VALUES ARE TO BE SET TO 1)
OR TYPE 0 IF NON-DEFAULT VALUES ARE TO BE ENTERED

If a 0 is typed at this stage, a further message appears:

NOW n INTEGER VALUES MUST BE SUPPLIED
WHICH MUST NOT LIE OUTSIDE LIMITS 0 1000000

The message is largely self-explanatory. The idea is that in some applications it may (for example) be desirable to pay particularly strong attention to species with high abundance. If so, then weights 1 1 1 1 3 could be supplied. Note that the number of values supplied should coincide with the number of cut levels, i.e. if there are 5 cut levels (as in the default case), then 5 weights should be supplied.

In many applications it is not necessary to use weights. However, with vegetation data taken from very large stands, it may sometimes be advisable to give presences and absences relatively low weight, as they may be due to anomalous small patches of terrain that occupy little of the stand area.

2.12 Indicator potential for cut levels

Message on console:

ENTER INDICATOR POTENTIALS FOR CUT LEVELS.
FOR EXAMPLE INDICATOR POTENTIALS 1 0 0 1 0 SIGNIFY
THAT PSEUDOSPECIES AT LEVELS 1 AND 4 CAN BE USED AS
INDICATORS, BUT THAT THOSE AT OTHER LEVELS CANNOT.
IN THE DEFAULT CASE, ALL PSEUDOSPECIES ARE AVAILABLE
AS INDICATORS.
TYPE -1 FOR DEFAULT VALUES (I.E. IF ALL VALUES ARE TO BE SET TO 1)
OR TYPE 0 IF NON-DEFAULT VALUES ARE TO BE ENTERED

If a 0 is typed at this stage, a further message appears:

NOW n INTEGER VALUES MUST BE SUPPLIED
NONE OF WHICH MUST LIE OUTSIDE LIMITS 0 1

The most common variation required here is that all the indicators should be real species and not pseudospecies. In this case, values 1 0 0 0 0 should be entered.

2.13 Omission of species from list of potential indicators

Message on console:

DO YOU WISH TO OMIT SOME SPECIES FROM LIST OF
POTENTIAL INDICATORS? SPECIES OMITTED FROM THIS LIST
ARE USED IN THE CALCULATION, BUT CANNOT APPEAR AS INDICATORS.
ENTER NUMBERS (NOT NAMES) OF ITEMS TO BE OMITTED
ONE PER CARD, ENDING LIST WITH A -1.
OTHER NEGATIVE NUMBERS DENOTE SEQUENCES. FOR EXAMPLE
A 4 FOLLOWED BY A -8 OMITS ITEMS 4 THROUGH -8.

This facility is to prevent particular species from being used in the indicator ordination. For example, if the indicator criterion is to be applied by relatively inexperienced field workers, it may be more useful if difficult taxonomic groups are omitted. Sequences of species are handled in the same way as sequences of stands (compare Section 2.3). Note that when species are barred from being indicators, they are nevertheless carried in the analysis, and will be printed out as usual in the final tabulation.

2.14 Format of the data matrix

The format used for input to TWINSPAN is the same as that for DECORANA (Hill, 1979), and is essentially the same as the "restricted condensed

format, input by samples" of ORDIFLEX (Gauch, 1977). Any matrix already in this format can be used directly by TWINSPAN, and any matrix in another format compatible with ORDIFLEX can be converted to the required format by the Cornell Ecology Program CONDENSE (Singer & Gauch, 1979).

Card 1 is a title card. This is reproduced at the front of the printed output and also at the front of the output to device 7.

Card 2 specifies two items:

1. The FORTRAN format in which the data have been written, using columns 1-60; and
2. The maximum number of couplets per card, this number being right-justified to column 70.

Users who are unfamiliar with FORTRAN format statements should consult a programmer who has used the language, or failing this any FORTRAN primer, or the documentation for ORDIFLEX (Gauch, 1977).

Card 3 is the beginning of the data matrix. The first item on each card is the index number of the sample. These index numbers must be in ascending order, except that an index number of zero terminates the matrix. Each species is identified by a number, and this is followed by a quantity. For example, the input

```
FUNNY LITTLE EXAMPLE  
(I2,2(I2,F4.1))  
1 1 1.3 3 1.0  
1 4 1.0  
2 1 .1 2 .2  
0  
SPEC 1 SPEC 2 SPEC 3 SPEC 4  
SAMP 1 SAMP 2
```

2 in column
70 of card 2

→ 2

signifies that sample 1 contains species 1 with abundance 1.3, species 3 with abundance 1.0, and species 4 with abundance 1.0. Sample 2 contains species 1 with abundance 0.1 and species 2 with abundance 0.2. Written out as a matrix, the above data are as follows:

Species	1	2	3	4
Sample				
1	1.3	0	1.0	1.0
2	0.1	0.2	0	0

Species names must be supplied next. These are abbreviated to binomials, each consisting of a 4-letter generic name and a 4-letter specific name, and are punched without spaces, so that ten 8-letter species names occupy an 80-column card.

Sample names are the last item to be supplied. Each of these is 8 letters long, so 10 sample names occupy an 80-column card.

Species names and sample names have been included in their correct place in the example above. A more substantial example is given in Fig. 2.

With the default analysis the lowest cut level is zero. In this case species entered with quantity zero are interpreted as present, but in small quantity, which is arbitrarily represented by a zero. A clear distinction needs to be maintained, therefore, between species that are absent--i.e. which are not listed among the species present in the sample--and species that are present but in small quantity that is represented by a zero.

2.15 Restrictions on reading the data matrix

As supplied, the program is dimensioned for up to 500 samples, 800 species, and 14500 non-zero items in the data matrix. These values can be increased by altering the declaration statements at the front of the main program (see Section 4.1).

The sample numbers must be ascending but need not be sequential. Samples can therefore be edited out of the matrix without making other changes. (But beware that the number of species names supplied still coincides with the highest species number in the data matrix.) Note well: this freedom may not apply to other Cornell Ecology Programs; for many of them the sample numbers must be sequential.

Species with no occurrences are permissible, and cause no difficulty. They do, however, use up array space and slightly slow down the calculation.

12 24TENNESSEE WATERSHED (G & G 1972 ORNL-IBP-71-10). TCTENA** S
 (I2,2X,8(I3,F6.2)) 8
 1 1 60.32 8 1.65 9 1.75 12 30.13
 2 1 72.95 8 1.58 9 0.92 12 0.79
 3 1 81.21 10 0.75 12 6.94
 4 1 90.23 2 0.87 8 1.70 10 3.80 12 7.07
 5 1 57.50 2 1.36 7 3.98 8 4.79 9 4.12 10 1.96 12 7.07
 6 1 74.63 2 0.20 4 0.87 5 8.69 6 1.01 8 6.15 10 8.95 11 0.35
 7 2 14.48 3 7.00 4 3.36 5 7.21 6 3.28 8 13.75 10 15.07 12 33.05
 8 2 5.51 6 11.80 9 0.96 11 5.32 12 37.05
 9 1 0.83 2 12.95 4 5.05 6 7.71 8 1.40 10 1.41 12 64.03
 10 2 17.31 3 3.98 5 4.19 6 14.29 9 2.21 11 1.79 12 41.83
 11 2 4.06 8 0.35 10 16.11 11 2.52 12 36.44
 12 2 4.26 4 5.14 5 8.25 6 0.75 8 4.94 10 1.01 12 19.09
 13 2 17.52 5 12.56 8 5.59 9 1.84 10 9.44 11 2.96 12 5.60
 14 2 19.43 3 2.82 4 0.83 5 14.91 6 16.45 9 0.55 10 16.40 11 9.19
 15 2 16.41 5 32.02 6 52.83 8 4.53 10 5.86 11 4.98
 16 2 34.57 3 3.65 4 1.10 5 15.23 6 9.69 8 4.51 9 8.08 10 11.24
 17 11 4.47 12 3.49
 17 2 50.76 3 18.27 4 2.18 5 6.13 6 10.22 8 4.24 9 2.21 10 2.78
 17 11 9.00
 18 2 11.75 3 3.89 5 13.20 6 22.79 8 10.27 9 1.81 10 10.79 11 19.86
 18 12 9.32
 19 1 8.44 2 5.50 5 46.06 6 7.25 8 1.97 9 7.55 10 8.04 11 12.75
 20 1 3.82 2 2.27 3 7.88 5 39.30 6 14.14 8 8.38 9 0.35 11 18.07
 21 2 1.77 5 65.58 6 3.67 8 1.87 10 1.92 11 3.54
 22 1 7.70 2 6.21 3 3.32 5 36.27 6 4.26 11 3.95
 23 1 1.10 2 9.95 3 7.11 4 19.21 5 40.07 6 1.77 8 4.91 9 1.27
 23 10 7.86 11 19.53
 24 3 10.04 5 46.32 7 13.45 11 0.92
 00
 PINU ECHCARY SPPQUER VELQUER RUBQUER PRIQUER ALBQUER COCNYSS SYLCORN FLOOXYD ARB
 ACER RUBLIRI TUL
 PINE 01PINE 02PINE 03PINE 04PINE 05PINE 06YELPR 07YELPR 08YELPR 09YELPR 10
 YELPR 11YELPR 120AKHK 130AKHK 140AKHK 150AKHK 160AKHK 170AKHK 18CHNOK 19CHNOK 20
 CHNOK 21CHNOK 22CHNOK 23CHNOK 24

Fig. 2 Data matrix in format acceptable to both TWINSPAN and ORDIFLEX.

3. INTERPRETATION OF THE OUTPUT

3.1 Reading the data matrix

The first part of the output (Fig. 3) concerns the input of the data matrix and the omission of samples. It is self-explanatory except for the print-out of the data matrix. Only the beginning and end of the matrix are printed, and all quantities are multiplied by 1000 and rounded to the nearest integer. Values of -1 are used to indicate the end of a sample. Thus a record

```
1 46000 5 2500 -1 1 56000 6 4500 -1
```

would refer to two samples, one with species 1 having quantity 46.0 and species 5 having quantity 2.5, and the other with species 1 having quantity 56.0 and species 6 having quantity 4.5. The matrix is printed out in the form that is held in the computer in order to remind the user of two things:

1. How the length of the raw data array relates to the number of non-zero items in the data matrix; and
2. That quantities smaller than 0.001 are lost in roundoff.

3.2 Entry of input parameters

This part of the output is also self-explanatory, and if the program is run interactively it merely repeats what was printed out on the console at run time. If the program is not being run interactively, then the input parameters should be checked carefully for correctness.

After the input parameters, three important statistics on the data are printed out:

1. Length of data array after defining pseudospecies;
2. Total number of species and pseudospecies; and
3. Number of species, excluding pseudospecies and ones with no occurrences.

With big problems these statistics may be relevant to determining whether a species classification can be made as well as the sample classification (see Section 4.2), or whether the user can afford to use more cut levels for the pseudospecies (Section 4.1).

94 25MEADOW RELEVES (MUELLER-DOMBOIS & ELLENBERG 1974 182-183). TCHEAD** S

NUMBER OF SAMPLES 25

NUMBER OF SPECIES 94

LENGTH OF RAW DATA ARRAY 1601

1	6000	7	100	10	1000	11	50000	12	100	13	1000
16	2000	20	100	21	100	23	1000	26	5000	27	1000
32	2000	34	15000	37	3000	44	1000	45	100	48	3000
52	100	57	1000	67	1000	68	100	69	4000	75	100
77	100	81	100	85	100	86	1000	88	100	89	100
91	100	-1	1	100	2	100	7	1000	10	1000	20
1000	21	100	23	100	25	100	26	5000	27	1000	32
1000	33	5000	37	2000	44	1000	51	1000	59	100	67
1000	68	100	69	74000	75	100	81	2000	84	2000	86
100	88	100	90	2000	-1	1	3000	7	2000	8	100
.....
2000	-1	1	3000	3	100	7	30000	9	5000	11	10000
12	100	13	100	20	2000	23	6000	25	100	26	8000
33	2000	34	2000	37	3000	39	100	44	1000	45	100
54	100	67	100	68	100	69	20000	74	100	75	100
79	4000	85	100	88	2000	90	2000	91	100	92	100
94	100	-1	1	100	2	100	4	1000	5	100	6
100	7	35000	13	100	15	2000	17	1000	24	3000	26
18000	33	3000	37	2000	41	100	45	100	46	15000	55
100	56	100	58	4000	67	100	69	10000	71	4000	75
100	77	1000	78	100	85	100	89	100	91	1000	-1

SPECIES NAMES

1 ACHI MILLI	2 AJUG REPTI	3 ALCH VULG	4 ALOP PRATI	5 ANGE SILVI	6 ANTH SILVI	7 ARRH ELATI	8 BELL PERE
9 BRAC PINNI	10 BRIZ MEDII	11 BROM EREC	12 CAMP GLOMI	13 CAMP ROTWI	14 CARD PRATI	15 CARE ACUTI	16 CARE FLAC
17 CARE GRACI	18 CARE HIRTI	19 CARE PANII	20 CENT JACEI	21 CERA CAESI	22 CHEN ALBUI	23 CHRY LEUCI	24 CIRS OLER
25 CREP BIENI	26 DACT GLOMI	27 DAUC CARO	28 DESC CAESI	29 DIAN SUPEI	30 EUPH ODONI	31 FEST ARUMI	32 FEST OVIN
33 FEST PRATI	34 FEST RUBRI	35 FILI ULMAI	36 GALI BOREI	37 GALI MOLLI	38 GALI ULIGI	39 GALI VERUJ	40 GERA PRAT
41 GEUM RIVAI	42 GLEC HEDEI	43 GLYC FLUI	44 HELI PUBEI	45 HERA SPHOI	46 HOLC LANAI	47 KNAU ARVEI	48 KOEL PYRA
49 LAMI ALBUI	50 LATH PRATI	51 LEON HISPI	52 LINU CATHI	53 LOLI PEREI	54 LOTU CORNI	55 LYCH FLOSI	56 LYSI NUMM
57 MEDII LUPU	58 MELA DIURI	59 MYOS ARVEI	60 MYOS PALUI	61 PAST SATII	62 PHAL ARUNI	63 PHLE PRATI	64 PHRA COMM
65 PIMP MAGNI	66 PIMP SAXII	67 PLAN LANC	68 PLAN MEDII	69 POA PRATI	70 POA TRIVI	71 POLY BISTI	72 POLY CONV
73 POTE REPTI	74 PRUN VULGI	75 RANU ACERI	76 RANU REPEI	77 RUME ACETI	78 RUME CRISI	79 SALV PRATI	80 SANG OFFI
81 SCAB COLU	82 SENE JACOI	83 SILA PRATI	84 SILE INFRI	85 TARA OFFI	86 THYM SERPI	87 TRAG PRATI	88 TRIF PRAT
89 TRIF REPEI	90 TRIS FLAVI	91 VERO CHAMI	92 VICI CRACI	93 VICI SEPII	94 VIOL HIRTI		

SAMPLE NAMES

1	1		2	2		3	3		4	4		5	5		6	6		7	7		8	8	
9	9		10	10		11	11		12	12		13	13		14	14		15	15		16	16	
17	17		18	18		19	19		20	20		21	21		22	22		23	23		24	24	
25	25																						

DO YOU WISH TO OMIT SOME SAMPLES?

ENTER NUMBERS (NOT NAMES) OF ITEMS TO BE OMITTED

ONE PER CARD, ENDING LIST WITH A -1.

OTHER NEGATIVE NUMBERS DENOTE SEQUENCES. FOR EXAMPLE

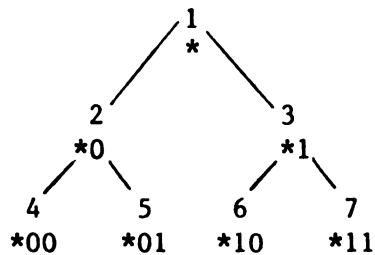
A 4 FOLLOWED BY A -8 OMITS ITEMS 4 THROUGH 8.

-1

Fig. 3 Lineprinter output derived from reading data matrix into TWINSPLAN. Sample names are here the same as sample numbers, but there is no necessity for this.

3.3 Classification of the samples

Divisions are made successively, according to the scheme below.



Each group is represented by two numbers, one in decimal, one in binary notation. If, in the hierarchy above, the symbol * is replaced by the number 1, then the decimal and binary numbers can be seen to be identical. Thus 10 is the representation of 2 in binary code, 110 represents 6, etc.

In general, group n is divided to obtain group $2n$ (the "negative" group) and group $2n+1$ (the "positive" group). The binary representation of the nodes of the hierarchy is more directly interpretable than the decimal representation, 0 denoting a left arm and 1 denoting a right arm.

Consider the output for a dichotomy line by line (refer to Fig. 4).

Line 1 gives the number of the group to be divided, both in decimal and in binary notation, together with the number of elements that it contains.

Line 2 gives information on the primary ordination (reciprocal averaging). Each iteration involves eight passes of the data.

Lines 3-5 specify the indicator ordination. Each sample is given an "indicator score" found by adding +1 for each positive indicator and -1 for each negative indicator that it contains. The sign (+ or -) of each indicator follows immediately after its name in the list of indicators. The division derived from the indicator ordination is specified on line 5. The two values printed are the maximum indicator score for a sample to be assigned to the negative group, and the minimum score for it to be assigned to the positive group. If these values are A and B respectively, then B is always $1+A$.

DIVISION 1 (N= 25) I.E. GROUP *

EIGENVALUE 0.308 AT ITERATION 1

INDICATORS, TOGETHER WITH THEIR SIGN

CIRS OLER1(-) BROM EREC1(+) HOLC LANAI(-) GEUM RIVAL(-)

MAXIMUM INDICATOR SCORE FOR NEGATIVE GROUP -1 MINIMUM INDICATOR SCORE FOR POSITIVE GROUP 0

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
20	21	22	23	24	25													

ITEMS IN NEGATIVE GROUP 2 (N= 17) I.E. GROUP *0

5 6 7 8 11 12 13 14 16 17 18 19

BORDERLINE NEGATIVES (N= 1)

16

ITEMS IN POSITIVE GROUP 3 (N= 8) I.E. GROUP *1

1 2 3 4 9 10 15 24

NEGATIVE PREFERENTIALS

AJUG REPT1(14, 2)	ALOP PRAT1(9, 0)	ANGE SILVI1(6, 0)	CARE ACUT1(5, 0)	CIRS OLER1(14, 0)	DESC CAES1(7, 0)
FILI ULMAL1(4, 0)	GEUM RIVAL1(14, 0)	GLEC HEDE1(6, 0)	HOLC LANAI1(12, 0)	LATH PRAT1(9, 1)	LYCH FLOS1(7, 0)
LYSI NUMMI1(8, 0)	MELA DIUR1(11, 0)	PIMP MAGNI1(4, 0)	POLY BIST1(4, 0)	RUME ACET1(16, 3)	TRAG PRAT1(6, 1)
ALOP PRAT2(8, 0)	CARE ACUT2(4, 0)	CIRS OLER2(8, 0)	DESC CAES2(6, 0)	HELI PUBE2(9, 1)	HOLC LANAI2(6, 0)
PLAN LANC2(8, 0)	RUME ACET2(4, 0)	TARA OFFI2(4, 0)	ALOP PRAT3(5, 0)	CIRS OLER3(4, 0)	DESC CAES3(4, 0)
GALI MOLL3(10, 2)	HELI PUBE3(4, 0)	TRIS FLAV3(9, 1)	ARRH ELAT4(12, 2)	CIRS OLER4(4, 0)	DACT GLOM4(12, 1)
FEST PRAT4(8, 0)	ARRH ELAT5(7, 1)				

POSITIVE PREFERENTIALS

BRIZ MED1(1, 4)	<u>BROM EREC1(0, 7)</u>	CAMP GLOM1(2, 5)	CAMP ROTU1(6, 7)	CARE FLAC1(3, 3)	DIAN SUPE1(2, 2)
FEST OVIN1(0, 4)	GALI BORE1(2, 2)	GALI VERUI1(0, 2)	KNAU ARVE1(1, 2)	KOEL PYRA1(0, 4)	LEON HISPI1(2, 2)
LINU CATH1(1, 4)	LOTU CORN1(2, 2)	PIMP SAXI1(0, 2)	SALV PRAT1(0, 5)	SCAB COLU1(0, 5)	SILE INFIL1(1, 2)
THYM SERP1(0, 5)	VICI CRAC1(2, 3)	VICI SEPIL1(3, 4)	VIOL HIRT1(1, 5)	BROM EREC2(0, 7)	FEST OVIN2(0, 2)
FEST RUBR2(4, 6)	KOEL PYRA2(0, 4)	SALV PRAT2(0, 4)	VIOL HIRT2(0, 2)	BROM EREC3(0, 7)	CHRY LEUC3(0, 2)
FEST RUBR3(0, 2)	<u>BROM EREC4(0, 7)</u>	POA PRAT4(5, 5)	BROM EREC5(0, 6)	POA PRAT5(1, 2)	

NON-PREFERENTIALS

ACHI MILL1(13, 8)	ANTH SILVI1(4, 1)	ARRH ELAT1(17, 8)	BELL PERE1(8, 3)	CENT JACE1(9, 5)	CERA CAES1(8, 5)
CHRY LEUC1(15, 8)	CREP BIEN1(12, 3)	DACT GLOM1(17, 8)	DAUC CARO1(13, 7)	FEST PRAT1(17, 6)	FEST RUBR1(9, 6)
GALI MOLL1(17, 8)	HELI PUBE1(11, 5)	HERA SPHO1(11, 3)	MEDI LUPU1(11, 6)	PLAN LANC1(17, 8)	PLAN MED1(8, 6)
POA PRAT1(17, 8)	PRUN VULG1(9, 3)	RANU ACER1(16, 7)	SENE JACO1(4, 2)	TARA OFFI1(13, 6)	TRIF PRAT1(12, 6)
TRIF REPE1(11, 3)	TRIS FLAV1(10, 5)	VERO CHAM1(15, 7)	ACHI MILL2(7, 6)	ARRH ELAT2(17, 6)	CARE FLAC2(3, 2)
CENT JACE2(6, 2)	CHRY LEUC2(5, 4)	DACT GLOM2(16, 8)	FEST PRAT2(17, 6)	GALI MOLL2(16, 7)	POA PRAT2(15, 8)
RANU ACER2(4, 1)	TRIS FLAV2(10, 5)	ACHI MILL3(4, 2)	ARRH ELAT3(13, 4)	DACT GLOM3(15, 7)	FEST PRAT3(10, 3)
POA PRAT3(10, 7)	GALI MOLL4(4, 1)				

END OF LEVEL 1

Fig. 4 First dichotomy resulting from application of TWINSPLAN to the meadow data considered by Mueller-Dombois & Ellenberg (1974, pp. 182-183). The indicators have been underlined. It may seem surprising that the pseudospecies GALI BORE1(2, 2) is a positive preferential. The reason for this is that it occurs in 2/8 = 25% of the samples in the positive group, and in 2/17 = 11.8% of the samples in the negative group. Hence it is more than twice as likely to occur in the positive group than in the negative group.

In the example, only those samples with an indicator score of -1 or less are included in the negative group. Referring to the two-way table (Fig. 7), it can be seen that the borderline sample (number 16) contains only one indicator, Geum rivale. This gives it an indicator score of -1, assigning it to the negative group. Sample 2, on the other hand, contains no indicators at all, and hence has an indicator score of 0, assigning it to the positive group. At the other extreme, sample 8 contains all three of the negative indicators and hence has a score of -3.

The indicators are listed in an approximate order of effectiveness. Thus in Fig. 4, Cirsium oleraceum (negative) is a better indicator than Bromus erectus (positive), etc. It may seem strange that Geum rivale, dividing (14,0), is a worse indicator than Holcus lanatus, dividing (12,0). The reason for this is that the measure of effectiveness pays less attention to samples near the center of the refined ordination than to those at the extremes, and furthermore, that it measures effectiveness for a division at the center of the refined ordination rather than for the slightly off-center dividing line that is actually chosen (see below). Technically, the effectiveness of indicators is measured by the absolute value of the "preference index," defined in the remarks under parameter FEEBLE in Section 4.10.

Scatter diagram relating indicator ordination to refined ordination is not printed out in the default case, but is included here to show what is available. The lineprinter output is highly schematic, and corresponds to the relation shown in Fig. 5. To interpret the lineprinter output in Fig. 4, the following points should be noted.

1. For ease of computing the ordination is divided into segments. These have no special significance, but are merely a convenient way of calculating where to locate the zone of indifference. There are five possible positions for the zone of indifference such that it lies entirely within the critical zone (see Fig. 5). The location of the zone of indifference is selected to minimize the number of misclassified samples (see below for how these are defined).
2. Segments of the refined ordination are shown along top of scatter

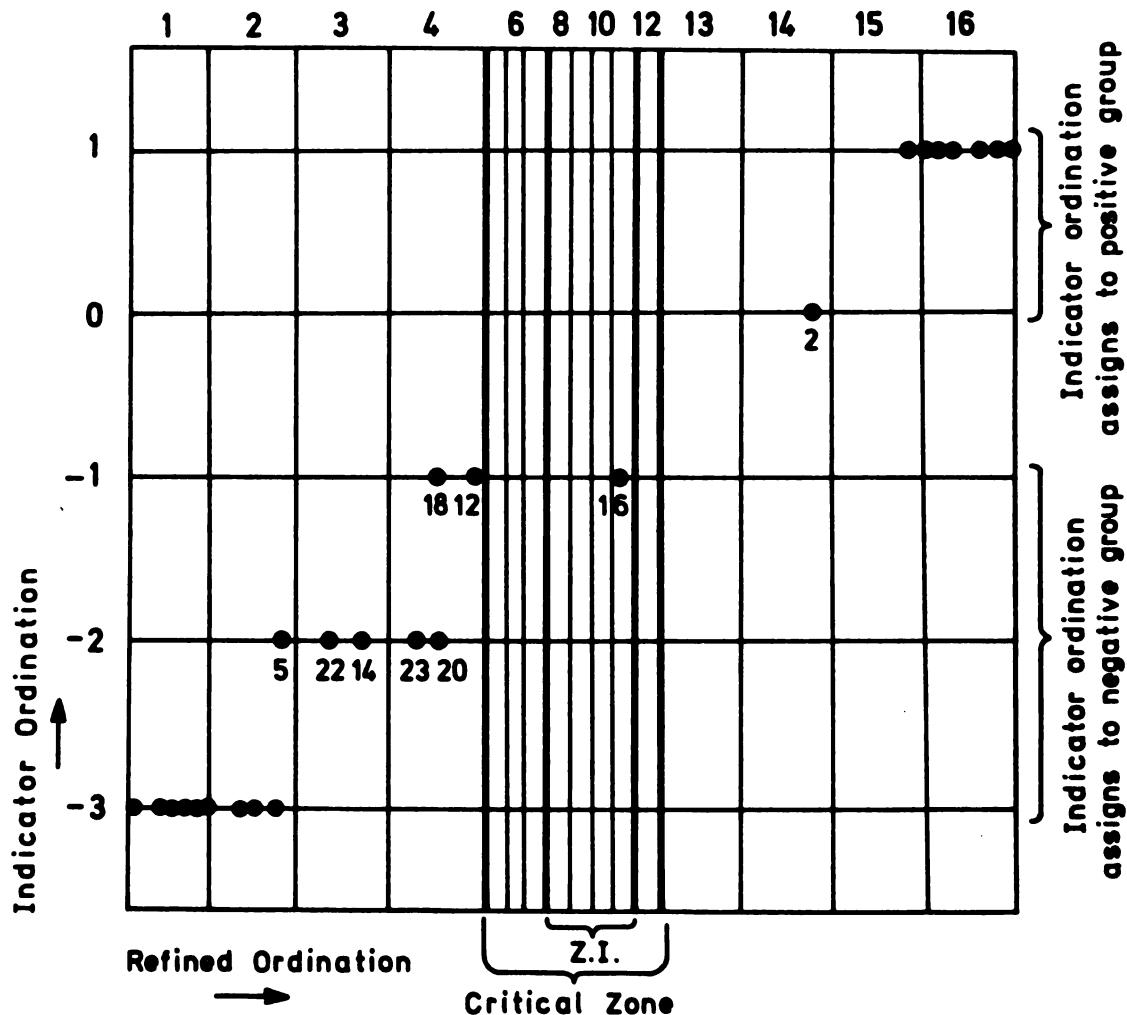


Fig. 5 Relation between indicator ordination and refined ordination (same dichotomy as Fig. 4). The segments of the refined ordination are indicated along the top of the diagram, and should be compared with the scatter diagram in Fig. 4. Segments 5-12 constitute the "critical zone," and segments 8-11 constitute the "zone of indifference" (abbreviated Z.I.) in the diagram. Samples are designated as "borderline cases" if they lie in the zone of indifference. In this example, only sample 16 is borderline.

diagram, which is not to scale. The critical zone (segments 5-12) is 20% of the length of the whole ordination. The length of the segments within the critical zone is one-quarter of the length of the segments outside it.

3. Indicator scores are shown along the right of the diagram. Indicator scores for assignment to the positive group are separated from those for assignment to the negative group by a row of stars.
4. The zone of indifference (here comprising segments 8-11) is indicated by columns of stars.
5. Values printed in the diagram indicate numbers of samples in each category. Thus the value 6 appears at the bottom left of the scatter diagram, corresponding to the six samples shown in Fig. 5 that occur in segment 1 and have indicator score -3.

Items in negative group are listed. The samples are specified by their names, which in this case are the same as their numbers.

Borderline negatives are those items that are in the negative group and which also lie in the zone of indifference (see Figs. 5 and 6). In Fig. 5, sample 16 is a borderline negative, and is assigned to the negative group because it has an indicator score of -1. The refined ordination is highly polarized, so that there are few borderline cases. In this respect, TWINSPAN differs from the version of indicator species analysis published by Hill, Bunce & Shaw (1975; see especially p. 602). In the original version there was no process of refinement, the "refined" and primary ordinations being identical; borderline cases were accordingly more numerous. A borderline case is assigned to its class according to its indicator score, the refined ordination being held in this case to be indecisive.

Misclassified negatives are those samples lying to the left of the zone of indifference but whose indicator score would assign them to the positive arm of the dichotomy (Fig. 6). This is regarded as a failure on the part of the indicator ordination to reproduce the dichotomy accurately, and hence as a "misclassification."

1 2 3 4 5 6 7** 8 91011**1213141516****

Misclassified **Border- ** Positives** 1
Negatives **line +ve** ** 0

 **Border- ** **-1
 line **Misclassif-2
Negatives **negative**Positives **-3

Zone of
Indifference

1 2 3 4 5 6 7** 8 91011**1213141516****

0 0 0 0 0 0** 0 0 0 0** 0 0 0 1 6** 1
0 0 0 0 0 0** 0 0 0 0** 0 0 1 0 0** 0

0 0 0 2 0 0 0** 0 0 0 1** 0 0 0 0 0**-1
0 1 2 2 0 0 0** 0 0 0 0** 0 0 0 0 0**-2
6 3 0 0 0 0 0** 0 0 0 0** 0 0 0 0 0**-3

Fig. 6 Categories of samples in relation to indicator scores and zones of the refined ordination. The scatter diagram in Fig. 4 is reproduced here for the sake of comparison. Indicator scores are shown along the right-hand margin. Samples with indicator score -1 or less are assigned to the negative group unless they lie to the right of the zone of indifference, in which case they are categorized as "misclassified positives." Samples in the zone of indifference are categorized according to their indicator score.

Items in positive group, borderline positives, and misclassified positives are defined as for negatives.

Negative preferentials are those pseudospecies and species that are at least twice as likely to occur on the negative side as on the positive side. Only those that occur in at least 20% of the samples on the negative side are listed. Values given in brackets are actual numbers of occurrences, so that the entry

FEST PRAT4(8, 0)

signifies that Festuca pratensis occurs with abundance 4 or more (i.e. with cover at least 10%) in 8 samples on the negative side of the dichotomy and in no samples on the positive side of the dichotomy. This can be checked by referring to the two-way table in Fig. 7. It should be noted that when there is a very uneven split, negative preferentials can easily occur in more samples on the positive side of the dichotomy than on the negative side. Suppose, for example, that the split is into 2 samples on the negative side and 10 on the positive. Then a species that occurs in 2 samples on the negative side and 4 samples on the positive side is a good negative preferential. It has 100% frequency on the negative side and 40% frequency on the positive side, and is therefore 2.5 times as likely to occur on the negative side as on the positive side.

Positive preferentials are defined as for negative preferentials; non-preferentials are pseudospecies and species that are reasonably common and which are not preferentials. Here again, only those that achieve 20% frequency on one side or the other are listed.

3.4 Classification of the species

Species are classified by TWINSPAN in much the same way as samples. However, there is an important difference in that the species classification is made in the light of the sample classification, and not using the raw data. In fact, the species classification is made on the basis of fidelity, i.e. using the degree to which species are confined to particular groups of samples. For example, in Fig. 7, Cirsium oleraceum

and Phalaris arundinacea are both completely faithful to group *0 of the sample hierarchy--i.e. they have no occurrences outside this group. However, Phalaris is also completely faithful to group *00 of the sample hierarchy, whereas Cirsium is not.

Species are assigned attributes according to differing degrees of fidelity to sample groups (see Section 4.4 for details). For example, both Cirsium oleraceum and Phalaris arundinacea will be registered as having the attribute "very highly faithful to group *00," but Phalaris is also very highly faithful to group *00, whereas Cirsium is not. Indeed, Cirsium occurs widely outside group *00, in group *01.

Because the species classification is made on the basis of fidelity to groups defined by the sample classification, altering the level of division of the classification (cf. Section 2.8) may alter the species classification even if no other changes are made.

The species classification differs from the sample classification in that "indicator characters" are of little interest. It is scarcely of much significance to know that (for example) "low level of fidelity to sample group X" is a preferential attribute of a particular group of species. Consequently, the output has been shortened for the species classification, to present only the dichotomies, without the additional information that is supplied during the sample classification.

If the data set is very large, there may be no room for the new matrix of species' attributes as well as the original matrix. If so, then the species classification is abandoned (see Section 4.2 for conditions under which this will happen).

3.5 Order of species and samples

These orderings are printed out for two purposes. In the first place, the final table may not contain all the less common species, so that the orderings may be convenient as an indication of where the rarer species would come in the arrangement. Secondly and more important, the final tabulation refers to the samples by number, not by name as elsewhere in

the program. The reason for this is that it is difficult to write the names vertically, owing to the poor character-manipulation of the FORTRAN language. A previous version of TWINSPAN that did write the names vertically was found to be machine-specific, and was therefore unsuitable for issue as a general-purpose program.

Users with programming ability, who wish to print sample names in the table, should use the vectors INAME1, INAME2 instead of INFLAG in subroutine TABOUT.

3.6 Tabular arrangement of species and samples

The tabular arrangement (Fig. 7) needs little explanation. The hierarchy is shown in binary notation (omitting the '*'s) along the right-hand and bottom margins. Numbers and names of species, and numbers of the samples, are shown along the left-hand and top margins.

The samples in Fig. 7 are rather homogeneous, and there are only two major groups. Sample 19 is anomalous in having several rare species abundant in it. Group *010 differs from *011 in having more of Carex acutiformis, Cirsium oleraceum, Deschampsia caespitosa, and Holcus lanatus, and less of Achillea millefolium and Trisetum flavescens; but the difference is a small one. Likewise, group *10 differs little from *11, the absence of Koeleria pyramidata being the main distinguishing feature. Further discussion of this set of data can be found in Maarel, Janssen & Louppen (1978).

3.7 Output to device 7

This output is intended to be read by machines rather than people. It lists the number, name, and class of each sample and each species. The classes are specified in both decimal and binary notation. (See Section 3.3 for an explanation of the class numbering.)

The purpose of writing out the solutions in machine-readable form is twofold. In some applications it may be desirable to subject a group derived from the classification to further analysis, e.g. to an ordination. Members of the selected group can be picked out by specifying their class

Fig. 7 Two-way table resulting from an application of TWINSPAN to the meadow data considered by Mueller-Dombois & Ellenberg (1974, pp. 182-183). Values denote categories of abundance defined by the pseudospecies cut levels. In this case the cut levels were the default values 0, 2, 5, 10, 20, and the raw data specified abundance by percent biomass. Values therefore denote categories of percent biomass:

1, 0-1% 2, 2-4% 3, 5-9%
 4, 10-19% 5, 20-100%.

Vertical lines separate classes of samples at level 3; horizontal lines separate classes of species at level 4. Note that the arrangement is approximately on the positive diagonal.

number and scanning the list of samples and classes. Alternatively, it may be desired to tabulate the classification in another form. The list of classes can be used to do this.

4. TECHNICAL DETAILS OF THE PROGRAM AND ALGORITHM

4.1 Dimensioning

All subroutines have variable dimensioning so that only the main program needs to be modified. There are three dimensions that may need to be modified:

NDAT (= 30000)
MMAX (= 500)
NMAX (= 800).

NDAT is the length of the data array. The data are stored as indicated in Fig. 3, and are never expressed as a two-dimensional array except in the final tabulation. Hence, the magnitude of the calculation increases only linearly with the amount of data supplied. It depends mainly on the number of non-zero items in the matrix, and to a much lesser extent on the number of species or samples. It does, however, depend also on the number of pseudospecies, as information on each pseudospecies is stored separately. With large problems it is possible to overrun the array storage by asking for a large number of pseudospecies. In such cases an error message is printed, and the user has two options: either to reduce the number of pseudospecies, or to increase the value of NDAT.

To increase the value of NDAT it is necessary to modify lines 22, 37, and 49 of the program. It will be seen in line 22 that JDAT is half the length of IDAT; this proportionality should be maintained. Also, in line 37, JDAT(1) is equivalenced to IDAT(NDAT/2+1), and this relationship should be maintained. The intention here is to allow the sample data to overrun into space that is not required until the species classification is worked on. In some cases it is possible for NDAT to be big enough to allow the samples to be classified, but not big enough to allow the species classification to be made as well (see Section 4.2 for details).

MMAX is the maximum number of samples that can be considered. In the listing supplied here, MMAX is set to 500. Users with larger problems should modify lines 23-25, 32, 33, 50. If the number of species is greater

than MMAX, then the species classification is impossible. In this case a warning message is printed, and the calculation will terminate at the end of the sample classification.

NMAX is the maximum number of species and pseudospecies that can be handled. For example, if a problem has 700 species, and if the user wishes to use 3 levels of pseudospecies, then he will probably need to set NMAX to about 1200. The theoretical maximum of 2100 will not be required, as not all species will attain the abundance demanded by the top cut level. If the program needs a larger value of NMAX, then an appropriate error message is printed. To modify NMAX, lines 26-28, 34, and 51 will need to be altered.

NMAX must also exceed the maximum species number in the data matrix. This may be relevant if there are many species with no occurrences.

4.2 Dimensions necessary for species classification

To make the species classification, three conditions must be satisfied.

1. There must be enough room in the array JDAT for a secondary data matrix specifying fidelity attributes of the species (see Section 4.4). This condition is rarely likely to be violated.
2. The data when held in pseudospecies form must not occupy more than half the array IDAT. If they do occupy more than this, a warning message is printed.
3. The number of species (after eliminating those with no occurrences) must not exceed MMAX, the upper bound on the number of samples. If it does exceed MMAX, a warning message is printed.

In addition, a fourth condition must be met if the full sample classification is to be used in classifying the species.

4. Let the largest number designating a group of samples be ICMAX. Usually, ICMAX is approximately $2^{**}(\text{LEVMAX}+1)$, where LEVMAX is the maximum level of division. Then NMAX must be greater than ICMAX*3. If this condition is not met, the lower-level groups of the sample classification are automatically amalgamated. This should not affect the output or the species classification

too seriously, but it will mean that the lowest-level sample groups are lost in the final tabulation.

Conditions 2 and 3 are the most likely to be violated, and the user will then get a warning. (Note that this is printed before the sample classification.) However, if the species classification is not made and no warning is printed, then condition 1 may be suspected of having failed, and NDAT should be increased. However, a simpler and better strategy is probably to make separate analyses of subsets of the data, using the output to device 7 to divide the samples into appropriate subsets (cf. Section 3.7).

4.3 Structure of the main program

The main program performs the following activities.

1. Values are set for a number of parameters (cf. Section 4.10).
2. The $MM \times NSPEC$ data matrix $A = [a_{ij}]$ is entered (subroutine QUIKIN). There are MM samples and NSPEC species. The matrix is stored in a linear array IDAT, in the form indicated in Section 3.1.
3. Parameters are supplied by the user (subroutine ENTER).
4. Quantities are converted to pseudospecies using cut levels supplied by the user (subroutine PSEUDO; cf. Section 4.5). The total number of pseudospecies is NN, and NSPEC is revised downwards if there are species with no occurrences in the matrix.
5. The samples are classified (subroutine CLASS; see Section 4.6).
6. Attributes are defined for the species, using the degree of fidelity (sensu Braun-Blanquet) of species to the groups of samples derived by the sample classification (see Section 4.4 for details). A data matrix for the species is put in JDAT, and a version of the original data matrix is retained in IDAT.
7. The species are classified (second application of subroutine CLASS).
8. The species and sample classifications are converted to orderings. This may involve reversing the order of the species, so as to get the data matrix on the positive diagonal.
9. The data matrix is rearranged so that within each sample the species appear in the correct order obtained at Activity 8 above (subroutine JORDER).
10. The final two-way table is printed out (subroutine TABOUT).

4.4 Conversion of data matrix for species classification

Activity 6 above is basic to the species classification. The rationale behind using fidelity is that species should be considered as similar or dissimilar not according to whether they are common or rare, but according to whether they occur in similar sample groups. Of course, an exceedingly common species such as Arrhenatherum elatius in Fig. 7 will not be highly faithful to any group, and will therefore not appear similar to any exceedingly rare species. However, A. elatius is classified in the group 011 along with several relatively uncommon species such as Potentilla reptans and Senecio jacobaea. These species resemble Arrhenatherum in that they show a poor relation to the classification of the samples.

Fidelity is defined as follows. Consider the class of samples whose index is IC (refer to Section 3.3 for an explanation of the index numbers of the classes). Then the fidelity of species J to class IC is defined by the ratio

$$\text{RAT}(\text{IC}, \text{J}) = \frac{\text{mean occurrence of J in class IC}}{\text{mean occurrence of J in individuals not in IC}} .$$

In this context, the mean occurrence of J is defined to be the unweighted mean value of b_{ij} for those samples i that occur in the group under consideration; where the matrix elements b_{ij} are those printed out in the final tabulation (i.e. $b_{ij} = C$ if species j achieves cut level C in sample i).

Using the fidelity ratio RAT, it is possible to define attributes for the species. Species J is deemed to have attribute F(IC,K) if $\text{RAT}(\text{IC}, \text{J})$ is greater than or equal to the limit SPE(K). Three values of SPE are used here (lines 100-102 of source code), and are:

$$\text{SPE1} = 0.8; \text{SPE2} = 2.0; \text{SPE3} = 6.0.$$

In the species classification, both the species and their attributes are given differing weights, as follows.

1. Extra weight for high fidelity. A parameter WTHIGH, set at the beginning of the program to 2.0, is used to give extra weight to

the higher fidelities. In this case, attributes of the form $F(IC,2)$ and $F(IC,3)$ are given twice the weight of $F(IC,1)$.

2. Extra weight for commoner species. In most applications it is undesirable to have numerous small groups of similar rare species segregating off at an early stage of the classification. This can be avoided by weighting the species according to their abundance, in which case the rarer species are unlikely to dominate the classification, and will tend to align themselves with commoner ones. The weighting used here is

$$w_j = \sum_i b_{ij} ;$$

where the matrix elements b_{ij} are (as in the definition of RAT) those that appear in the final tabulation.

3. Extra weight for larger groups and for the higher levels of the hierarchy in the sample classification. Weights of the attributes $F(IC,K)$ corresponding to the class IC are multiplied by

$$w_{IC} = 2^{-L/2} \sum_{i \in IC} \sum_j b_{ij} ,$$

where the summation $i \in IC$ is taken over those samples i that belong to group IC and L is the level of group IC. In effect, this means that each level down the hierarchy is given weight $\sqrt{2}$ less than that above, and that less attention is paid to fidelity to small groups than to large ones.

4.5 Subroutine PSEUDO

This constructs pseudospecies for the sample classification (cf. Section 1.7), and needs little comment except to note that the real species to which the pseudospecies are attached are identified by the vectors JNFLAG and INDPOT. Specifically, JNFLAG(INDPOT(J)) is the number of the real species corresponding to pseudospecies J. If J is a bottom-level pseudospecies (usually this will mean that it is a real species), then INDPOT(J) = J. INDPOT is so called because it is used to determine the indicator potential of a species for the indicator ordination. If either

INDPOT(J) or INDPOT(INDPOT(J)) is negative, then J is barred from being used in the indicator ordination. This device is used to ensure that no one real species is used more than once in the indicator ordination-- i.e. if Poa annua 2 is an indicator, then Poa annua 1, P. annua 3, etc., are barred from being indicators in the same dichotomy. There is no constraint preventing a species from being used as an indicator in many different dichotomies.

4.6 Subroutine CLASS

This subroutine makes the classification. It may be divided into the following activities.

1. A parameter IEND is set equal to $2^{**}(\text{LEVMAX}+2)$, which is bigger than the largest index number of any class that can result from divisions down to level LEVMAX. Any individuals that lack all attributes are identified at the outset and assigned to class IEND. Later on in the program, IEND is added to the index number of any class that is not to be divided further.
2. Various housekeeping activities are undertaken (subroutines DECODE, UPDATE, and RECODE). The name of subroutine UPDATE expresses its function well. It takes a newly made dichotomy, updates the class register ICLASS to allow for the new dichotomy, and looks for a new class to be divided. If no class remains to be divided, a switch ISTOP is set to 1. The subroutines RECODE and DECODE establish and remove a temporary numbering of the individuals and attributes for use during a particular dichotomy. The attributes are recoded so that they run consecutively from 1 to N, and the individuals are renumbered to run from 1 to M.
3. The rarer attributes are downweighted according to the following rules:
 - (a) All attributes with frequency greater than FRQLIM (= 0.2) are given unit weight;
 - (b) Attributes with frequency less than FRQLIM are given weights on a linear sliding scale in proportion to their frequency, starting with weight CWTMIN (= 0.01) for very rare attributes and rising to weight 1.0 for attributes with frequency FRQLIM. The purpose of this downweighting is to reduce the effect of the

- rarer attributes. It is possible to modify the degree of down-weighting by changing parameter values (see Section 4.10).
4. The individuals are ordinated by reciprocal averaging (Hill, 1973) in subroutine RA. Note that in addition to weights applied at Activity 3, the reciprocal averaging is weighted by weights RRWT for the individuals and CCWT for the attributes. In the sample classification the weights RRWT are set to 1.0, as are the weights CCWT in the standard analysis. However, CCWT may be varied by the user so as to be different for different cut levels (see Section 2.11). In the classification of the species, both RRWT and CCWT are set to non-unit values (see Section 4.4).
 5. The primary ordination is refined by application of subroutine POLISH (see Section 4.7).
 6. The order of the two groups resulting from the dichotomy is considered, and is arranged to fit in as well as possible with the rest of the table (for details see Section 4.8).
 7. The indicator ordination is defined in accordance with the ideas presented by Hill, Bunce & Shaw (1975)--subroutines ZONEUP, INDSCO, TOPIND, CODESC, TABLE, and FIND. If the maximum number of indicators is set to zero (cf. Section 2.6), then several of these subroutines are omitted. They are always omitted in the classification of the species.
 8. Information on the dichotomy is printed out (subroutine REPORT).

4.7 Refinement of the ordination

Refinement of the ordination is performed by subroutine POLISH. Essentially, this amounts to the application of a transfer algorithm (Gower, 1974; synonymously an iterative relocation algorithm--see Cormack, 1971), using a discriminant function to make the transfers. The primary ordination is converted to a dichotomy by dividing it at its center of gravity, and the frequencies of attributes on the positive and negative sides of the dichotomy are then used to construct a new ordination (and hence a new dichotomy). There is allowance both for borderline cases and for prior weighting of the individuals (for details see Section 4.10 and subroutine INDSCO), but this is ignored in the description given below.

The discriminant function is derived by adding together two ordinations. The first of these is additive, the second is gotten by taking a mean. The first ordination will generally be the dominant partner at the lower levels of the hierarchy; the second will be dominant at the higher levels.

The first ordination is gotten by adding together the "preference scores" of the commoner and more preferential species. A preference score of +1 is given to each attribute that is at least three times more frequent on the positive side as on the negative side, and which is commoner than a certain minimum. Common negative preferentials are treated similarly, and are given a score of -1. Attributes that are rarer, or which are less markedly preferential, are downweighted accordingly. In symbols, let an attribute J have frequency AYY on the positive side, and frequency AY on the negative. Define

$$\text{PREF} = (\text{AYY} - \text{AY}) / (\text{AYY} + \text{AY})$$

$$\text{FREQ} = \text{AYY} + \text{AY}$$

Then the score given to the attribute is

$$\text{Preference score} = (\text{FREQ}/\text{FRQLIM}) * (\text{PREF}/\text{PRLIM})^{**5} .$$

Note that this downweighting is only applied when either FREQ falls below the threshold FRQLIM (= 0.2; this is--unnecessarily--the same parameter as is used in Section 4.6, Activity 3), or when the absolute value of PREF falls below PRLIM (= (RATLIM - 1)/(RATLIM + 1), where RATLIM is the frequency-ratio limit, here set to 3.0). Clearly, rare attributes are mildly downweighted, but attributes with an insufficient preference ratio are greatly so.

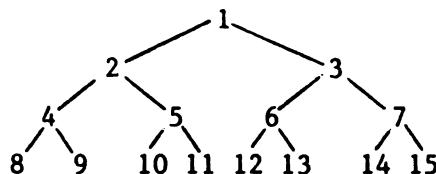
The first ordination is completed by standardizing the values so that the maximum absolute value is 1.

The other ordination in POLISH is gotten simply by taking the mean value of PREF/PRLIM (truncated to a maximum absolute value of 1) in the individual. This allows rarer attributes to express an opinion as well.

Finally, the refined ordination is gotten by adding these two. In practice, this process polarizes the ordination quite strongly, so that there are few borderline cases (cf. Section 3.3).

4.8 Ordering of dichotomies

Consider the following hierarchy, and in particular the question of how to order the pairs 10,11 and 14,15.



To get as smooth an arrangement as possible, we want 10 to be like 9, and 11 to be like 12; and we want 14 to be like 13, whereas 15 should be relatively unlike 13. One way to achieve this aim would be first to complete all the divisions at a particular level of the hierarchy, and then to try swivelling the dichotomies to see which ordering is the most satisfactory. An alternative approach, adopted here, is to try to see whether 10 or 11 more closely resembles 4, and whether 14 or 15 more closely resembles 6. This has the advantage that groups at level 4 are compared with those at level 3, so that the correct order can be determined at the time each dichotomy is formed. It has the added advantage that it compares the new groups with what are likely to be relatively large groups, so that order is determined more by general relationships than by particular accidents.

The basic ordering criterion in TWINSPLAN, then, is to take two groups $(2n)$ and $(2n+1)$ and to compare them with $(n-1)$ if n is odd and with $(n+1)$ if n is even. The comparison is made by setting up a discriminant function to distinguish $(2n)$ from $(2n+1)$, and then asking where this discriminant would place the comparison group (which is either $(n-1)$ or $(n+1)$, according as n is odd or even). This comparison is made by subroutine CLOSER.

This simple ordering criterion is complicated by the fact that the groups $(2n)$ and $(2n+1)$ are compared not only with one or other of $(n-1)$

and $(n+1)$, but also with one or other of $(n/2 - 1)$ and $(n/2 + 1)$. That is to say, not only is the local order considered, but also the relation to groups two stages back in the hierarchy. Let $YSCOR1$, $YSCOR2$ be the values of the discriminant function for comparisons one and two levels back in the hierarchy. Then the final discriminant is

$$YSCORE = YSCOR1 + W*YSCOR2 ,$$

where W is a weight function set to $+0.5$ or -0.5 (the sign depending on the remainder when n is divided by 4). This device pushes responsibility for the order even further back in the hierarchy, with the aim of achieving greater stability in the ordering. In particular, if the division at level 2 was effectively orthogonal to that at level 3, then the order dictated by relations at level 2 may be irrelevant to those at level 3; whereas relations at level 1 may still indicate a clear preference.

The discriminant function used in subroutine CLOSER is as follows.
Consider the ratio

$$PREF = ABS(AY0 - AY1)/(AY0 + AY1) ,$$

where $AY0$ is the frequency of the attribute on the negative side of the dichotomy, and $AY1$ is the frequency on the positive side. Now take

$$PREF = PREF/0.3 ,$$

and set all those values of $PREF$ that are bigger than 1 to 1. In this way we obtain a function that is 1 for all attributes that have a frequency ratio greater than about 1.9, and is less than 1 for lower values, dropping to 0 for complete indifference. (The value 1.9 arises because $0.3 \approx (1.9 - 1)/(1.9 + 1)$.) Now each attribute is scored for negativeness (PNEG), indifference (PIND), or positiveness (PPOS), as follows.

Case 1. The attribute is more frequent on the negative side.

Then $PNEG = PREF^{**4}$; $PIND = 1 - PNEG$; $PPOS = 0$.

Case 2. The attribute is more frequent on the positive side.

Then $PNEG = 0$; $PIND = 1 - PPOS$; $PPOS = PREF^{**4}$.

Using this scoring system, any individual or group of individuals can be assessed for negativeness, positiveness, or indifference; and in

particular, the positive and negative sides of the dichotomy can be so assessed. Let the mean scores per individual on the negative side of the dichotomy be XONEG, XOIND, and XOPOS; and let the mean scores per individual on the positive side be X1NEG, X1IND, and X1POS. The idea is now to assign scores that pay as little attention as possible to the indifferent attributes, while still requiring that the mean score on the negative side should be equal and opposite to that on the positive side. This is achieved by scoring -1 for negativeness, +1 for positiveness, and adjusting the score for indifference to make up the difference. In symbols, the score for indifference is

$$(XONEG + X1NEG - XOPOS - X1POS) / (XOIND + X1IND)$$
.

The only additional requirement is that the score for indifference should not exceed +1 or be less than -1. If it does exceed these limits, then it is reduced to +1 or -1 as appropriate.

There is a small additional feature that needs to be noticed, namely that in the discriminant function for ordering groups of samples, the various pseudospecies corresponding to a given real species are treated merely as additional occurrences of the one real species. (They are recognized as belonging to one another through the vector INDPOT; see Section 4.5.) For the species classification, on the other hand,

$$INDPOT(J) = J$$

for all values of J. Hence the differing degrees of fidelity are not recognized as belonging to one another.

The score for an individual is calculated by adding

$$PPOS - PNEG + (\text{Score for indifference} * PIND)$$

for all the attributes possessed by the individual. Finally, the score for a whole group is gotten by adding together the scores of all the individuals that it contains.

4.9 Subroutine RA

The algorithm used here for reciprocal averaging is a direct iteration algorithm comparable to that used by Hill (1973), but less susceptible

to round-off error. The essence of the algorithm is to represent the two-way averaging iteration approximately by a 4×4 symmetric tridiagonal matrix. The top eigenvector of the 4-dimensional approximation is then sought, and is used as a trial vector for the next iteration.

4.10 Significance of parameter values

Lines 53-101 of the program are devoted to setting the values of various parameters. Many of these parameters have already been mentioned, but are considered briefly again here to explain the result of varying them.

NUMTAB determines the number of samples printed out per page of the two-way table. This may need to be reduced for lineprinters with less than 124 characters per line (see Section 2.1).

RARE determines which species are to be included in the listing of preferential species at each dichotomy (see Section 3.3). At present this list is rather long (cf. Fig. 4), and it could be shortened by setting RARE to 0.4. In this case, only species and pseudospecies occurring in at least 40% of the samples on one or other side of the dichotomy would be listed.

FEEBLE is used for rejecting indicators that are too feeble. An initial dichotomy is made by dividing the refined ordination at its center. Attributes are then graded on the basis of their preference index, where

$$\text{Preference index} = \frac{\text{chance of occurring on positive side of dichotomy}}{\text{chance of occurring on negative side of dichotomy}}$$

For example, an attribute that occurs in 30% of samples on the positive side of the dichotomy and in 10% of samples on the negative side has a preference index of $0.3 - 0.1 = 0.2$. With the setting FEEBLE = 0.1 as supplied, such an attribute is then available as an indicator. Two points should be noted.

1. The absolute value of the preference index is used in applying FEEBLE, as the intention is to identify all good indicators, whether positive or negative.

2. Some allowance is made for borderline samples, giving them less weight than samples that are definitely positive or negative. For samples outside the borderline zone defined by CRCUT (see below), however, the criterion described above is applied exactly.

FRQLIM is a limit used for downweighting rare attributes, both in subroutine WEIGHT (see Section 4.6, Activity 3), and in subroutine POLISH (see Section 4.7). Lowering the value of FRQLIM will mean that fewer attributes are downweighted. This will make the analysis more inclined to separate small, deviant groups of samples at an early stage in the hierarchy.

TOL is the tolerance for deciding whether an eigenvector has been obtained in subroutine RA. Let x be a trial vector, standardized to unit length, and let Lx be the result of applying the two-way averaging of reciprocal averaging, also standardized to unit length. Then x is counted as an eigenvector when length of $(x - Lx)$ is less than TOL. There is little point in working to a high accuracy with the primary ordination, and the supplied value of TOL (= 0.003) appears to be quite adequate.

RATLIM is a limit used in subroutine POLISH (see Section 4.7). As supplied, it is set to 3.0, so that attributes that are less than three times as frequent on one side of the dichotomy as on the other are given less weight in refining the ordination.

REPLIM is a similar limit used in subroutine REPORT. As supplied it is set to 2.0, so that attributes that are less than twice as frequent on one side of the dichotomy as on the other are counted as non-preferentials (cf. Section 3.3). Raising the value of REPLIM will reduce the number of species and pseudospecies that are counted as preferentials, but will not affect the number of these that are printed out in all (unlike RARE; see comments above).

IREWT is the number of times that subroutine POLISH is applied per dichotomy. The value of 2 used here was found to be the best empirically,

but if the method of refining the ordination is changed, there is no reason why another value of IREW_T might not be better.

ICWEXP is power of downweighting for species not up to limit FRQLIM. Rarer species can be more heavily downweighted by raising the value of ICWEXP (see lines 1530 and 1751 of source code).

IPREXP is a similar exponent for preferentiality in POLISH (see Section 4.7). In fact the power of downweighting used in POLISH is one greater than IPREXP, which explains why an exponent **5 is mentioned in Section 4.7, whereas IPREXP is set to 4.

CWTMIN is a minimum weight to be given to attributes in subroutine WEIGHT. See Section 4.6, Activity 3.

CRCUT determines the length of the zone used in defining borderline cases for the purposes of refining the ordination. (It was mentioned in Section 4.7 that borderline cases are given special treatment.) As supplied, CRCUT is set to 0.2. This means that samples in the middle 20% of the range of the ordination are downweighted in determining preferentiality, from zero weight at the center of the ordination, rising on a linear scale to unit weight at distances 40% and 60% from the end.

CRLONG determines the length of the critical zone, for the purposes of defining misclassifications, etc. As supplied, CRLONG is 0.2. This is why the critical zone in Fig. 4 (Section 3.3) is 20% of the total length of the ordination. If it is felt that too many samples are being classified as "borderline cases" then this can be corrected by lowering the value of CRLONG.

MZCRIT, MZIND, and MZOUT are best interpreted by reference to Fig. 4. MZCRIT is the number of segments in the critical zone (here set to 8). MZIND is the number of segments in the zone of indifference (here set to 4), and MZOUT is the number of segments on each side of the diagram outside the critical zone (here set to 4 also). One way to reduce the number of samples that are classified as "borderline cases" would be to

reduce the value of MZIND from 4 to 2, halving the length of the zone of indifference, so that it drops from 10% to 5% of the total length of the ordination.

WTHIGH, SPE1, SPE2, and SPE3 are parameters used in converting the data matrix for the species classification (see Section 4.4 for details).

4.11 Outlook

TWINSPAN is long and rather complicated. It would be a better program if it were simpler, concentrating more directly on users' particular requirements. For example, many users will be more interested in getting a good two-way table than in the niceties of the indicator ordination. The indicator ordination requires several hundred lines of programming code, and could perhaps be omitted in a future edition.

Another feature that could perhaps be improved is the process of refining the ordination. The discriminant function used to order the dichotomies is more elegant than that used in subroutine POLISH, and could well be used also to refine the ordination. This modification, also, would save programming code.

It would be desirable to look more than two levels back in determining the order of dichotomies (cf. Section 4.8). Thus, if two small and aberrant groups are successively split off a larger group, and if the larger group is split again, then the ordering of the two groups resulting from the third dichotomy will be determined by their similarity to the two small groups split off previously, and not to the general trend of the two-way table.

For the present, however, the program is being allowed to stand. A more careful study of discriminant functions needs to be made before any substantial progress is possible.

5. ACKNOWLEDGMENTS

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Dr. R. H. Whittaker invited me to visit Cornell, and encouraged me to pursue the possibilities of numerical classification. I am indebted to him, to H. G. Gauch, Jr., and to Dr. P. L. Marks for the provision of test data and much helpful criticism.

Finally I should like to thank S. B. Singer for his expert advice about the Cornell computer and for writing several utility programs which have been indispensable in preparing TWINSPAN.

6. REFERENCES

- Cormack, R. M. (1971). A review of classification. J. R. Statist. Soc., Series A, 134, 321-367.
- Gauch, H. G. (1977). ORDIFLEX -- A flexible computer program for four ordination techniques: weighted averages, polar ordination, principal components analysis, and reciprocal averaging, Release B. Ecology and Systematics, Cornell University, Ithaca, New York 14850.
- Gower, J. C. (1974). Maximal predictive classification. Biometrics 30, 643-654.
- Hill, M. O. (1973). Reciprocal averaging: an eigenvector method of ordination. J. Ecol. 61, 237-249.
- Hill, M. O. (1977). Use of simple discriminant functions to classify quantitative phytosociological data. First International Symposium on Data Analysis and Informatics, Versailles 7-9 Sept. 1977 (Ed. by E. Diday, L. Lebart, J. P. Pages, & R. Tomassone), Vol. 1, pp. 181-199. Institut de Recherche d'Informatique et d'Automatique, Domaine de Voluceau, Rocquencourt, B.P. 105, 78150 le Chesnay, France.
- Hill, M. O. (1979). DECORANA -- A FORTRAN program for detrended correspondence analysis and reciprocal averaging. Ecology and Systematics, Cornell University, Ithaca, New York 14850.
- Hill, M. O., Bunce, R. G. H., & Shaw, M. W. (1975). Indicator species analysis, a divisive polythetic method of classification, and its application to a survey of native pinewoods in Scotland. J. Ecol. 63, 597-613.
- Maarel, E. van der, Janssen, J. G. M., & Louppen, J. M. W. (1978). TABORD, a program for structuring phytosociological tables. Vegetatio 38, 143-156.
- Mueller-Dombois, D. & Ellenberg, H. (1974). Aims and Methods of Vegetation Ecology. Wiley, New York.
- Singer, S. B. & Gauch, H. G. (1979). CONDENSE -- Convert data matrices from any ORDIFLEX format into a condensed format by samples. Ecology and Systematics, Cornell University, Ithaca, New York 14850.
- Westhoff, V. & Maarel, E. van der. (1973). The Braun-Blanquet approach. Ordination and Classification of Communities (Ed. by R. H. Whittaker), pp. 617-726. Junk, The Hague.

7. LISTING OF SOURCE CODE

C CORNELL ECOLOGY PROGRAM TWINSPLAN - WRITTEN BY M.O. HILL, JULY 1979. TWI 1
C SOURCE CODE AND ACCOMPANYING DOCUMENTATION ARE AVAILABLE FROM TWI 2
C HUGH G. GAUCH, JR., ECOLOGY AND SYSTEMATICS, CORNELL UNIVERSITY TWI 3
C ITHACA, NEW YORK 14850. TWI 4
C TWI 5
C PERFORMS TWO-WAY INDICATOR SPECIES ANALYSIS, ARRANGING SPECIES AND TWI 6
C SAMPLES IN A TWO-WAY TABLE AT THE END. TWI 7
C TWI 8
C DIMENSIONED FOR 500 SAMPLES, 800 SPECIES, 14500 NON-ZERO ITEMS IN TWI 9
C DATA-MATRIX. IDAT IS DIMENSIONED 30000=2*(14500+500). JDAT IS TWI 10
C DIMENSIONED WITH HALF THE LENGTH OF IDAT, AND THE FRONT OF IT IS TWI 11
C EQUIVALENCED TO THE MIDDLE OF IDAT. TWI 12
C TWI 13
C INPUT DEVICES ARE 3 (INPUT PARAMETERS - SHOULD BE A COMPUTER TWI 14
C TERMINAL IF POSSIBLE), 5 (DATA-MATRIX). TWI 15
C TWI 16
C OUTPUT DEVICES ARE 4 (MONITORING - SHOULD BE CONSOLE OF COMPUTER TWI 17
C TERMINAL IF POSSIBLE; OTHERWISE THIS OUTPUT IS DISPENSABLE); TWI 18
C 7 (COPY OF SOLUTION IN MACHINE-READABLE FORM - OPTIONAL AND ONLY TWI 19
C SUPPLIED IF REQUESTED); AND LINEPRINTER USING FORTRAN WORD PRINT. TWI 20
C TWI 21
INTEGER IDAT(30000),JDAT(15000) TWI 22
INTEGER IX(500),ICLASS(500),IIROW(500),IADDR(500) TWI 23
INTEGER INFLAG(500),INAME1(500),INAME2(500),IZONE(500) TWI 24
INTEGER IX1(500),IX2(500),IX3(500) TWI 25
INTEGER IY(800),JJCOL(800),INDPOT(800),INDORD(800) TWI 26
INTEGER JNFLAG(800),JNAME1(800),JNAME2(800),JNAM(800) TWI 27
INTEGER IY0(800),IY1(800),IY2(800) TWI 28
INTEGER ITEM(200) TWI 29
INTEGER LCUT(10),LWGT(10),LIND(10) TWI 30
INTEGER INDSIG(20),IPICT(104,25) TWI 31
REAL X(500),XX(500),RTOT(500),RRWT(500),ROWWGT(500) TWI 32
REAL X3(500),X4(500),X5(500) TWI 33
REAL Y(800),YY(800),CTOT(800),CCWT(800),COLWGT(800) TWI 34
REAL TOT(511),TOTJ(511) TWI 35
REAL TITLE(20) TWI 36
EQUIVALENCE (IDAT(15001),JDAT(1)) TWI 37
EQUIVALENCE (IX1(1),X3(1)),(IX2(1),X4(1)),(IX3(1),X5(1)) TWI 38
EQUIVALENCE (IY0(1),COLWGT(1)),(IY1(1),CTOT(1)),(IY2(1),CCWT(1)) TWI 39
COMMON/PICT/MZ,MS,I2D,IIZD,ISD,ISHIFT,MZIND,MZCRIT, TWI 40
1MZOUT,NIND TWI 41
COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM TWI 42
COMMON/SWITCH/IFAIL,IDIAGR,ISTOP,IREWLT TWI 43
COMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX TWI 44
CCMON/ARBS/CWTMIN,CRLONG,CRCUT TWI 45
COMMON/WORK/ITEM TWI 46
CCMON/SECOND/ISEC TWI 47
COMMON/SPESPE/WTHIGH,SPE1,SPE2,SPE3 TWI 48
NDAT=30000 TWI 49
MMAX=500 TWI 50
NMAX=800 TWI 51
C THIS IS THE END OF DECLARATIONS AND SETTING OF DIMENSIONS. TWI 52
NUMTAB=100 TWI 53
C NUMTAB IS MAXIMUM NUMBER OF SAMPLES PER PAGE IN FINAL TABULATION TWI 54
RARE=0.2 TWI 55

C RARE IS LIMIT OF RARITY FOR USE IN LISTING PREFERENTIAL SPECIES	TWI	56
FEEBLE=0.1	TWI	57
C INDICATORS WITH PREFERENCE INDEX LESS THAN FEEBLE ARE REJECTED	TWI	58
FRQLIM=0.2	TWI	59
C FRQLIM IS LIMIT OF RARITY FOR DOWNWEIGHTING. SPECIES OCCURRING IN	TWI	60
C A PROPORTION OF SAMPLES LESS THAN FRQLIM ARE DOWNWEIGHTED IN THE	TWI	61
C ORDINATION.	TWI	62
TOL=0.003	TWI	63
C TOL IS TOLERANCE FOR DECIDING WHETHER A TRIAL VECTOR IS EIGENVECTOR	TWI	64
RATLIM=3.0	TWI	65
C RATLIM IS THE FREQUENCY-RATIO LIMIT USED IN SUBROUTINE POLISH	TWI	66
REPLIM=2.0	TWI	67
C REPLIM IS THE FREQUENCY-RATIO LIMIT USED IN SUBROUTINE REPORT	TWI	68
IREWT=2	TWI	69
C IREWDT IS NUMBER OF TIMES SUBROUTINE POLISH TO BE CALLED PER DIVISION	TWI	70
ICWEXP=1	TWI	71
C ICWEXP IS POWER (I.E. EXPONENT) OF DOWNWEIGHTING FOR SPECIES NOT UP	TWI	72
C TO RARITY LIMIT FRQLIM.	TWI	73
IPREXP=4	TWI	74
C IPREXP IS POWER (I.E. EXPONENT) OF DOWNWEIGHTING USED FOR	TWI	75
C PREFERENTIALITY IN SUBROUTINE POLISH.	TWI	76
CWTMIN=0.01	TWI	77
C CWTMIN IS SMALLEST MULTIPLIER THAN CAN BE USED IN DOWNWEIGHTING.	TWI	78
C NO SPECIES CAN BE DOWNWEIGHTED BY A FACTOR MORE THAN CWTMIN.	TWI	79
CRCUT=0.2	TWI	80
C CRCUT IS LENGTH OF CRITICAL ZONE USED IN SUBROUTINE POLISH	TWI	81
CRLONG=0.2	TWI	82
C CRLONG IS PROPORTION OF TOTAL LENGTH OF ORDINATION OCCUPIED BY	TWI	83
C CRITICAL ZONE, FOR PURPOSES OF DEFINING MISCLASSIFICATIONS ETC.	TWI	84
MZCRIT=8	TWI	85
C MZCRIT IS NUMBER OF SEGMENTS IN CRITICAL ZONE.	TWI	86
MZIND=4	TWI	87
C MZIND IS THE NUMBER OF SEGMENTS IN THE ZONE OF INDIFFERENCE	TWI	88
MZOUT=4	TWI	89
C MZOUT IS THE NUMBER OF SEGMENTS TO BE RECOGNIZED OUTSIDE CRITICAL	TWI	90
C ZONE. HENCE NUMBER OF ZONES IN DIAGRAM OF DIVISION (AVAILABLE	TWI	91
C AS AN OPTION) IS 2*MZOUT+MZCRIT.	TWI	92
WTHIGH=2.0	TWI	93
C WTHIGH IS WEIGHTING GIVEN TO CHARACTERISTICS OF SPECIES THAT	TWI	94
C CORRESPOND TO HIGHER LEVELS OF FIDELITY - I.E. TO THOSE	TWI	95
C CHARACTERISTICS CORRESPONDING TO FIDELITY LEVELS SPE2,SPE3.	TWI	96
SPE1=0.8	TWI	97
SPE2=2.0	TWI	98
SPE3=6.0	TWI	99
C SPEC1,2,3 ARE THRESHOLDS OF FIDELITY USED IN DEFINING THE	TWI	100
C ATTRIBUTES OF SPECIES FOR THE SPECIES-CLASSIFICATION.	TWI	101
CALL QUIKINI(MMAX,NMAX,NDAT,MM,NSPEC,INAME1,	TWI	102
1INAME2,JNAME1,JNAME2,1DAT,IADDR,MDAT,TITLE,INFLAG)	TWI	103
CALL ENTER(NSPEC,NL,L CUT,MMIN,MIND,JMAX,LEVMAX,	TWI	104
1IDIAGR,IPUNCH,LWGT,LIND,INDORD)	TWI	105
MZ=MZCRIT+2*MZOUT	TWI	106
MMZ=MZ+4	TWI	107
MMS=MIND+4	TWI	108
DO 50 J=1,NSPEC	TWI	109
50 JNFLAG(J)=J	TWI	110

```
CALL PSEUDO(MM,NN,NMAX,NL,NDAT,NSPEC,IDAT,LCUT,           TWI 111
1JNFLAG,JNAME1,JNAME2,JNAM,INDPOT,IY)                   TWI 112
IF(MMIN.GT.MM) STOP                                     TWI 113
IF(LEVMAX.EQ.0) STOP                                     TWI 114
IF(NSPEC.GT.MMAX) PRINT 1000,MMAX                      TWI 115
IF(NSPEC.GT.MMAX) WRITE(4,1000) MMAX                  TWI 116
1000 FORMAT(//1X,'*** BEWARE *** NUMBER OF SPECIES EXCEEDS',   TWI 117
1' MMAX (=' ,I4,') .'/                                TWI 118
21X,'NO SPECIES CLASSIFICATION IS POSSIBLE.')        TWI 119
DO 60 II=1,MM                                         TWI 120
60 RRWT(II)=1.0                                      TWI 121
DO 165 JJ=1,NN                                         TWI 122
IL=JNAM(JJ)                                           TWI 123
165 CCWT(JJ)=FLOAT(LWGT(IL))+1.0E-5                 TWI 124
DO 170 JJ=1,NN                                         TWI 125
J=IABS(INDPOT(JJ))                                    TWI 126
JJJ=JNFLAG(J)                                         TWI 127
170 IF(INDORD(JJJ).EQ.0) INDPOT(JJ)=-J               TWI 128
PRINT 1011                                            TWI 129
1011 FORMAT('1')                                       TWI 130
ISEC=1                                               TWI 131
IF(IPUNCH.EQ.1) WRITE(7,1200) (TITLE(IT),IT=1,20)      TWI 132
1200 FORMAT(1X,19A4,A3//1X,'SAMPLE CLASSIFICATION')
CALL CLASS(MM,NN,NDAT,MIND,MMZ,MMS,IX,ICLASS,IIROW,IADDR,
1INDPOT,INDORD,IZONE,IY,JCOL, IDAT,INDSIG,IPICT,X,XX,RTOT,
2RRWT,ROWWGT,Y,YY,CTOT,CCWT,COLWGT,INAME1,INAME2,JNAME1,
3JNAME2,JNAM,X3,X4,X5,LIND,INFLAG,IPUNCH)
ID=IADDR(MM)                                         TWI 137
415 IF(IDAT(ID).EQ.-1) GOTO 420                      TWI 138
ID=ID+1                                              TWI 139
GOTO 415                                             TWI 140
420 NDAT=NDAT/2                                       TWI 141
IF (ID.GT.NDAT) GOTO 999                           TWI 142
IID=NDAT                                         TWI 143
DO 422 J=1,NSPEC                                     TWI 144
422 Y(J)=0.0                                         TWI 145
425 IDAT(IID)=IDAT(ID)                            TWI 146
IID=IID-1                                         TWI 147
ID=ID-1                                           TWI 148
IF(ID.EQ.0) GOTO 426                            TWI 149
GOTO 425                                         TWI 150
426 JJJ=0                                           TWI 151
ICMAX=0                                         TWI 152
DO 431 II=1,MM                                     TWI 153
IC=ICLASS(II)                                    TWI 154
427 IF(IC*3.LE.NMAX) GOTO 428                     TWI 155
IC=IC/2                                         TWI 156
GOTO 427                                         TWI 157
428 ICLASS(II)=IC                                 TWI 158
431 IF(IC.GT.ICMAX) ICMAX=IC                    TWI 159
IF (NSPEC.GT.MMAX) GOTO 999                      TWI 160
DO 432 IC=1,ICMAX                               TWI 161
432 TOT(IC)=0.0                                  TWI 162
JD=0                                              TWI 163
IBIG=10000                                         TWI 164
                                         TWI 165
```

```
DO 440 II=1,MM          TWI 166
  IC=ICLASS(II)          TWI 167
433 IID=IID+1           TWI 168
  JJ=IDAT(IID)          TWI 169
  IF(JJ.EQ.-1) GOTO 437  TWI 170
  J=IABS(INDPOT(JJ))    TWI 171
  JD=JD+1               TWI 172
  IF(JD.GT.NDAT) GOTO 999 TWI 173
  JDAT(JD)=J*IBIG+IC    TWI 174
  TOT(IC)=TOT(IC)+1.0   TWI 175
  IF(J.EQ.JJJ) GOTO 435  TWI 176
  JJJ=J                  TWI 177
  Y(J)=Y(J)+CCWT(J)*RRWT(II) TWI 178
  ID=ID+1               TWI 179
  ICAT(ID)=J             TWI 180
  IC=ID+1               TWI 181
  IDAT(ID)=1             TWI 182
  IF(ID.GE.IID) GOTO 999 TWI 183
  GOTO 433               TWI 184
435 IDAT(ID)=IDAT(ID)+1 TWI 185
  GOTO 433               TWI 186
437 ID=ID+1               TWI 187
  IDAT(ID)=JJ             TWI 188
  JJJ=0                  TWI 189
440 CONTINUE              TWI 190
  DO 441 J=1,NSPEC        TWI 191
441 RRWT(J)=Y(J)+0.00001 TWI 192
C WE NOW WORK ON SPECIES CLASSIFICATION      TWI 193
  IIBIG=IBIG-1            TWI 194
  DO 442 J=1,NSPEC        TWI 195
  JD=JD+1                TWI 196
  IF(JD.GT.NDAT) GOTO 999 TWI 197
442 JDAT(JD)=J*IBIG+IIBIG  TWI 198
  CALL ISORT(JDAT,JD)     TWI 199
C MOVE MATRIX UP TO BACK OF ARRAY            TWI 200
  JJD=NDAT                TWI 201
445 JDAT(JJD)=JDAT(JD)      TWI 202
  JJD=JJD-1                TWI 203
  JD=JD-1                  TWI 204
  IF(JD.EQ.0) GOTO 450      TWI 205
  GOTO 445                TWI 206
C WE NOW RECONSTITUTE THE MATRIX AND PROCEED TO ACCUMULATE  TWI 207
C THE TOTALS FOR THE SAMPLE CATEGORIES        TWI 208
450 IC=ICMAX                TWI 209
455 ICC=IC/2                 TWI 210
  TOT(ICC)=TOT(ICC)+TOT(IC)  TWI 211
  IC=IC-1                  TWI 212
  IF(IC.GT.1) GOTO 455      TWI 213
  DO 470 JJJ=1,NSPEC        TWI 214
  DO 456 IC=1,ICMAX         TWI 215
456 TOTJ(IC)=0.0             TWI 216
458 JJD=JJD+1               TWI 217
  JJ=JDAT(JJD)              TWI 218
  J=JJ/IBIG                 TWI 219
  IC=JJ-J*IBIG               TWI 220

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```

```
IF(IC.EQ.IIBIG) GOTO 460          TWI 221
TOTJ(IC)=TOTJ(IC)+1.0            TWI 222
GOTO 458                        TWI 223
460 IC=ICMAX                      TWI 224
462 IIC=IC/2                       TWI 225
TOTJ(IIC)=TOTJ(IIC)+TOTJ(IC)      TWI 226
IC=IC-1                          TWI 227
IF(IC.GT.1) GOTO 462              TWI 228
DO 464 IC=1,ICMAX                TWI 229
RAT=TOTJ(IC)*(TOT(1)-TOT(IC))/(TOT(IC)*(TOTJ(1)-
1TOTJ(IC))+1.0E-7)                TWI 230
IF (RAT.LT.SPE1) GOTO 464          TWI 231
JD=JD+1                          TWI 232
JCAT(JD)=IC*3-2                  TWI 233
IF (RAT.LT.SPE2) GOTO 464          TWI 234
JD=JD+1                          TWI 235
JDAT(JD)=IC*3-1                  TWI 236
IF (RAT.LT.SPE3) GOTO 464          TWI 237
JD=JD+1                          TWI 238
JCAT(JD)=IC*3                  TWI 239
464 CONTINUE                      TWI 240
JD=JD+1                          TWI 241
IF(JD.GE.JJD) GOTO 999             TWI 242
JDAT(JD)=-1                      TWI 243
TWI 244
470 CCNTINUE                      TWI 245
C JDAT NOW CONTAINS DESIRED SPECIES INFORMATION. IT IS NOW A    TWI 246
C MATTER OF FIXING UP WEIGHTS AND CLASSIFYING SPECIES           TWI 247
DO 480 IC=1,ICMAX                TWI 248
WEIGHT=1.0                      TWI 249
IIC=IC                          TWI 250
473 IIC=IIC*2                     TWI 251
IF(IIC.GT.1023) GOTO 476          TWI 252
WEIGHT=WEIGHT*1.414               TWI 253
GOTO 473                        TWI 254
476 WEIGHT=WEIGHT*TOT(IC)         TWI 255
IIC=3*IC                         TWI 256
CCWT(IIC-2)=WEIGHT               TWI 257
INDPOT(IIC-2)=IIC-2               TWI 258
CCWT(IIC-1)=WEIGHT*WTHIGH        TWI 259
INDPOT(IIC-1)=IIC-1               TWI 260
CCWT(IIC)=WEIGHT*WTHIGH          TWI 261
INDPOT(IIC)=IIC                  TWI 262
480 CCNTINUE                      TWI 263
PRINT 1011                        TWI 264
ISEC=2                           TWI 265
IF(IPUNCH.EQ.1) WRITE(7,1300)      TWI 266
1300 FORMAT(//1X,'SPECIES CLASSIFICATION')      TWI 267
CALL CLASS(NSPEC,3*ICMAX,NDAT,0,MMZ,MMS,      TWI 268
1IX,JNAM,IIROW,IADDR,INDPOT,INDORD,IZONE,IY,JCOL,JDAT,      TWI 269
2INDSIG,IPICT,X,XX,RTOT,RRWT,ROWWGT,Y,YY,CTOT,CCWT,COLWGT,  TWI 270
3JNAME1,JNAME2,I NAME1,I NAME2,I NAME1,X3,X4,X5,LIND,JNFLAG,IPUNCH)  TWI 271
CALL CLORD(MM,LEVMAX,ICLASS,IX)          TWI 272
CALL CLORD(NSPEC,LEVMAX,JNAM,IY)          TWI 273
C WE HAVE NOW ESSENTIALLY GOT SAMPLES AND SPECIES ORDERED, BUT   TWI 274
C THE SPECIES ORDER MAY NEED TO BE REVERSED TO GET THINGS ON THE  TWI 275
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C POSITIVE DIAGONAL. ALSO WE ACCUMULATE SPECIES TOTALS IN INDPOT.      TWI 276
DO 500 II=1,MM              TWI 277
I=IX(II)                   TWI 278
IX1(II)=INFLAG(I)          TWI 279
IX2(II)=INAME1(I)          TWI 280
IX3(II)=INAME2(I)          TWI 281
500 X(I)=FLOAT(II)          TWI 282
DO 510 JJ=1,NSPEC          TWI 283
INDPOT(JJ)=0                TWI 284
J=IY(JJ)                   TWI 285
510 Y(J)=FLOAT(JJ)          TWI 286
TOT1=0.0                   TWI 287
TOTX=0.0                   TWI 288
TOTY=0.0                   TWI 289
TOTXY=0.0                  TWI 290
ID=0                       TWI 291
DO 520 II=1,MM              TWI 292
AX=X(II)                   TWI 293
512 ID=ID+1                 TWI 294
J=IDAT(ID)                 TWI 295
IF(J.EQ.-1) GOTO 520        TWI 296
AY=Y(J)                     TWI 297
ID=ID+1                   TWI 298
JQ=IDAT(ID)                TWI 299
INDPOT(J)=INDPOT(J)+JQ     TWI 300
Q=FLOAT(JQ)                 TWI 301
TOT1=TOT1+Q                 TWI 302
TOTX=TOTX+Q*AX              TWI 303
TOTY=TOTY+Q*AY              TWI 304
TOTXY=TOTXY+Q*AX*AY         TWI 305
GOTO 512                   TWI 306
520 LCNTINUE                TWI 307
TOTXY=TOTXY-TOTX*TOTY/TOT1  TWI 308
IF(TOTXY.GT.0.0) GOTO 537   TWI 309
C IN THIS CASE TOTXY IS NEGATIVE, I.E. IX AND IY ARE ORDERS      TWI 310
C THAT ARE NEGATIVELY CORRELATED, AND IY MUST BE REVERSED          TWI 311
DO 530 J=1,NSPEC            TWI 312
530 JJCOL(J)=IY(NSPEC-J+1)  TWI 313
DO 535 J=1,NSPEC            TWI 314
535 IY(J)=JJCOL(J)          TWI 315
537 JMIN=1                  TWI 316
DO 540 J=1,NSPEC            TWI 317
JJ=IY(J)                   TWI 318
IF(INDPOT(JJ).EQ.0) JMIN=JMIN+1  TWI 319
IY0(J)=JNFLAG(JJ)           TWI 320
IY1(J)=JNAME1(JJ)           TWI 321
540 IY2(J)=JNAME2(JJ)       TWI 322
C WE WANT TO PRINT OUT ONLY JMAX COMMONEST SPECIES      TWI 323
PRINT 1500                 TWI 324
1500 FORMAT(//1X,'ORDER OF SPECIES INCLUDING RARER ONES')    TWI 325
PRINT 1501,(IY0(J),IY1(J),IY2(J),J=JMIN,NSPEC)             TWI 326
1501 FORMAT((1X,7(I4,1X,A4,1X,A4,'|'),I4,1X,A4,1X,A4))    TWI 327
PRINT 1502                 TWI 328
1502 FORMAT(//1X,'ORDER OF SAMPLES')                         TWI 329
PRINT 1503,(IX1(II),IX2(II),IX3(II),II=1,MM)               TWI 330

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```
1503 FORMAT((1X,7(I4,1X,2A4,' |'),I4,1X,2A4))           TWI 331
      IF(JMAX.EQ.0) STOP                                TWI 332
      IBIG=5000                                         TWI 333
      DO 542 J=1,NSPEC                                 TWI 334
      JJCOL(J)=IBIG*NSPEC                            TWI 335
      Y(J)=1.0                                         TWI 336
 542 INDPOT(J)=-IBIG*INDPOT(J)-IBIG+J               TWI 337
      CALL ISORT(INDPOT,NSPEC)                         TWI 338
      IF(JMAX.GT.NSPEC-JMIN+1) JMAX=NSPEC-JMIN+1       TWI 339
      DO 546 JJ=1,JMAX                               TWI 340
      J=-INDPOT(JJ)                                    TWI 341
      J=(J/IBIG)*IBIG-J+IBIG                         TWI 342
 546 JJCOL(J)=0                                     TWI 343
      DO 547 JJ=1,NSPEC                             TWI 344
      J=IY(JJ)                                       TWI 345
 547 IY(JJ)=JJ*IBIG+J+JJCOL(J)                      TWI 346
      CALL ISORT(IY,NSPEC)                           TWI 347
      DO 548 JJ=1,NSPEC                             TWI 348
      J=IY(JJ)                                       TWI 349
 548 IY(JJ)=J-(J/IBIG)*IBIG                        TWI 350
 550 CALL JORDER(MM,NSPEC,NDAT,IDAT,
     1IADDR,IY,JJCOL,INDPOT)                         TWI 351
      ITIMES=MM/NUMTAB                                TWI 352
      IF(MM.NE.ITIMES*NUMTAB)ITIMES=ITIMES+1          TWI 353
      DO 600 IT=1,ITIMES                            TWI 354
      IMIN=(IT-1)*NUMTAB+1                           TWI 355
      IMAX=IT*NUMTAB                                  TWI 356
      IF(IMAX.GT.MM) IMAX=MM                         TWI 357
 600 CALL TABOUT(MM,NSPEC,IMIN,IMAX,1,JMAX,NDAT,IDAT,
     1IADDR,IX,IY,INFLAG,JNFLAG,JNAME1,JNAME2,ICLASS,JNAM) TWI 358
      STOP                                            TWI 359
 999 PRINT 1999                                      TWI 360
      WRITE(4,1999)                                    TWI 361
 1999 FORMAT(1X,'** STOP ** INSUFFICIENT ARRAY SPACE TO CONTINUE') TWI 362
      STOP                                            TWI 363
      END                                             TWI 364
C
C      SUBROUTINE QUIKIN(MMAX,NMAX,NDAT,MM,N,IAME1,
C      1NAME2,JNAME1,JNAME2,IDAT,IBEGIN,NID,TITLE,INFLAG)
C READS IN DATA MATRIX BY SAMPLES, FOLLOWED BY ROW AND CCOLUMN NAMES IN TWI 365
C SPECIFIED FORMAT. THE SAMPLES MUST BE IN ASCENDING NUMERICAL TWI 366
C ORDER. SECOND HALF OF SUBROUTINE CONSISTS OF AN OPTION TO DELETE TWI 367
C UNWANTED SAMPLES FROM THE ANALYSIS.                          TWI 368
      INTEGER IAME1(MMAX),NAME2(MMAX),JNAME1(NMAX),
     1JNAME2(NMAX),IDAT(NDAT)                         TWI 369
      INTEGER IBEGIN(MMAX),INFLAG(MMAX)                TWI 370
      REAL AITEM(100),FMT(15),TITLE(20)                TWI 371
      INTEGER ITEM(200)                                TWI 372
      CCOMMON/WORK/ITEM                                TWI 373
      PRINT 1002                                       TWI 374
      WRITE(4,1002)                                    TWI 375
      READ(5,1000) (TITLE(IT),IT=1,20)                TWI 376
 1000 FCRMAT(20A4)                                   TWI 377
      PRINT 2000,(TITLE(IT),IT=1,20)                  TWI 378
      WRITE(4,2001) (TITLE(IT),IT=1,20)                TWI 379
                                         TWI 380
                                         TWI 381
                                         TWI 382
                                         TWI 383
                                         TWI 384
                                         TWI 385
```

DATA PARADIGM

2001	FORMAT(//1X,19A4,A3//)	TWI	386
2000	FORMAT(//1X,20A4//)	TWI	387
	READ(5,1001) (FMT(IT),IT=1,15),NITEM	TWI	388
1001	FORMAT(15A4,8X,I2)	TWI	389
1002	FORMAT(//1X,'READING DATA MATRIX FROM DEVICE 5')	TWI	390
	DO 10 II=1,MMAX	TWI	391
10	INFLAG(II)=0	TWI	392
	ID=0	TWI	393
	N=0	TWI	394
	III=1	TWI	395
	ITEM(NITEM+1)=0	TWI	396
40	READ(5,FMT) II,(ITEM(IT),AITEM(IT),IT=1,NITEM)	TWI	397
	IF(II.EQ.III) GOTO 50	TWI	398
	ID=ID+1	TWI	399
	IDAT(ID)=-1	TWI	400
	IF(II.EQ.0) GOTO 100	TWI	401
	IBEGIN(II)=ID+1	TWI	402
	IF(II.LT.III) GOTO 999	TWI	403
	III=II	TWI	404
50	IT=0	TWI	405
55	IT=IT+1	TWI	406
	J=ITEM(IT)	TWI	407
	IF(J) 997,40,56	TWI	408
56	IF(J.GT.N) N=J	TWI	409
	INFLAG(II)=1	TWI	410
	ID=ID+1	TWI	411
	IDAT(ID)=J	TWI	412
	AIJ=AITEM(IT)	TWI	413
	IF(AIJ.LT.0.0) GOTO 997	TWI	414
	ID=ID+1	TWI	415
	IDAT(ID)=IFIX(AIJ*1000.0+0.5)	TWI	416
	IF(ID.GT.NDAT-3) GOTO 996	TWI	417
	GOTO 55	TWI	418
100	MM=III	TWI	419
	IF(MM.GT.MMAX) GOTO 995	TWI	420
	IF(N.GT.NMAX) GOTO 994	TWI	421
	NID=ID	TWI	422
	IBEGIN(1)=1	TWI	423
C	WE NOW READ THE SPECIES NAMES FOLLOWED BY THE SAMPLE NAMES	TWI	424
	READ (5,1100) (JNAME1(J),JNAME2(J),J=1,N)	TWI	425
	READ (5,1100) (INAME1(I),INAME2(I),I=1,MM)	TWI	426
1100	FCRFORMAT(20A4)	TWI	427
	WRITE(4,1101) MM,N,NID	TWI	428
	PRINT 1101,MM,N,NID	TWI	429
1101	FORMAT('ONUMBER OF SAMPLES',I6/	TWI	430
	1'ONUMBER OF SPECIES',I6/	TWI	431
	2'OLENGTH OF RAW DATA ARRAY',I6)	TWI	432
	IID=120	TWI	433
	IF(NID.LT.IID) IID=NID	TWI	434
	PRINT 1102,(IDAT(ID),ID=1,IID)	TWI	435
1102	FORMAT(1X,12I10)	TWI	436
	IF(NID.LE.120) GOTO 150	TWI	437
	IF(NID.LE.240) GOTO 140	TWI	438
	PRINT 1103	TWI	439
1103	FORMAT(1X,6X,28('...'),'..')	TWI	440

SEE ERRA1:

140	IID=NID-119	TWI	441
	IF(IID.LT.121) IID=121	TWI	442
	PRINT 1102,(IDAT(ID),ID=IID,NID)	TWI	443
150	PRINT 1104	TWI	444
1104	FORMAT('OSPECIES NAMES')	TWI	445
	PRINT 1105,(J,JNAME1(J),JNAME2(J),J=1,N)	TWI	446
1105	FORMAT((1X,7(I4,1X,A4,1X,A4,' '),I4,1X,A4,1X,A4))	TWI	447
	PRINT 1106	TWI	448
1106	FORMAT('OSAMPLE NAMES')	TWI	449
	PRINT 1107,(II,INAME1(II),INAME2(II),II=1,MM)	TWI	450
1107	FORMAT((1X,7(I4,1X,2A4,' '),I4,1X,2A4))	TWI	451
	WRITE(4,2100)	TWI	452
	PRINT 2100	TWI	453
2100	FORMAT(//1X,'DO YOU WISH TO OMIT SOME SAMPLES?')	TWI	454
	CALL OMIT(INFLAG,MM)	TWI	455
	I=0	TWI	456
	ID=0	TWI	457
DO 400	II=1,MM	TWI	458
	IF(INFLAG(II).NE.1) GOTO 400	TWI	459
I=I+1		TWI	460
	INAME1(I)=INAME1(II)	TWI	461
	INAME2(I)=INAME2(II)	TWI	462
	INFLAG(I)=II	TWI	463
	IID=IBEGIN(II)	TWI	464
350	ID=ID+1	TWI	465
	IDAT(ID)=IDAT(IID)	TWI	466
	IF(IDAT(IID).EQ.-1) GOTO 400	TWI	467
	IID=IID+1	TWI	468
	GOTO 350	TWI	469
400	CONTINUE	TWI	470
	MM=I	TWI	471
	NID=ID	TWI	472
	RETURN	TWI	473
994	PRINT 1994,N,NMAX	TWI	474
	WRITE(4,1994) N,NMAX	TWI	475
1994	FORMAT(1X,'***ABORT*** LARGEST SPECIES NUMBER ',	TWI	476
	1'IS',I4,'BUT SHOULD NOT EXCEED',I4)	TWI	477
	STOP	TWI	478
995	PRINT 1995,M,MMAX	TWI	479
	WRITE(4,1995) M,MMAX	TWI	480
1995	FORMAT(1X,'***ABORT*** LARGEST SAMPLE NUMBER ',	TWI	481
	1'IS',I4,'BUT SHOULD NOT EXCEED',I4)	TWI	482
	STOP	TWI	483
996	PRINT 1996,II	TWI	484
	WRITE(4,1996) II	TWI	485
1996	FORMAT(1X,'***ABORT*** NO MORE SPACE FOR DATA ',	TWI	486
	1'MATRIX, STOPPED AT SAMPLE',I4)	TWI	487
	STOP	TWI	488
997	PRINT 1997,II	TWI	489
	WRITE(4,1997) II	TWI	490
1997	FORMAT(1X,'***ABORT*** NEGATIVE NUMBER FOUND IN SAMPLE',I4)	TWI	491
	STOP	TWI	492
999	PRINT 1999,III	TWI	493
	WRITE(4,1999) III	TWI	494
1999	FORMAT(1X,'***ABORT*** NON-SEQUENTIAL SAMPLE',	TWI	495

```
1' NUMBER FOUND AFTER SAMPLE',I4)           TWI  496
STOP                                         TWI  497
END                                           TWI  498
C                                             TWI  499
      SUBROUTINE ENTER(NSPEC,NL,LCUT,MMIN,MIND,JMAX,LEVMAX,
1IDIAGR,IPUNCH,LWGT,LIND,INDPOT)           TWI  500
C THIS SUBROUTINE ENTERS PARAMETERS FROM DEVICE 3   TWI  501
      INTEGER LCUT(9),LWGT(9),LIND(9),INDPOT(NSPEC)    TWI  502
      REAL CUT(9)                                     TWI  503
      WRITE(4,2000)                                   TWI  504
      PRINT 2000                                      TWI  505
2000 FORMAT('1','NOW ENTER INPUT PARAMETERS'/1X)    TWI  507
10 PRINT 2005                                      TWI  508
      WRITE(4,2005)                                   TWI  509
2005 FORMAT(1X,'ENTER NUMBER (NOT EXCEEDING 9) OF PSEUDOSPCIES CUT ',
1'LEVELS'/1X,'OR TYPE -1 FOR DEFAULT CUT LEVELS',
2', WHICH ARE 0 2 5 10 20.')                TWI  510
      READ(3,*) NL                                    TWI  511
      PRINT 2010,NL                                 TWI  512
      WRITE(4,2010) NL                               TWI  513
2010 FORMAT(1X,'ANSWER = ',I2)                   TWI  514
      IF(NL.EQ.-1) GOTO 18                         TWI  515
      IF(NL.GE.1.AND.NL.LE.9) GOTO 13             TWI  516
      WRITE(4,2013)
      PRINT 2013
2013 FORMAT(1X,'*** BEWARE *** VALUE LIES OUTSIDE LIMITS',
1' 1,9 - TRY AGAIN')
      GOTO 10
13 WRITE(4,2015) NL
      PRINT 2015,NL
2015 FORMAT('NOW ENTER',I4,' CUT LEVELS')
      READ(3,*) (CUT(IL),IL=1,NL)
      IF(NL.EQ.1) GOTO 20
      DO 14 IL=2,NL
      IF(CUT(IL-1).GT.CUT(IL)) GOTO 16
14 CCNTINUE
      GOTO 20
16 PRINT 2016
      WRITE(4,2016)
2016 FORMAT(1X,'*** BEWARE *** CUT LEVELS OUT OF ORDER - TRY',
1' AGAIN')
      GOTO 10
18 NL=5
      CUT(1)=0.0
      CUT(2)=2.0
      CUT(3)=5.0
      CUT(4)=10.0
      CUT(5)=20.0
20 PRINT 2020,(CUT(IL),IL=1,NL)
      WRITE(4,2020) (CUT(IL),IL=1,NL)
2020 FORMAT('CUT LEVELS'/1X,9F8.2)
      DO 30 IL=1,NL
30 LCUT(IL)=IFIX(1000.0*CUT(IL)+0.5)
      WRITE(4,2030)
      PRINT 2030
                                         TWI  547
                                         TWI  548
                                         TWI  549
                                         TWI  550
```

ONE EDITION

2030 FORMAT('ENTER MINIMUM GROUP SIZE FOR DIVISION')	TWI	551
CALL PARA(MMIN,0,10000,5)	TWI	552
WRITE(4,2040)	TWI	553
PRINT 2040	TWI	554
2040 FORMAT('ENTER MAXIMUM NUMBER OF INDICATORS PER DIVISION')	TWI	555
CALL PARA(MIND,0,15,7)	TWI	556
WRITE(4,2050)	TWI	557
PRINT 2050	TWI	558
2050 FORMAT('ENTER MAXIMUM NUMBER OF SPECIES IN FINAL TABULATION')	TWI	559
CALL PARA(JMAX,0,1000,100)	TWI	560
WRITE(4,2060)	TWI	561
PRINT 2060	TWI	562
2060 FORMAT('ENTER MAXIMUM LEVEL OF DIVISIONS')	TWI	563
CALL PARA(LEVMAX,0,15,6)	TWI	564
WRITE(4,2070)	TWI	565
PRINT 2070	TWI	566
2070 FORMAT('TYPE 1 IF DIAGRAMS OF DIVISIONS ARE WANTED.')	TWI	567
CALL PARA(IDIAGR,0,1,0)	TWI	568
WRITE(4,2075)	TWI	569
PRINT 2075	TWI	570
2075 FORMAT('TYPE 1 IF MACHINE-READABLE COPY OF SOLUTION ', 1'TO BE WRITTEN TO DEVICE 7')	TWI	571
CALL PARA(IPUNCH,0,1,0)	TWI	572
WRITE(4,2080)	TWI	573
PRINT 2080	TWI	574
2080 FORMAT('ENTER WEIGHTS FOR LEVELS OF PSEUDOSPECIES.'/ 11X,'FOR EXAMPLE WEIGHTS 1 2 2 2 SIGNIFY THAT PSEUDOSPECIES'/ 21X,'CORRESPONDING TO 3 HIGHER CUT LEVELS ARE TO BE GIVEN TWICE'/ 31X,'THE WEIGHT OF PSEUDOSPECIES AT THE LOWEST LEVEL.')	TWI	575
CALL VPARA(LWGT,NL,0,1000000)	TWI	576
WRITE(4,2090)	TWI	577
PRINT 2090	TWI	578
2090 FORMAT('ENTER INDICATOR POTENTIALS FOR CUT LEVELS.'/ 11X,'FOR EXAMPLE INDICATOR POTENTIALS 1 0 0 1 0 SIGNIFY '/ 21X,'THAT PSEUDOSPECIES AT LEVELS 1 AND 4 CAN BE USED AS'/ 31X,'INDICATORS, BUT THAT THOSE AT OTHER LEVELS CANNOT.'/ 41X,'IN THE DEFAULT CASE, ALL PSEUDOSPECIES ARE AVAILABLE'/ 51X,'AS INDICATORS.')	TWI	579
CALL VPARA(LIND,NL,0,1)	TWI	580
WRITE(4,2100)	TWI	581
PRINT 2100	TWI	582
2100 FORMAT('DO YOU WISH TO OMIT SOME SPECIES FROM LIST OF'/ 11X,'POTENTIAL INDICATORS? SPECIES OMITTED FROM THIS LIST'/ 21X,'ARE USED IN THE CALCULATION, BUT CANNOT APPEAR AS'/ 3' INDICATORS.')	TWI	583
DO 50 J=1,NSPEC	TWI	584
50 INDPO(TJ)=1	TWI	585
CALL OMIT(INDPO,NSPEC)	TWI	586
RETURN	TWI	587
END	TWI	588
C	TWI	589
SUBROUTINE OMIT(INFLAG,MM)	TWI	590
C STARTING WITH A VECTOR INFLAG, SETS TO ZERO ANY ITEMS THAT ARE	TWI	591
C TYPED IN FROM TERMINAL.	TWI	592
INTEGER INFLAG(MM)	TWI	593
	TWI	594
	TWI	595
	TWI	596
	TWI	597
	TWI	598
	TWI	599
	TWI	600
	TWI	601
	TWI	602
	TWI	603
	TWI	604
	TWI	605

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      WRITE(4,2100)                                     TWI  606
      PRINT 2100                                      TWI  607
2100 FORMAT(1X,'ENTER NUMBERS (NOT NAMES) OF ITEMS TO BE ',
1' OMITTED'/1X,'ONE PER CARD, ENDING LIST WITH A -1./'
21X,'OTHER NEGATIVE NUMBERS DENOTE SEQUENCES. FOR EXAMPLE '
31X,'A 4 FOLLOWED BY A -8 OMITS ITEMS 4 THROUGH 8.')
      NCNONO=MM+1                                     TWI  608
      NCNONO=MM+1                                     TWI  609
200 READ(3,*) NONO                                 TWI  610
      IF(NONO.EQ.-1) GOTO 300                         TWI  611
      IF(NONO.GT.MM) GOTO 260                         TWI  612
      IF(NONO.LT.1) GOTO 250                         TWI  613
      PRINT 2101,NONO                                TWI  614
      WRITE(4,2101) NONO                            TWI  615
2101 FORMAT(1X,'OMIT ITEM',I6)                     TWI  616
      INFLAG(NONO)=0                               TWI  617
      NCNONO=NONO                                TWI  618
      GOTO 200                                     TWI  619
250 IF(NONO.LT.-MM) GOTO 260                      TWI  620
      IF(NONO.GT.-NONONO-1) GOTO 260                TWI  621
      NCNO=-NONO                                TWI  622
      DO 255 II=NONONO,NCNO                        TWI  623
255 INFLAG(II)=0                                 TWI  624
      NONO=-NONO                                TWI  625
      WRITE(4,2104) NONONO,NONO                      TWI  626
      PRINT 2105,NCNO                            TWI  627
2104 FORMAT(1X,'OMIT ITEMS',2I5)                  TWI  628
2105 FORMAT('+',9X,'S',5X,I5)                    TWI  629
      NONONO=MM+1                               TWI  630
      GOTO 200                                     TWI  631
260 PRINT 2103,NONO                            TWI  632
      WRITE(4,2103) NONO                           TWI  633
2103 FORMAT(1X,'*** BEWARE *** INADMISSIBLE NUMBER',I6,' - IGNORED')
      GOTO 200                                     TWI  634
300 PRINT 2102,NONO                            TWI  635
      WRITE(4,2102) NONO                           TWI  636
2102 FORMAT(1X,9X,I6)                           TWI  637
      RETURN                                       TWI  638
      END                                           TWI  639
C
C      SUBROUTINE PARA(K,MIN,MAX,KK)
C      READS PARAMETER K, CHECKS BOUNDS MIN AND MAX, AND SETS THE VALUE OF
C      K TO THE DEFAULT VALUE KK IF K.EQ.-1.
10   WRITE(4,1000) KK,MIN,MAX                      TWI  640
      PRINT 1000,KK,MIN,MAX                        TWI  641
1000 FORMAT(1X,'TYPE -1 FOR DEFAULT VALUE ( =',I3,'); OTHERWISE',
1' TYPE (INTEGER) VALUE REQUIRED/',
21X,'WHICH MUST NOT LIE OUTSIDE LIMITS',2I6)
      READ(3,*) K                                  TWI  642
      PRINT 1001,K                                TWI  643
      WRITE(4,1001) K                            TWI  644
1001 FORMAT(1X,'ANSWER=',I5)                      TWI  645
      IF(K.GE.MIN.AND.K.LE.MAX) RETURN            TWI  646
      IF(K.EQ.-1) GOTO 20                          TWI  647
      PRINT 1002,MIN,MAX                         TWI  648
      WRITE(4,1002) MIN,MAX                       TWI  649

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1002 FORMAT(1X,'*** BEWARE ***  VALUE LIES OUTSIDE ALLOWED LIMITS', TWI 661
 12I6,' - TRY AGAIN') TWI 662
      GOTO 10 TWI 663
 20 K=KK TWI 664
      PRINT 1003,K TWI 665
      WRITE(4,1003) K TWI 666
1003 FORMAT(1X,'VALUE SET TO',I5,' BY DEFAULT') TWI 667
      RETURN TWI 668
      END TWI 669
C TWI 670
      SUBROUTINE VPARA(IX,M,MIN,MAX)
C READS IN PARAMETERS IX AND CHECKS WHETHER IN RANGE MIN,MAX TWI 671
      INTEGER IX(M) TWI 672
 50 PRINT 1000 TWI 673
      WRITE(4,1000) TWI 674
1000 FORMAT(1X,'TYPE -1 FOR DEFAULT VALUES (I.E. IF ALL VALUES TO BE',
 1' SET TO 1)'/1X,'OR TYPE 0 IF NON-DEFAULT VALUES ARE TO BE',
 2' ENTERED')
      READ (3,*) K TWI 678
      PRINT 1001,K TWI 679
      WRITE(4,1001) K TWI 680
1001 FORMAT(1X,'ANSWER=',I5) TWI 681
      IF(K.NE.0) GOTO 100 TWI 682
      PRINT 1002,M,MIN,MAX TWI 683
      WRITE(4,1002) M,MIN,MAX TWI 684
1002 FORMAT(1X,'NOW',I5,' INTEGER VALUES MUST BE SUPPLIED'/
 11X,'NONE OF WHICH MUST LIE OUTSIDE LIMITS',2I9) TWI 685
      READ(3,*) (IX(I),I=1,M) TWI 686
      WRITE(4,1003) TWI 687
      PRINT 1003 TWI 688
1003 FORMAT(1X,'ANSWER = ')
      WRITE(4,1004) (IX(I),I=1,M) TWI 689
      PRINT 1005,(IX(I),I=1,M) TWI 690
1004 FCRMAT(1X,20I4) TWI 691
1005 FORMAT(1X,20I6) TWI 692
 60 DO 60 I=1,M TWI 693
      K=IX(I)
      IF(K.LT.MIN.OR.K.GT.MAX) GOTO 150 TWI 694
 60 CCNTINUE TWI 695
      RETURN TWI 696
100 IF(K.EQ.-1) GOTO 200 TWI 697
150 WRITE(4,1010) K TWI 698
      PRINT 1010,K TWI 699
1010 FORMAT(1X,'*** BEWARE ***  UNACCEPTABLE VALUE',
 1I6,2X,' - TRY AGAIN')
      GOTO 50 TWI 700
200 DO 300 I=1,M TWI 701
300 IX(I)=1 TWI 702
      RETURN TWI 703
      END TWI 704
C TWI 705
      SUBROUTINE ISORT(IX,N)
      INTEGER IX(N)
C EFFICIENT (HEAP-SORT) IN SITU SORTING OF VECTOR IX(N)
      DO 10 I=1,N TWI 706
      END TWI 707
C TWI 708
      SUBROUTINE ISORT(IX,N)
      INTEGER IX(N)
C EFFICIENT (HEAP-SORT) IN SITU SORTING OF VECTOR IX(N)
      DO 10 I=1,N TWI 709
      END TWI 710
C TWI 711
      SUBROUTINE ISORT(IX,N)
      INTEGER IX(N)
C EFFICIENT (HEAP-SORT) IN SITU SORTING OF VECTOR IX(N)
      DO 10 I=1,N TWI 712
      END TWI 713
C TWI 714
      SUBROUTINE ISORT(IX,N)
      INTEGER IX(N)
C EFFICIENT (HEAP-SORT) IN SITU SORTING OF VECTOR IX(N)
      DO 10 I=1,N TWI 715
      END TWI 716
```

```
J=I           TWI 716
JX=IX(J)      TWI 717
5 IF(J.EQ.1) GOTO 8   TWI 718
JJ=J/2        TWI 719
JJX=IX(JJ)    TWI 720
IF(JJX.GE.JX) GOTO 8   TWI 721
IX(J)=JJX    TWI 722
J=JJ         TWI 723
GOTO 5       TWI 724
8 IX(J)=JX    TWI 725
10 CONTINUE   TWI 726
  I=N          TWI 727
  GOTO 14     TWI 728
12 IX(J)=JX    TWI 729
14 IF(I.EQ.1) RETURN  TWI 730
  JX=IX(I)    TWI 731
  IX(I)=IX(1)  TWI 732
  I=I-1       TWI 733
  J=1          TWI 734
  JJ=2         TWI 735
15 IF(I-JJ) 12,17,16  TWI 736
16 JJX=IX(JJ)    TWI 737
  JJJ=JJ+1     TWI 738
  JJJX=IX(JJJ)  TWI 739
  IF(JJX.GE.JJJX) GOTO 18  TWI 740
  IF(JX.GE.JJJX) GOTO 12  TWI 741
  IX(J)=JJJX   TWI 742
  J=JJJ        TWI 743
  JJ=J*2       TWI 744
  GOTO 15     TWI 745
17 JJJX=IX(JJ)  TWI 746
18 IF(JX.GE.JJJX) GOTO 12  TWI 747
  IX(J)=JJX   TWI 748
  J=JJ         TWI 749
  JJ=J*2       TWI 750
  GOTO 15     TWI 751
  END         TWI 752
C           TWI 753
C SUBROUTINE CLORD(MM,LEVMAX,ICLASS,IX)  TWI 754
C ORDERS SAMPLES IN ORDER OF CLASSIFICATION IN ICLASS, MAKING  TWI 755
C ALLOWANCES FOR DIFFERING LEVELS OF DIVISION  TWI 756
  INTEGER ICLASS(MM),IX(MM)  TWI 757
  IEND=2**LEVMAX+2  TWI 758
  IBIG=10000  TWI 759
  DO 10 II=1,MM  TWI 760
  IC=ICLASS(II)  TWI 761
  IF(IC.EQ.0) GOTO 7  TWI 762
5  IC=IC*2  TWI 763
  IF(IC.LT.IEND) GOTO 5  TWI 764
  IX(II)=IBIG*IC+II  TWI 765
  GOTO 10  TWI 766
7  IX(II)=II  TWI 767
10 CONTINUE  TWI 768
  CALL ISORT(IX,MM)  TWI 769
  DO 12 II=1,MM  TWI 770
```

```
IC=IX(II)          TWI 771
12 IX(II)=IC-(IC/IBIG)*IBIG   TWI 772
    RETURN           TWI 773
    END             TWI 774
C               TWI 775
    SUBROUTINE JORDER(M,N,NDAT,IDAT,IADDR,
1JORD,IY,IIY)      TWI 776
C ORDERS DATA MATRIX WITHIN SAMPLES SO AS TO ACCORD WITH A TWI 778
C SPECIFIED ORDER JORD. ADDRESSES OF BEGINNING OF EACH TWI 779
C SAMPLE ARE STORED IN IADDR(I). NOTE THAT QUANTITATIVE TWI 780
C DATA IN EXCESS OF 10000 ARE DISALLOWED. TWI 781
    INTEGER IDAT(NDAT),IADDR(M),JORD(N),IY(N),IIY(N)
    IBIG=10000        TWI 782
    DO 10 J=1,N       TWI 783
    JJ=JORD(J)         TWI 784
10 IY(JJ)=J          TWI 785
    ID=1              TWI 786
    DO 50 I=1,M       TWI 787
    IJ=0              TWI 788
    IADDR(I)=ID       TWI 789
24 J=IDAT(ID)        TWI 790
    ID=ID+1           TWI 791
    IF(J.EQ.-1) GOTO 26  TWI 792
    JJ=IY(J)*IBIG     TWI 793
    JJQ=IDAT(ID)      TWI 794
    ID=ID+1           TWI 795
    JJ=JJ+JJQ          TWI 796
    IJ=IJ+1           TWI 797
    IIY(IJ)=JJ         TWI 798
    GOTO 24            TWI 799
26 IF(IJ.EQ.0) GOTO 50  TWI 800
    CALL ISORT(IIY,IJ)  TWI 801
    ID=IADDR(I)         TWI 802
    DO 40 IIJ=1,IJ       TWI 803
    JJ=IIY(IIJ)          TWI 804
    J=JJ/IBIG            TWI 805
    IDAT(ID)=JORD(J)    TWI 806
    ID=ID+1              TWI 807
    JJQ=JJ-J*IBIG        TWI 808
    IDAT(ID)=JJQ         TWI 809
    ID=ID+1              TWI 810
40 CCNTINUE          TWI 811
    ID=ID+1              TWI 812
50 CONTINUE           TWI 813
    RETURN              TWI 814
    END                TWI 815
C               TWI 816
    SUBROUTINE TABOUT(M,N,IMIN,IMAX,JMIN,JMAX,NDAT,
1IDAT,IADDR,IORD,JORD,INFLAG,          TWI 817
2JNFLAG,JNAME1,JNAME2,ICLASS,JCLASS)  TWI 818
C PRINTS OUT ROWS AND COLUMNS OF MATRIX IN ARBITRARY TWI 819
C ORDERS IORD, JORD (BUT NOTE THAT ROWS ARE ASSUMED TO BE TWI 820
C ALREADY ORDERED INTERNALLY BY JORD). THE MATRIX IS TWI 821
C TRANSPOSED FOR PRINTING OUT, AS IS CONVENTIONAL IN TWI 822
C PHYTOSOCIOLOGY. THE LIMITS IMIN ETC ARE THE LIMITS TWI 823
TWI 824
TWI 825
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C OUTSIDE WHICH MATRIX IS NOT PRINTED.          TWI 826
  INTEGER IDAT(NDAT),IADDR(M),IORD(M),JORD(N),
  ITEM(200),IOUT(10)                         TWI 827
  INTEGER BLANK,STAR,DASH                     TWI 828
  INTEGER INFLAG(M),JNFLAG(N),JNAME1(N),JNAME2(N) TWI 829
  INTEGER ICLASS(M),JCLASS(N)                  TWI 830
  COMMON/WORK/ITEM                           TWI 831
  DATA BLANK/' '/,STAR/**/                  TWI 832
  DATA DASH/-/-/                            TWI 833
  DATA IOUT/'0','1','2','3','4','5','6','7','8','9'/
  DO 10 I=1,200                                TWI 834
10  ITEM(I)=BLANK                           TWI 835
    NNAME=BLANK                           TWI 836
    PRINT 1002                           TWI 837
1002 FORMAT('1')                           TWI 838
    DO 30 IL=1,4                           TWI 839
      IDIV=10***(4-IL)
      III=2
      DO 20 I=IMIN,IMAX
        III=III+1
        II=IORD(I)
        IORIG=INFLAG(II)
        IR=IORIG/IDIV
        IF(IR.GE.1) ITEM(III)=IOUT(IR-(IR/10)*10+1)
20  CCNTINUE                           TWI 840
30  PRINT 1000,(ITEM(II),II=1,III)          TWI 841
    PRINT 1001                           TWI 842
1001 FORMAT(1X)
    IF(JMAX.LT.JMIN) RETURN               TWI 843
    DO 110 J=JMIN,JMAX                   TWI 844
      JJ=JCRD(J)
      CALL BIN(ITEM,JCLASS(JJ))
      DO 90 II=1,6                         TWI 845
90  ITEM(150+II)=ITEM(1+II)                TWI 846
    ITEM(1)=JNFLAG(JJ)
    ITEM(2)=JNAME1(JJ)
    ITEM(3)=JNAME2(JJ)
    III=3
    DO 100 I=IMIN,IMAX
      II=IORD(I)
      ID=IADDR(II)
53  JJJ=IDAT(ID)
    IF(JJJ.NE.JJ) GOTO 98
    ID=ID+1
    JQ=IDAT(ID)
    ID=ID+1
    IADDR(II)=ID
    III=III+1
    IF(JQ.GE.10) GOTO 96
    ITEM(III)=IOUT(JQ+1)
    GOTO 100
96  ITEM(III)=STAR
    GOTO 100
98  III=III+1
    ITEM(III)=DASH                         TWI 847
                                         TWI 848
                                         TWI 849
                                         TWI 850
                                         TWI 851
                                         TWI 852
                                         TWI 853
                                         TWI 854
                                         TWI 855
                                         TWI 856
                                         TWI 857
                                         TWI 858
                                         TWI 859
                                         TWI 860
                                         TWI 861
                                         TWI 862
                                         TWI 863
                                         TWI 864
                                         TWI 865
                                         TWI 866
                                         TWI 867
                                         TWI 868
                                         TWI 869
                                         TWI 870
                                         TWI 871
                                         TWI 872
                                         TWI 873
                                         TWI 874
                                         TWI 875
                                         TWI 876
                                         TWI 877
                                         TWI 878
                                         TWI 879
                                         TWI 880
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100 CONTINUE          TWI  881
    III=III+1          TWI  882
    ITEM(III)=BLANK    TWI  883
    III=III+1          TWI  884
    ITEM(III)=BLANK    TWI  885
    DO 105 II=1,6      TWI  886
    III=III+1          TWI  887
105 ITEM(III)=ITEM(150+II)  TWI  888
    PRINT 2000,(ITEM(II),II=1,III)  TWI  889
1000 FORMAT(1X,5X,A4,1X,A4,2X,108A1)  TWI  890
2000 FORMAT(1X,I4,1X,A4,1X,A4,2X,108A1)  TWI  891
110 CONTINUE          TWI  892
    PRINT 1001          TWI  893
    ITEM(49)=BLANK     TWI  894
    ITEM(50)=BLANK     TWI  895
    DO 220 LEVEL=1,6   TWI  896
    III=50             TWI  897
    DO 210 I=IMIN,IMAX TWI  898
    II=IORD(I)         TWI  899
    IC=ICLASS(II)      TWI  900
    CALL BIN(ITEM,IC)  TWI  901
    III=III+1          TWI  902
210 ITEM(III)=ITEM(1+LEVEL)  TWI  903
    PRINT 1000,(ITEM(II),II=49,III)  TWI  904
220 CCNTINUE          TWI  905
    RETURN             TWI  906
    END                TWI  907
C                                     TWI  908
    SUBROUTINE PSEUDO(MM,NN,NMAX,NL,NDAT,NSPEC,IDAT,LCUT,
    1JNFLAG,JNAME1,JNAME2,JNAM,INDPOT,IIY)
C SETS UP PSEUDOSPECIES DATA IN IDAT, AND ALSO VECTOR INDPOT,
C WHICH STORES THE REAL SPECIES NUMBERS CORRESPONDING TO THE
C PSEUDOSPECIES. PSEUDOSPECIES CUT LEVELS ARE IN LCUT.
C SPECIES NAMES IN JNAME1,2. JNAM STORES LEVELS OF PSEUDOSPECIES.
    INTEGER JT0P(10),IDAT(NDAT),LCUT(NL),JNAME1(NMAX),JNAME2(NMAX),
    1JNFLAG(NMAX),JNAM(NMAX),INDPOT(NMAX),IIY(NMAX)
    IBIG=50000
    ID=0
    IDMAX=0
    DO 20 II=1,MM
15  ID=ID+1
    IDMAX=IDMAX+1
    J=IDAT(ID)
    IF(J.EQ.-1) GOTO 20
    ID=ID+1
    JQ=IDAT(ID)
    JJQ=0
    DO 16 IL=1,NL
    IF(JQ.GE.LCUT(IL))JJQ=JJQ+1
16  CCNTINUE
    IDAT(ID)=JJQ
    IDMAX=IDMAX+JJQ-1
    GOTO 15
20  CCNTINUE
    IF(IDMAX.LE.NDAT) GOTO 40
    TWI  909
    TWI  910
    TWI  911
    TWI  912
    TWI  913
    TWI  914
    TWI  915
    TWI  916
    TWI  917
    TWI  918
    TWI  919
    TWI  920
    TWI  921
    TWI  922
    TWI  923
    TWI  924
    TWI  925
    TWI  926
    TWI  927
    TWI  928
    TWI  929
    TWI  930
    TWI  931
    TWI  932
    TWI  933
    TWI  934
    TWI  935
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BEE ERRATA

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IDMAX=IDMAX+50                                TWI 936
PRINT 1001, IDMAX                            TWI 937
WRITE(4,1001) IDMAX                          TWI 938
1001 FORMAT(1X,'***ERROR*** DATA ARRAY NEEDS',I6,' ELEMENTS') TWI 939
      STOP                                     TWI 940
      40 IID=NDAT                             TWI 941
      80 IDAT(IID)=IDAT(ID)                   TWI 942
          ID=ID-1                           TWI 943
          IID=IID-1                         TWI 944
          IF(ID.GT.0) GOTO 80                 TWI 945
          DO 90 II=1,MM                      TWI 946
          JJJ=0                           TWI 947
      84 IID=IID+1                           TWI 948
          J=IDAT(IID)                         TWI 949
          IF(J.EQ.-1) GOTO 87                 TWI 950
          IID=IID+1                         TWI 951
          JJQ=IDAT(IID)                       TWI 952
          IF(JJQ.EQ.0) GOTO 84                 TWI 953
          DO 86 JQ=1,JJQ                      TWI 954
          JJJ=JJJ+1                         TWI 955
      86 IIY(JJJ)=J+IBIG*(JQ-1)             TWI 956
          GOTO 84                           TWI 957
      87 DO 88 JJ=1,JJJ                      TWI 958
          ID=ID+1                           TWI 959
      88 IDAT(ID)=IIY(JJ)                   TWI 960
          ID=ID+1                           TWI 961
          IDAT(ID)=-1                      TWI 962
          IF(ID.LE.IID) GOTO 90              TWI 963
          IDMAX=IDMAX+300                   TWI 964
          PRINT 1001, IDMAX                  TWI 965
          STOP                               TWI 966
      90 CCNTINUE                           TWI 967
C WE NOW HAVE THE DESIRED INFORMATION IN IDAT. ALL WE HAVE
C TO DO IS TO CONDENSE THE NUMBERING           TWI 968
      IP=0                                 TWI 969
      DO 140 IL=1,NL                      TWI 970
      DO 100 JJJ=1,NSPEC                   TWI 971
      100 IIY(JJJ)=0                      TWI 972
          ID=0                               TWI 973
          DO 110 II=1,MM                   TWI 974
      103 ID=ID+1                           TWI 975
          JJJ=IDAT(ID)                     TWI 976
          IF(JJJ.EQ.-1) GOTO 110            TWI 977
          IF(JJJ.GE.IBIG) GOTO 103          TWI 978
          IF(JJJ.LT.-1) GOTO 103            TWI 979
          IIY(JJJ)=IIY(JJJ)+1              TWI 980
          GOTO 103                           TWI 981
      110 CCNTINUE                           TWI 982
          DO 120 JJJ=1,NSPEC                TWI 983
          IF(IIY(JJJ).EQ.0) GOTO 120          TWI 984
          IP=IP+1                           TWI 985
          INDPT(IP)=JJJ                     TWI 986
          IIY(JJJ)=IP                      TWI 987
      120 CCNTINUE                           TWI 988
          ID=0                               TWI 989
                                         TWI 990
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DO 130 II=1,MM          TWI  991
123 ID=ID+1             TWI  992
  JJJ=IDAT(ID)          TWI  993
  IF(JJJ.EQ.-1) GOTO 130 TWI  994
  IF(JJJ.GE.IBIG) GOTO 127 TWI  995
  IF(JJJ.LT.-1) GOTO 123 TWI  996
  JJJ=IIY(JJJ)           TWI  997
127 ICAT(ID)=JJJ-IBIG   TWI  998
  GOTO 123              TWI  999
130 CONTINUE             TWI 1000
  JT0P(IL)=IP           TWI 1001
140 CONTINUE             TWI 1002
  ID=0                  TWI 1003
  DO 150 II=1,MM         TWI 1004
145 ID=ID+1             TWI 1005
  JJ=IDAT(ID)           TWI 1006
  IF(JJ.EQ.-1) GOTO 150 TWI 1007
  IDAT(ID)=JJ+IBIG     TWI 1008
  GOTO 145              TWI 1009
150 CCNTINUE             TWI 1010
  PRINT 1002,ID          TWI 1011
  WRITE(4,1002) ID        TWI 1012
1002 FORMAT(//1X,'LENGTH OF DATA ARRAY AFTER DEFINING PSEUDOSPECIES',TWI 1013
  I17)
  IF(ID*2.GT.NDAT) PRINT 1003,NDAT      TWI 1014
  IF(ID*2.GT.NDAT) WRITE(4,1003) NDAT    TWI 1015
1003 FORMAT(1X,'*** BEWARE *** THIS EXCEEDS ONE-HALF LENGTH OF',TWI 1016
  1' DATA ARRAY (=',I6,')'/      TWI 1017
  21X,'NO SPECIES CLASSIFICATION CAN BE PERFORMED')
  NN=IP                  TWI 1018
  IF(NN.LE.NMAX) GOTO 155      TWI 1019
  PRINT 1000,NN              TWI 1020
  WRITE(4,1000) NN            TWI 1021
1000 FORMAT(1X,'***ERROR*** - NEEDS DIMENSIONING FOR',I6,      TWI 1022
  1' PSEUDOSPECIES')
  STOP                   TWI 1023
155 NSPEC=JT0P(1)          TWI 1024
  PRINT 1004,NN            TWI 1025
  WRITE(4,1004) NN          TWI 1026
1004 FORMAT(//1X,'TOTAL NUMBER OF SPECIES AND PSEUDOSPECIES',I7) TWI 1027
  PRINT 1005,NSPEC          TWI 1028
  WRITE(4,1005) NSPEC        TWI 1029
1005 FORMAT(//1X,'NUMBER OF SPECIES, EXCLUDING PSEUDOSPECIES ',TWI 1030
  1'AND ONES WITH NO OCCURRENCES',I5)
  DO 160 JJ=1,NSPEC        TWI 1031
  J=INDPOT(JJ)             TWI 1032
  IIY(J)=JJ                TWI 1033
  JNFLAG(JJ)=JNFLAG(J)     TWI 1034
  JNAME1(JJ)=JNAME1(J)     TWI 1035
160 JNAME2(JJ)=JNAME2(J)   TWI 1036
  DO 170 JJ=1,NN            TWI 1037
  J=INDPOT(JJ)             TWI 1038
170 INDPOT(JJ)=IIY(J)      TWI 1039
  JJJ=2                    TWI 1040
  IL=0                     TWI 1041

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SEE EPARATE

SEE LARVA

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DO 180 JJ=1,NN          TWI 1046
J=IABS(INDPOT(JJ))      TWI 1047
JNFLAG(JJ)=JNFLAG(J)    TWI 1048
JNAME1(JJ)=JNAME1(J)    TWI 1049
JNAME2(JJ)=JNAME2(J)    TWI 1050
IF(J.LE.JJJ) IL=IL+1    TWI 1051
JNAM(JJ)=IL              TWI 1052
180 JJJ=J                TWI 1053
RETURN                  TWI 1054
END                      TWI 1055
C                         TWI 1056
      SUBROUTINE CLASS(MM,NN,NDAT,MIND,MMZ,MMS,IX,ICLASS,
1 IIROW,IADDR,INDPOT,INDORD,IZONE,IY,JJCOL,IDAT,
2 INDSIG,IPICT,X,XX,RTOT,RRWT,ROWWGT,Y,YY,CTOT,CCWT,
3 COLWGT,I NAME1,I NAME2,JNAME1,JNAME2,JNAM,X3,X4,X5,LIND,
4 INFLAG,IPUNCH)          TWI 1057
C THIS DOES THE CLASSIFICATION. IF ISEC.EQ.2 IT DOES A SPECIES   TWI 1058
C CLASSIFICATION, PRINTING OUT LESS INFORMATION THAN IN THE CASE WHERE   TWI 1059
C ISEC.EQ.1.               TWI 1060
      INTEGER PLUS,IX(MM),ICLASS(MM),IIROW(MM),IADDR(MM),INFLAG(MM),   TWI 1061
1 INDPOT(NN),INDORD(NN),IZONE(MM),IY(NN),                           TWI 1062
2 JJCOL(NN),IDAT(NDAT),INDSIG(MMS),                                TWI 1063
3 IPICT(MMZ,MMS),ITEM(200)                                         TWI 1064
      INTEGER LIND(10)
      INTEGER I NAME1(MM),I NAME2(MM),JNAME1(NN),
1 JNAME2(NN),JNAM(NN)
      REAL X(MM),XX(MM),RTOT(MM),RRWT(MM),ROWWGT(MM),Y(NN),YY(NN),
1 CTOT(NN),CCWT(NN),COLWGT(NN)
      REAL X3(MM),X4(MM),X5(MM)
      DATA PLUS/'+'/,MINUS/'-'/
      CCOMMON/WORK/ITEM
      COMMON/PICT/MZ,MS,IZD,IIZD,ISD,ISHIFT,MZIND,
1 MZCRIT,MZOUT,NIND
      COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM
      CCOMMON/SECCND/ISEC
      CCOMMON/SWITCH/IFAIL,IDIAGR,ISTOP,IREWT
      CCOMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX
      COMMON/ARBS/CWTMIN,CRLONG,CRCUT
      IC=1
      LEVEL=1
      ISTOP=0
      IEND=2**LEVMAX+2
      ID=1
      DO 5 II=1,MM
      IADDR(II)=ID
4  JJ=IDAT(ID)
      ID=ID+1
      IF(JJ.EQ.-1) GOTO 5
      GOTO 4
5  CCNTINUE
      DO 10 II=1,MM
10  ICLASS(II)=IC
      DO 15 II=1,MM
      ID=IADDR(II)
      IF(IDAT(ID).EQ.-1) ICLASS(II)=IEND
      TWI 1065
      TWI 1066
      TWI 1067
      TWI 1068
      TWI 1069
      TWI 1070
      TWI 1071
      TWI 1072
      TWI 1073
      TWI 1074
      TWI 1075
      TWI 1076
      TWI 1077
      TWI 1078
      TWI 1079
      TWI 1080
      TWI 1081
      TWI 1082
      TWI 1083
      TWI 1084
      TWI 1085
      TWI 1086
      TWI 1087
      TWI 1088
      TWI 1089
      TWI 1090
      TWI 1091
      TWI 1092
      TWI 1093
      TWI 1094
      TWI 1095
      TWI 1096
      TWI 1097
      TWI 1098
      TWI 1099
      TWI 1100
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15 CCNTINUE          TWI 1101
      GOTO 30          TWI 1102
20 CALL DECODE(M,MM,N,NN,NDAT,IIROW,IADDR,IDAT,JJCOL)  TWI 1103
      CALL UPDATE(M,MM,IC,LEVEL,ICLASS,IIROW,IX)          TWI 1104
      IF(ISTOP.EQ.1) GOTO 500                           TWI 1105
30 CALL RECODE(M,MM,N,NN,NDAT,IC,ICLASS,IIROW,JJCOL,IADDR,IDAT)  TWI 1106
      CALL BIN(IITEM,IC)          TWI 1107
      PRINT 1010          TWI 1108
1010 FORMAT(1X,30('*****'))          TWI 1109
      PRINT 1000,IC,M,(IITEM(IT),IT=1,30)          TWI 1110
      WRITE(4,1000) IC,M,(IITEM(IT),IT=1,30)          TWI 1111
1000 FORMAT('ODIVISION',I5,' (N='',I4,'',          TWI 1112
      19X,'I.E. GROUP ',30A1)          TWI 1113
      IF(M.GE.MMIN) GOTO 40          TWI 1114
      PRINT 1001          TWI 1115
1001 FORMAT(' DIVISION FAILS - THERE ARE TOO FEW ITEMS')  TWI 1116
      IFAIL=1          TWI 1117
      GOTO 20          TWI 1118
40 CALL WEIGHT(M,MM,N,NN,NDAT,X,Y,IIROW,IADDR,          TWI 1119
      IRRWT,ROWWGT,JJCOL,CCWT,COLWGT,IDAT)          TWI 1120
      CALL RA(M,MM,N,NN,NDAT,EIG,X,XX,X3,X4,X5,Y,RTOT,          TWI 1121
      ICTOT,IIROW,IADDR,RCWGT,COLWGT,IDAT)          TWI 1122
      IF(IFAIL.EQ.1) GOTO 20          TWI 1123
      CALL XMAXMI(X,AXMAX,AXMIN,M)          TWI 1124
      IF(AXMAX.GT.-AXMIN) GOTO 45          TWI 1125
      DO 44 I=1,M          TWI 1126
44 X(I)=-X(I)          TWI 1127
      AX=AXMAX          TWI 1128
      AXMAX=-AXMIN          TWI 1129
      AXMIN=-AX          TWI 1130
45 CRMIN=AXMIN*CRLONG          TWI 1131
      CRMAX=AXMAX*CRLONG          TWI 1132
      CRMID=0.5*(CRMIN+CRMAX)          TWI 1133
      CRHALF=0.5*(CRMAX-CRMIN)          TWI 1134
      IF(IREWWT.EQ.0) GOTO 48          TWI 1135
      DO 47 IRE=1,IREWT          TWI 1136
      DO 46 I=1,M          TWI 1137
      AX=(X(I)-CRMID)/CRHALF          TWI 1138
      IF(AX.GT.1.0) AX=1.0          TWI 1139
      IF(AX.LT.-1.0) AX=-1.0          TWI 1140
46 XX(I)=AX          TWI 1141
      CALL POLISH(M,MM,N,NN,NDAT,X,XX,Y,YY,IIROW,JJCOL,RTOT,          TWI 1142
      1ROWWGT,CCWT,CULWGT,IADDR,IDAT)          TWI 1143
      CALL XMAXMI(X,AXMAX,AXMIN,M)          TWI 1144
      CRMID=0.5*(AXMIN+AXMAX)          TWI 1145
      CRHALF=0.5*CRLONG*(AXMAX-AXMIN)          TWI 1146
      CRMIN=CRMID-CRHALF          TWI 1147
47 CRMAX=CRMID+CRHALF          TWI 1148
48 IF(IC.EQ.1) GOTO 63          TWI 1149
C WE NOW FIND OUT WHICH OF THE DERIVED GROUPS MORE CLOSELY RESEMBLES  TWI 1150
C THE BROTHER OF THE GROUP THAT HAS JUST BEEN DIVIDED          TWI 1151
C TO DO THIS WE REQUIRE SPECIES TOTALS (IGNORING PSEUDOSPECIES WEIGHTS)  TWI 1152
      SMALL=1.0E-7          TWI 1153
      DO 49 JJ=1,NN          TWI 1154
      Y(JJ)=SMALL          TWI 1155
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49 YY(JJ)=SMALL          TWI 1156
  TOT0=0.0                TWI 1157
  TOT1=0.0                TWI 1158
  DO 52 I=1,M              TWI 1159
  SI=IIRROW(I)            TWI 1160
  ID=IADDR(II)             TWI 1161
  IF(X(I).GE.CRMID) TOT1=TOT1+RRWT(II)  TWI 1162
  IF(X(I).LE.CRMID) TOT0=TOT0+RRWT(II)  TWI 1163
51 J=IDAT(ID)             TWI 1164
  ID=ID+1                 TWI 1165
  IF(J.EQ.-1) GOTO 52      TWI 1166
  JJ=JJCOL(J)              TWI 1167
  JJJ=IABS(INDPCT(JJ))    TWI 1168
  IF(X(I).GE.CRMID) YY(JJJ)=YY(JJJ)+RRWT(II)*CCWT(JJ)  TWI 1169
  IF(X(I).LE.CRMID) Y(JJJ)=Y(JJJ)+RRWT(II)*CCWT(JJ)  TWI 1170
  GOTO 51                  TWI 1171
52 CONTINUE                TWI 1172
  IIC=IC+1                 TWI 1173
  IF(IIC.EQ.(IIC/2)*2) IIC=IIC-2           TWI 1174
  CALL CLOSER(YSCOR1,IIC,MM,N,NN,NDAT,IEND,TOTO,TOT1,
  1IDAT,ICLASS,IADDR,INDPOT,JJCOL,RRWT,CCWT,COLWGT,Y,YY)  TWI 1175
  IIIIC=IC/2+1              TWI 1176
  IF(IIIIC.EQ.(IIIIC/2)*2) IIIIC=IIIIC-2           TWI 1177
  YSCOR2=0.0                 TWI 1178
  IF(IC.LE.3) GOTO 54        TWI 1179
  CALL CLOSER(YSCOR2,IIIIC,MM,N,NN,NDAT,IEND,TOTO,TOT1,
  1IDAT,ICLASS,IADDR,INDPOT,JJCOL,RRWT,CCWT,COLWGT,Y,YY)  TWI 1180
  54 IW=IC-(IC/4)*4          TWI 1181
  W=0.5                      TWI 1182
  IF(IW.EQ.1.OR.IW.EQ.2) W=-W           TWI 1183
  YSCORE=YSCOR1+W*YSCOR2         TWI 1184
  IF(IIC.NE.(IIC/2)*2) GOTO 559        TWI 1185
C IIC IS EVEN, SO WE WANT THE NEGATIVE GROUP TO BE CLOSER TO IT  TWI 1186
  IF(YSCORE.LE.0) GOTO 63            TWI 1187
  GOTO 60                      TWI 1188
C IN THIS CASE IIC IS ODD          TWI 1189
  559 IF(YSCORE.GE.0) GOTO 63        TWI 1190
  60 DO 61 I=1,M                 TWI 1191
  61 X(I)=-X(I)                 TWI 1192
  AX=AXMAX                     TWI 1193
  AXMAX=-AXMIN                 TWI 1194
  AXMIN=-AX                     TWI 1195
  AX=CRMAX                     TWI 1196
  CRMAX=-CRMIN                 TWI 1197
  CRMIN=-AX                     TWI 1198
  CRMID=-CRMID                 TWI 1199
  63 CALL ZONEUP(X,AXMIN,CRMIN,CRMAX,AXMAX,M,MZ,MZOUT,MZCRIT,IZONE)  TWI 1200
  IF(MIND.EQ.0) GOTG 100          TWI 1201
  CUT1=CRMID-0.5*(CRMAX-CRMIN)*FLOAT(MZIND)/(FLOAT(MZCRIT)+0.001)  TWI 1202
  CUT2=2.0*CRMID-CUT1            TWI 1203
  CALL INDSCC(M,MM,N,NN,NDAT,X,AXPOS,AXNEG,
  1CUT1,CUT2,Y,YY,IIRCW,ROWWGT,IADDR,IDAT)           TWI 1204
  DO 65 J=1,N                   TWI 1205
  65 Y(J)=YY(J)/AXPOS-Y(J)/AXNEG        TWI 1206
  CALL TOPIND(N,NN,MIND,FEEBLE,Y,INDORD,INDSIG,        TWI 1207
                                TWI 1208
                                TWI 1209
                                TWI 1210
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SEE ERRATA

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1JJCOL,INDPCT,JNAM,NIND,LIND)          TWI 1211
IF(NIND.EQ.0) GOTO 100                  TWI 1212
MIS=10000                               TWI 1213
CALL CODESC(M,MM,N,NN,NDAT,NIND,INDORD,IX,IY,IIROW,IADDR,IDAT)  TWI 1214
C NOW FIND BEST NUMBER OF INDICATORS AND BEST DIVISION POINT    TWI 1215
DO 70 IIND=1,NIND                      TWI 1216
CALL TABLE(M,MMZ,MMS,MIND,IX,IZONE,INDSIG,IPICT,O,IIND)        TWI 1217
CALL FIND(MMZ,MMS,IPICT,IZD1,ISD1,MISCL)                      TWI 1218
IF(MISCL.GE.MIS)GOTO 70                  TWI 1219
MIS=MISCL                               TWI 1220
ISD=ISD1                                TWI 1221
IZD=IZD1                                TWI 1222
IIZD=IZD-MZIND                          TWI 1223
IND=IIND                                 TWI 1224
70 CCNTINUE                               TWI 1225
CALL TABLE(M,MMZ,MMS,MIND,IX,IZONE,INDSIG,IPICT,1,IND)        TWI 1226
PRINT 1003                               TWI 1227
1003 FURMAT(1X,'INDICATORS, TOGETHER WITH THEIR SIGN')
IT=0                                     TWI 1228
DO 80 IIND=1,IND                      TWI 1229
J=INDORD(IIND)                         TWI 1230
JJ=JJCOL(J)                            TWI 1231
IT=IT+1                                 TWI 1232
ITEM(IT)=JNAME1(JJ)                     TWI 1233
IT=IT+1                                 TWI 1234
ITEM(IT)=JNAME2(JJ)                     TWI 1235
IT=IT+1                                 TWI 1236
ITEM(IT)=JNAM(JJ)                       TWI 1237
IT=IT+1                                 TWI 1238
ITEM(IT)=JNAM(JJ)                       TWI 1239
IT=IT+1                                 TWI 1240
IF(INDSIG(IIND).GT.0) GOTO 75          TWI 1241
ITEM(IT)=MINUS                           TWI 1242
GOTO 80                                 TWI 1243
75 ITEM(IT)=PLUS                         TWI 1244
80 CCNTINUE                               TWI 1245
PRINT 1004,(ITEM(IIT),IIT=1,IT)
1004 FORMAT(1X,7(A4,1X,A4,I1,'(',A1,')',3X))  TWI 1246
GOTO 120                                TWI 1247
100 DO 110 I=1,M                         TWI 1248
110 IX(I)=0                               TWI 1249
MS=1                                     TWI 1250
IZD=(1+MZ)/2                            TWI 1251
IIZD=IZD                                TWI 1252
ISD=0                                     TWI 1253
IF(MIND.EQ.0) NIND=0                     TWI 1254
IF(MIND.NE.0)PRINT 1006                  TWI 1255
1006 FORMAT(1X,'(NO INDICATORS)')
120 CALL REPORT(M,MM,N,NN,NDAT,MMZ,MMS,IC,X,XX,ROWWGT,
1Y,YY,COLWGT,IX,IZONE,IIROW,IY,JJCOL,IADDR,IDAT,IPICT,
2INAME1,INAME2,JNAME1,JNAME2,JNAM)      TWI 1257
DO 130 I=1,M                            TWI 1258
AX=X(I)                                 TWI 1259
IF(AX.GT.3.5) IX(I)=1                   TWI 1260
IF(AX.LT.3.5) IX(I)=0                   TWI 1261
130 CCNTINUE                               TWI 1262
GOTO 20                                 TWI 1263
130 CCNTINUE                               TWI 1264
GOTO 20                                 TWI 1265
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500 PRINT 1010          TWI 1266
  IF(IPUNCH.EQ.0) RETURN      TWI 1267
  NUMBER=0                  TWI 1268
  DO 510 II=1,MM            TWI 1269
    IF(ICLASS(II).NE.0) NUMBER=NUMBER+1
510 CONTINUE               TWI 1270
  WRITE(7,1020) NUMBER      TWI 1271
1020 FORMAT(1X,'THERE ARE',I6,' ITEMS')
  DO 550 II=1,MM            TWI 1272
    IC=ICLASS(II)
    IF(IC.EQ.0) GOTO 550      TWI 1273
    CALL BIN(ITEM,IC)
    IF(ISEC.EQ.2) GOTO 530      TWI 1274
    WRITE(7,1030) INFLAG(II),INAME1(II),INAME2(II),IC,
    1(ITEM(IT),IT=1,30)        TWI 1275
    GOTO 550                  TWI 1276
530 WRITE(7,1040) INFLAG(II),INAME1(II),INAME2(II),IC,
    1(ITEM(IT),IT=1,30)        TWI 1277
1030 FORMAT(1X,I4,1X,2A4,I7,3X,30A1)      TWI 1278
1040 FORMAT(1X,I4,1X,A4,1X,A4,I6,3X,30A1)      TWI 1279
550 CCNTINUE               TWI 1280
  RETURN                     TWI 1281
  END                        TWI 1282
C
C           SUBROUTINE CLOSER(YSCORE,IIC,MM,N,NN,NDAT,IEND,TOTO,TOT1,
1IDAT,ICLASS,IADDR,INDPOT,JJCOL,RRWT,CCWT,Y,Y0,Y1)      TWI 1283
C OBTAINS A QUANTITY YSCORE, TO DETERMINE WHICH ORDER THE TWO HALVES      TWI 1284
C OF DIVISION SHOULD BE PUT IN. IF YSCORE IS NEGATIVE, THEN GROUP      TWI 1285
C IIC MORE CLOSELY RESEMBLES THE NEGATIVE SIDE OF DIVISION, AND      TWI 1286
C VICE VERSA IF YSCORE IS POSITIVE.      TWI 1287
  INTEGER IDAT(NDAT),IADDR(MM),ICLASS(MM),INDPOT(NN),JJCOL(NN)      TWI 1288
  REAL RRWT(MM),CCWT(NN),Y(NN),Y0(NN),Y1(NN)      TWI 1289
  TGT=0.0                      TWI 1290
  DO 50 JJ=1,NN                TWI 1291
  50 Y(JJ)=1.0E-7              TWI 1292
  DO 54 II=1,MM                TWI 1293
    IIIC=ICLASS(II)
    IF(IIIC.GT.IEND) IIIC=IIIC-IEND      TWI 1294
    IF(IIIC.NE.IIC.AND.IIIC/2.NE.IIC.AND.IIIC/4.NE.IIC) GOTO 54      TWI 1295
    ID=IADDR(II)
    TOT=TOT+RRWT(II)
  53 JJ=IDAT(ID)
    ID=ID+1
    IF(JJ.EQ.-1) GOTO 54      TWI 1300
    JJJ=IABS(INDPOT(JJ))
    Y(JJJ)=Y(JJJ)+RRWT(II)*CCWT(JJ)      TWI 1301
    GOTO 53                  TWI 1302
  54 CCNTINUE               TWI 1303
    YNEG=0.0                  TWI 1304
    YIND=0.0                  TWI 1305
    YPOS=0.0                  TWI 1306
    X1NEG=0.0                 TWI 1307
    X1IND=0.0                 TWI 1308
    X1POS=0.0                 TWI 1309
    XONEG=0.0                 TWI 1310
    TWI 1311
    TWI 1312
  55 CCNTINUE               TWI 1313
    YNEG=0.0                  TWI 1314
    YIND=0.0                  TWI 1315
    YPOS=0.0                  TWI 1316
    X1NEG=0.0                 TWI 1317
    X1IND=0.0                 TWI 1318
    X1POS=0.0                 TWI 1319
    XONEG=0.0                 TWI 1320
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XOIND=0.0          TWI 1321
XOPOS=0.0          TWI 1322
DO 57 J=1,N        TWI 1323
JJ=JJCOL(J)        TWI 1324
JJJ=IABS(INDPOT(JJ)) TWI 1325
IF(JJ.NE.JJJ) GOTO 57 TWI 1326
AY=Y(JJ)/TOT      TWI 1327
AY0=YO(JJJ)/TOT0  TWI 1328
AY1=Y1(JJJ)/TOT1  TWI 1329
PREF=ABS(AY0-AY1)/(AY0+AY1) TWI 1330
PREF=PREF/0.3     TWI 1331
IF(PREF.GT.1.0) PREF=1.0 TWI 1332
PREF=PREF*PREF*PREF*PREF TWI 1333
IF(AY1.GT.AY0) GOTO 55 TWI 1334
PNEG=PREF         TWI 1335
PIND=(1.0-PREF)   TWI 1336
PPOS=0.0          TWI 1337
GOTO 56          TWI 1338
55 PNEG=0.0        TWI 1339
PIND=(1.0-PREF)   TWI 1340
PPOS=PREF         TWI 1341
56 YNEG=YNEG+PNEG*AY TWI 1342
YIND=YIND+PIND*AY TWI 1343
YPOS=YPOS+PPOS*AY TWI 1344
XONEG=XONEG+PNEG*AY0 TWI 1345
X1NEG=X1NEG+PNEG*AY1 TWI 1346
XOIND=XOIND+PIND*AY0 TWI 1347
X1IND=X1 IND+PIND*AY1 TWI 1348
XOPOS=XOPOS+PPOS*AY0 TWI 1349
X1POS=X1POS+PPOS*AY1 TWI 1350
57 CONTINUE        TWI 1351
XNEG=XONEG+X1NEG  TWI 1352
XIND=XOIND+X1IND  TWI 1353
XPOS=XOPOS+X1POS  TWI 1354
IF(XPOS.GT.XNEG)GOTO 58 TWI 1355
IF(XIND.LE.(XNEG-XPOS)) GOTO 59 TWI 1356
YIND=YIND*(XNEG-XPOS)/XIND TWI 1357
GOTO 59          TWI 1358
58 YIND=-YIND    TWI 1359
XIND=-XIND      TWI 1360
IF(-XIND.LE.(XPOS-XNEG)) GOTO 59 TWI 1361
YIND=YIND*(XPOS-XNEG)/(-XIND) TWI 1362
59 YSCORE=YPOS-YNEG+YIND TWI 1363
YScore=YScore*TOT  TWI 1364
RETURN           TWI 1365
END              TWI 1366
C               TWI 1367
SUBROUTINE DECODE(M,MM,N,NN,NDAT,IIROW,IADDR,IDAT,JJCOL) TWI 1368
C GOES THROUGH ROWS OF SUBMATRIX, BACKTRANSLATING J TO JJ, I.E. TWI 1369
C RETURNING TO COLUMN NUMBERING IN FULL MATRIX. TWI 1370
INTEGER IIROW(M),IADDR(MM),IDAT(NDAT),JJCOL(N) TWI 1371
DO 10 I=1,M      TWI 1372
II=IIROW(I)       TWI 1373
ID=IADDR(II)      TWI 1374
5 J=IDAT(ID)      TWI 1375
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IF(J.EQ.-1) GOTO 10          TWI 1376
IDAT(ID)=JJCOL(J)           TWI 1377
ID=ID+1                     TWI 1378
GOTO 5                      TWI 1379
10 CCNTINUE                 TWI 1380
RETURN                       TWI 1381
END                          TWI 1382
C
C      SUBROUTINE UPDATE(M,MM,IC,LEVEL,ICLASS,IIROW,IX)
C UPDATES CLASSIFICATION; IF DIVISION HAS FAILED, ADDS IEND TO
C CLASS NUMBER IC; FINDS NEW CLASS IC TO BE DIVIDED.           TWI 1383
INTEGER ICLASS(MM),IIROW(M),IX(M)                         TWI 1384
COMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX              TWI 1385
CCCOMMON/SWITCH/IFAIL,IDIAGR,ISTOP,IREWT                TWI 1386
IF(IFAIL.EQ.0) GOTO 20                                     TWI 1387
DO 10 I=1,M                                                 TWI 1388
II=IIROW(I)                                                TWI 1389
10 ICLASS(II)=ICLASS(II)+IEND                            TWI 1390
GOTO 40                                                    TWI 1391
C WE NOW APPLY THE DIVISION HELD IN IX(I) (=0 OR 1), AND UPDATE
C THE CLASS REGISTER FOR THE ROWS                           TWI 1392
C 20 DO 30 I=1,M                                         TWI 1393
II=IIROW(I)                                                TWI 1394
30 ICLASS(II)=ICLASS(II)*2+IX(I)                         TWI 1395
C WE NOW FIND THE NEXT CLASS TO BE DIVIDED AND SET ISTOP=1 IF
C WE HAVE COME TO THE END OF THE DIVISIONS                 TWI 1396
40 IC=IEND                                                 TWI 1397
DO 50 II=1,MM                                             TWI 1398
IIC=ICLASS(II)                                           TWI 1399
IF(IIC.LT.IC)IC=IIC                                       TWI 1400
50 CONTINUE                                                TWI 1401
IF(IC.GE.IEND) GOTO 60                                   TWI 1402
IF(2**LEVEL.GT.IC)GOTO 80                               TWI 1403
PRINT 1000,LEVEL                                         TWI 1404
1000 FORMAT('0   END OF LEVEL',I4,/)                      TWI 1405
LEVEL=LEVEL+1                                            TWI 1406
IF (LEVEL.LE.LEVMAX) GOTO 80                           TWI 1407
60 ISTOP=1                                               TWI 1408
PRINT 1001                                              TWI 1409
1001 FORMAT('0THIS IS THE END OF THE DIVISIONS REQUESTED')
DG 70 II=1,MM                                             TWI 1410
IC=ICLASS(II)                                           TWI 1411
IF(IC.GE.IEND)ICLASS(II)=IC-IEND                        TWI 1412
70 CCNTINUE                                              TWI 1413
80 RETURN                                                 TWI 1414
END                                                       TWI 1415
C
C      SUBROUTINE RECODE(M,MM,N,NN,NDAT,IC,ICLASS,IIROW,
1JJCOL,IADDR,IDAT)                                      TWI 1416
C GIVEN CLASS IC TO BE DIVIDED, CALCULATES M,N, THE NUMBERS OF
C ROWS AND COLUMNS IN SUBMATRIX DEFINED BY IC. RENUMBERS THESE
C WITH CONSECUTIVE NUMBERING, SETTING UP TRANSLATION VECTORS
C IIROW,JJCOL, TO GET BACK TO OLD NUMBERING WHEN NECESSARY.    TWI 1417
INTEGER ICLASS(MM),IIROW(MM),JJCOL(NN),IADDR(MM),IDAT(NDAT)  TWI 1418
I=0                                                       TWI 1419
TWI 1420
TWI 1421
TWI 1422
TWI 1423
TWI 1424
TWI 1425
TWI 1426
TWI 1427
TWI 1428
TWI 1429
TWI 1430
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DO 10 II=1,MM          TWI 1431
IF (ICLASS(II).NE.IC) GOTO 10      TWI 1432
I=I+1                          TWI 1433
IIROW(I)=II                      TWI 1434
10 CONTINUE                      TWI 1435
M=I                            TWI 1436
DO 20 JJ=1,NN                      TWI 1437
20 JJCOL(JJ)=0                      TWI 1438
C NOW ACCUMULATE CCOLUMN TOTALS IN JJCOL      TWI 1439
DO 30 I=1,M                      TWI 1440
II=IIROW(I)                      TWI 1441
ID=IADDR(II)                      TWI 1442
25 JJ=IDAT(ID)                      TWI 1443
IF(JJ.EQ.-1) GOTO 30      TWI 1444
JJCOL(JJ)=JJCOL(JJ)+1      TWI 1445
ID=ID+1                          TWI 1446
GOTO 25                          TWI 1447
30 CCNTINUE                      TWI 1448
J=0                            TWI 1449
DO 40 JJ=1,NN                      TWI 1450
IF(JJCOL(JJ).EQ.0) GOTO 40      TWI 1451
J=J+1                          TWI 1452
JJCOL(JJ)=J                      TWI 1453
40 CCNTINUE                      TWI 1454
N=J                            TWI 1455
C NOW GO THROUGH THE MATRIX RECODING THE COLUMNS      TWI 1456
DO 50 I=1,M                      TWI 1457
II=IIROW(I)                      TWI 1458
ID=IADDR(II)                      TWI 1459
45 JJ=IDAT(ID)                      TWI 1460
IF(JJ.EQ.-1) GOTO 50      TWI 1461
IDAT(ID)=JJCOL(JJ)      TWI 1462
ID=ID+1                          TWI 1463
GOTO 45                          TWI 1464
50 CCNTINUE                      TWI 1465
C NOW THE REVERSE CODING J TO JJ IS PUT IN JJCOL      TWI 1466
J=0                            TWI 1467
DO 60 JJ=1,NN                      TWI 1468
IF(JJCOL(JJ).EQ.0) GOTO 60      TWI 1469
J=J+1                          TWI 1470
JJCOL(J)=JJ                      TWI 1471
60 CONTINUE                      TWI 1472
RETURN                         TWI 1473
END                           TWI 1474
C
SUBROUTINE BIN(ITEM,IC)          TWI 1475
C CONVERTS CLASS NUMBER IC TO BINARY DIGITS, READY FOR PRINTING OUT      TWI 1476
INTEGER ITEM(200), BLANK,ZERO,ONE,STAR      TWI 1477
DATA BLANK/' '/,ZERO/'0'/,ONE/'1'/,STAR/'*'/
LEVEL=0                          TWI 1478
IIC=1                            TWI 1479
10 IF(IIC.GT.IC) GOTO 20      TWI 1480
IIC=IIC*2                        TWI 1481
LEVEL=LEVEL+1                      TWI 1482
GOTO 10                          TWI 1483
TWI 1484
TWI 1485
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20 DO 30 IT=1,30                                     TWI 1486
30 ITEM(IT)=BLANK                                 TWI 1487
    IT=LEVEL                                 TWI 1488
    IF(LEVEL.GT.30) IT=30                      TWI 1489
    IIC=IC                                    TWI 1490
40 IIIC=IIC/2                                    TWI 1491
    IIC=IIC-2*IIIC                           TWI 1492
    IF(IIC.EQ.0) ITEM(IT)=ZERO                TWI 1493
    IF(IIC.EQ.1) ITEM(IT)=ONE                 TWI 1494
    IIC=IIIC                                  TWI 1495
    IT=IT-1                                    TWI 1496
    IF(IT.GT.0) GOTO 40                      TWI 1497
    ITEM(1)=STAR                                TWI 1498
    RETURN                                     TWI 1499
    END                                         TWI 1500
C
C      SUBROUTINE WEIGHT(M,MM,N,NN,NDAT,X,Y,IIROW,IADDR,RRWT,ROWWGT,
1 JJCOL,CCWT,COLWGT,IDAT)                         TWI 1501
C      WEIGHTS COLUMNS ACCORDING TO HOW RARE THEY ARE. THOSE RARER THAN   TWI 1502
C      A PROPORTION FRQLIM ARE DOWNWEIGHTED IN PROPORTION TO THEIR       TWI 1503
C      SHORTFALL FROM FRQLIM, RAISED TO THE POWER ICWEXP. ROW-WEIGHTS     TWI 1504
C      FOR SUBMATRIX ARE PUT IN ROWWGT.                               TWI 1505
      REAL X(M),Y(N),RRWT(MM),ROWWGT(M),CCWT(NN),COLWGT(N)           TWI 1506
      INTEGER IIROW(M),IADDR(MM),JJCOL(NN),IDAT(NDAT)                  TWI 1507
      COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM               TWI 1508
      COMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX                   TWI 1509
      COMMON/ARBS/CWTMIN,CRLONG,CRCUT                            TWI 1510
      CWT=1.0-CWTMIN                                         TWI 1511
      IF(FRQLIM.LT.1.0E-10)FRQLIM=1.0E-10                     TWI 1512
      DO 10 J=1,N                                         TWI 1513
10  COLWGT(J)=1.0                                     TWI 1514
      TOT=0.0                                         TWI 1515
      DO 20 I=1,M                                     TWI 1516
      II=IIROW(I)                                 TWI 1517
      RR=RRWT(II)                                 TWI 1518
      ROWWGT(I)=RR                                TWI 1519
      TOT=TOT+RR                                 TWI 1520
      TOT=TOT+RR                                 TWI 1521
      TOT=TOT+RR                                 TWI 1522
20  X(I)=1.0                                         TWI 1523
      CALL XYMULT(M,MM,N,NN,NDAT,X,Y,IIROW,IADDR,ROWWGT,
1 COLWGT, IDAT)                                TWI 1524
      DO 30 J=1,N                                     TWI 1525
      JJ=JJCOL(J)                                 TWI 1526
      AY=Y(J)/TOT                                TWI 1527
      IF(AY.GE.FRQLIM)AY=FRQLIM                  TWI 1528
      COLWGT(J)=CCWT(JJ)*((AY/FRQLIM)**ICWEXP*CWT+CWTMIN)    TWI 1529
30  CCNTINUE                                         TWI 1530
      RETURN                                         TWI 1531
      END                                           TWI 1532
C
C      SUBROUTINE RA(M,MM,N,NN,NDAT,EIG,X,XX,X3,X4,X5,Y,
1 RTOT,CTOT,IIROW,IADDR,ROWWGT,COLWGT, IDAT)          TWI 1533
C      DOES RECIPROCAL AVERAGING, DERIVING EIGENVECTOR X AND EIGENVALUE   TWI 1534
C      EIG. ROWWGT, COLWGT ARE ROW AND COLUMN WEIGHTS. XX,X3,X4,X5,Y ARE   TWI 1535
C      WORKSPACE.                                         TWI 1536
      REAL X(M),XX(M),Y(N),RTOT(M),CTOT(N),ROWWGT(M),COLWGT(N)        TWI 1537
      TWI 1538
      TWI 1539
      TWI 1540
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REAL X3(M),X4(M),X5(M)                                TWI 1541
INTEGER IIROW(M),IADDR(MM),IDAT(NDAT)                 TWI 1542
COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM    TWI 1543
COMMON/SWITCH/IFAIL,IDIAGR,ISTOP,IREWT                TWI 1544
IFAIL=0                                                 TWI 1545
DO 10 I=1,M                                           TWI 1546
10 XX(I)=1.0                                         TWI 1547
DO 20 J=1,N                                           TWI 1548
20 Y(J)=1.0                                         TWI 1549
CALL XYMULT(M,MM,N,NN,NDAT,XX,CTOT,IIROW,IADDR,ROWWGT,
1COLWGT,IDAT)                                       TWI 1550
CALL YXMULT(M,MM,N,NN,NDAT,Y,RTOT,IIROW,IADDR,ROWWGT,
1COLWGT,IDAT)                                       TWI 1551
TOT=0.0                                                TWI 1552
DO 30 J=1,N                                           TWI 1553
30 TOT=TOT+CTOT(J)                                   TWI 1554
DO 40 I=1,M                                           TWI 1555
40 X(I)=FLOAT(I)                                     TWI 1556
X(1)=1.1                                             TWI 1557
C WE NOW HAVE TRIAL VECTOR X, DELIBERATELY MADE IRREGULAR TO AVOID
C NASTY COINCIDENCES                                 TWI 1558
TTOL=1.0E-5                                         TWI 1559
ICOUNT=0                                              TWI 1560
IITIM=0                                              TWI 1561
50 A=0.0                                              TWI 1562
C FIRST CENTER TO ZERO MEAN AND UNIT LENGTH          TWI 1563
DO 60 I=1,M                                           TWI 1564
60 A=A+X(I)*RTOT(I)                                 TWI 1565
A=A/TOT                                              TWI 1566
EX=0.0                                                TWI 1567
DO 70 I=1,M                                           TWI 1568
AX=X(I)-A                                           TWI 1569
EX=EX+AX*AX*RTOT(I)                                 TWI 1570
70 X(I)=AX                                         TWI 1571
EX=SQRT(EX)                                         TWI 1572
DO 80 I=1,M                                           TWI 1573
80 X(I)=X(I)/EX                                     TWI 1574
A11=0.0                                              TWI 1575
A12=0.0                                              TWI 1576
A22=0.0                                              TWI 1577
A23=0.0                                              TWI 1578
A33=0.0                                              TWI 1579
A34=0.0                                              TWI 1580
A44=0.0                                              TWI 1581
B13=0.0                                              TWI 1582
B14=0.0                                              TWI 1583
B24=0.0                                              TWI 1584
B24=0.0                                              TWI 1585
B24=0.0                                              TWI 1586
B24=0.0                                              TWI 1587
CALL XYMULT(M,MM,N,NN,NDAT,X,Y,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)                           TWI 1588
DO 81 J=1,N                                           TWI 1589
81 Y(J)=Y(J)/CTOT(J)                                 TWI 1590
CALL YXMULT(M,MM,N,NN,NDAT,Y,XX,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)                           TWI 1591
A=0.0                                                 TWI 1592
DO 82 I=1,M                                           TWI 1593
82 X(I)=A                                           TWI 1594
DO 83 I=1,M                                           TWI 1595
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```
AX=XX(I)
XX(I)=AX/RTOT(I)
A=A+AX
82 A11=A11+AX*X(I)
IF(A11.GT.TTOL) GOTO 83
IFAIL=1
RETURN
83 A=A/TOT
DO 84 I=1,M
AX=XX(I)-A-A11*X(I)
A12=A12+AX*AX*RTOT(I)
84 XX(I)=AX
A12=SQRT(A12)
DO 86 I=1,M
86 XX(I)=XX(I)/A12
IF(A12.LT.TOL) GOTO 200
IF(ICOUNT.GT.5) GOTO 200
ICOUNT=ICOUNT+1
CALL XYMULT(M,MM,N,NN,NDAT,XX,Y,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)
DO 91 J=1,N
91 Y(J)=Y(J)/CTOT(J)
CALL YXMULT(M,MM,N,NN,NDAT,Y,X3,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)
A=0.0
DO 92 I=1,M
AX=X3(I)
X3(I)=AX/RTOT(I)
A=A+AX
A22=A22+AX*XX(I)
92 B13=B13+AX*X(I)
IF(A22.LT.TTOL) GOTO 120
A=A/TOT
DO 94 I=1,M
AX=X3(I)-A-A22*XX(I)-B13*X(I)
A23=A23+AX*AX*RTOT(I)
94 X3(I)=AX
A23=SQRT(A23)
DO 96 I=1,M
96 X3(I)=X3(I)/A23
CALL XYMULT(M,MM,N,NN,NDAT,X3,Y,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)
DO 101 J=1,N
101 Y(J)=Y(J)/CTOT(J)
CALL YXMULT(M,MM,N,NN,NDAT,Y,X4,IIROW,
1IADDR,ROWWGT,COLWGT,IDAT)
A=0.0
DO 102 I=1,M
AX=X4(I)
X4(I)=AX/RTOT(I)
A=A+AX
A33=A33+AX*X3(I)
B14=B14+AX*X(I)
102 B24=B24+AX*XX(I)
IF(A33.LT.TTOL) GOTO 120
TWI 1596
TWI 1597
TWI 1598
TWI 1599
TWI 1600
TWI 1601
TWI 1602
TWI 1603
TWI 1604
TWI 1605
TWI 1606
TWI 1607
TWI 1608
TWI 1609
TWI 1610
TWI 1611
TWI 1612
TWI 1613
TWI 1614
TWI 1615
TWI 1616
TWI 1617
TWI 1618
TWI 1619
TWI 1620
TWI 1621
TWI 1622
TWI 1623
TWI 1624
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TWI 1629
TWI 1630
TWI 1631
TWI 1632
TWI 1633
TWI 1634
TWI 1635
TWI 1636
TWI 1637
TWI 1638
TWI 1639
TWI 1640
TWI 1641
TWI 1642
TWI 1643
TWI 1644
TWI 1645
TWI 1646
TWI 1647
TWI 1648
TWI 1649
TWI 1650
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```
A=A/TOT          TWI 1651
DO 104 I=1,M    TWI 1652
AX=X4(I)-(A+A33*X3(I)+B14*X(I)+B24*XX(I))
A34=A34+AX*AX*RTOT(I)  TWI 1653
104 X4(I)=AX    TWI 1654
A34=SQRT(A34)   TWI 1655
DO 106 I=1,M    TWI 1656
106 X4(I)=X4(I)/A34  TWI 1657
CALL XYMULT(M,MM,N,NN,NDAT,X4,Y,IIROW,
IIADDR,ROWWGT,COLWGT,IDAT)  TWI 1658
DO 111 J=1,N    TWI 1659
111 Y(J)=Y(J)/CTOT(J)  TWI 1660
CALL YXMULT(M,MM,N,NN,NDAT,Y,X5,IIROW,
IIADDR,ROWWGT,COLWGT,IDAT)  TWI 1661
DO 112 I=1,M    TWI 1662
112 A44=A44+X4(I)*X5(I)  TWI 1663
120 AX1=1.0      TWI 1664
AX2=0.1        TWI 1665
AX3=0.01       TWI 1666
AX4=0.001      TWI 1667
DO 130 ITIMES=1,100  TWI 1668
AXX1=A11*AX1+A12*AX2  TWI 1669
AXX2=A12*AX1+A22*AX2+A23*AX3  TWI 1670
AXX3=A23*AX2+A33*AX3+A34*AX4  TWI 1671
AXX4=A34*AX3+A44*AX4  TWI 1672
AX1=A11*AXX1+A12*AXX2  TWI 1673
AX2=A12*AXX1+A22*AXX2+A23*AXX3  TWI 1674
AX3=A23*AXX2+A33*AXX3+A34*AXX4  TWI 1675
AX4=A34*AXX3+A44*AXX4  TWI 1676
EX=SQRT(AX1**2+AX2**2+AX3**2+AX4**2)  TWI 1677
AX1=AX1/EX      TWI 1678
AX2=AX2/EX      TWI 1679
AX3=AX3/EX      TWI 1680
AX4=AX4/EX      TWI 1681
IF(ITIMES.NE.(ITIMES/5)*5) GOTO 130  TWI 1682
EXX=SQRT(EX)    TWI 1683
RESI=SQRT((AX1-AXX1/EXX)**2+(AX2-AXX2/EXX)**2+
1(AX3-AXX3/EXX)**2+(AX4-AXX4/EXX)**2)  TWI 1684
1F(RESI.LT.TOL*0.05) GOTO 135  TWI 1685
130 CONTINUE    TWI 1686
135 IITIM=IITIM+ITIMES  TWI 1687
DO 140 I=1,M    TWI 1688
140 X(I)=AX1*X(I)+AX2*XX(I)+AX3*X3(I)+AX4*X4(I)  TWI 1689
GOTO 50        TWI 1690
200 PRINT 1000,A11,ICOUNT  TWI 1691
1000 FORMAT(1X,'EIGENVALUE',F6.3,' AT ITERATION',I4)
IF(A12.GT.TOL) PRINT 1010,ICOUNT,A12,TOL  TWI 1692
1010 FORMAT(1X,'RA TROUBLE',I4,' ITERATIONS, AND RESIDUAL IS STILL ',
1F6.3,' INSTEAD OF ',F6.3,' (THE TOLERANCE)')
EIG=A11        TWI 1693
RETURN         TWI 1694
END            TWI 1695
C
SUBROUTINE XMAXMI(X,AXMAX,AXMIN,M)  TWI 1696
C CALCULATES MAXIMUM AND MINIMUM OF VECTOR X(M).  TWI 1697
TWI 1698
TWI 1699
TWI 1700
TWI 1701
TWI 1702
TWI 1703
TWI 1704
TWI 1705
```

```
REAL X(M)                                     TWI 1706
AXMAX=-1.0E10                                TWI 1707
AXMIN=-AXMAX                                 TWI 1708
DO 10 I=1,M                                    TWI 1709
AX=X(I)                                       TWI 1710
IF(AX.GT.AXMAX)AXMAX=AX                      TWI 1711
IF(AX.LT.AXMIN)AXMIN=AX                      TWI 1712
10 CONTINUE                                    TWI 1713
RETURN                                         TWI 1714
END                                            TWI 1715
C
C SUBROUTINE POLISH(M,MM,N,NN,NDAT,X,XX,Y,YY,
1 IROW,JCOL,RTOT,ROWWGT,CCWT,CCLWGT,IADDR,IDAT)    TWI 1716
C SUBROUTINE POLARIZES ORDINATION X SO THAT ORDINATION APPROXIMATES    TWI 1717
C BETTER TO A CLASSIFICATION. TO DO THIS, DERIVES TWO NEW           TWI 1718
C ORDINATIONS, TAKES THEIR SUM, AND PUTS BACK IN X.                  TWI 1719
C ORDINATION 1 IS ADDITIVE, DERIVED BY ADDITNG PREFERENCE SCORES IN    TWI 1720
C A SUMMATION IN WHICH ONLY GOOD (I.E. REASONABLY FREQUENT AND      TWI 1721
C REASONABLY PREFERENTIAL) SPECIES ARE GIVEN APPRECIABLE WEIGHT.     TWI 1722
C ORDINATION 2 TAKES THE MEAN PREFERENCE SCORE OF THE SPECIES IN     TWI 1723
C EACH STAND, UNWEIGHTED EXCEPT FOR BASIC WEIGHTS CCWT.             TWI 1724
C EACH STAND, UNWEIGHTED EXCEPT FOR BASIC WEIGHTS CCWT.             TWI 1725
C EACH STAND, UNWEIGHTED EXCEPT FOR BASIC WEIGHTS CCWT.             TWI 1726
INTEGER IROW(M),JCOL(N),IADDR(MM),IDAT(NDAT)          TWI 1727
REAL X(M),XX(M),Y(N),YY(N),RTOT(M),ROWWGT(M),CCWT(NN),COLWGT(N)    TWI 1728
COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM        TWI 1729
COMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX            TWI 1730
COMMON/ARBS/CWTMIN,CRLONG,CRCUT                     TWI 1731
CALL XMAXMI(X,AXMAX,AXMIN,M)                         TWI 1732
CUTMID=(AXMAX+AXMIN)*0.5                            TWI 1733
CUTHLF=(AXMAX-AXMIN)*0.5*CRCUT                     TWI 1734
CUT1=CUTMID-CUTHLF                                  TWI 1735
CUT2=CUTMID+CUTHLF                                  TWI 1736
CALL INDSCC(M,MM,N,NN,NDAT,X,AXPOS,AXNEG,CUT1,CUT2,    TWI 1737
1Y,YY,IROW,ROWWGT,IADDR,IDAT)                      TWI 1738
C PUT PREFERENCE SCORE IN YY AND CALCULATE APPROPRIATE COLUMN WEIGHTS    TWI 1739
PRLIM=(RATLIM-1.0)/(RATLIM+1.0)                    TWI 1740
DO 10 J=1,N                                         TWI 1741
JJ=JCOL(J)                                         TWI 1742
AY=Y(J)/AXNEG                                      TWI 1743
AYY=YY(J)/AXPOS                                     TWI 1744
PREF=(AYY-AY)/(AYY+AY)                             TWI 1745
FREQ=AYY+AY                                         TWI 1746
IF(FREQ.GT.FRQLIM) FREQ=FRQLIM                   TWI 1747
IF(PREF.GT.PRLIM)PREF=PRLIM                      TWI 1748
IF(PREF.LT.-PRLIM)PREF=-PRLIM                   TWI 1749
IF(ABS(PREF).LT.0.001)PREF=0.001                 TWI 1750
COLwGT(J)=CCWT(J)*(FREQ/FRQLIM)**ICWEXP*(ABS(PREF)/PRLIM)**IPREXP    TWI 1751
Y(J)=PREF/PRLIM                                    TWI 1752
10 CONTINUE                                         TWI 1753
CALL YXMULT(M,MM,N,NN,NDAT,Y,X,IROW,IADDR,ROWWGT,COLWGT,IDAT)    TWI 1754
DO 20 I=1,M                                         TWI 1755
20 X(I)=X(I)/ROWWGT(I)                           TWI 1756
CALL XMAXMI(X,AXMAX,AXMIN,M)                      TWI 1757
IF(ABS(AXMIN).GT.AXMAX)AXMAX=ABS(AXMIN)          TWI 1758
DO 30 I=1,M                                         TWI 1759
30 X(I)=X(I)/AXMAX                               TWI 1760
```

```
C WE NOW ABOLISH COLUMN WEIGHTS           TWI 1761
DO 40 J=1,N                           TWI 1762
  YY(J)=1.0                         TWI 1763
  JJ=JJCOL(J)                      TWI 1764
40 COLWGT(J)=CCWT(JJ)                 TWI 1765
  CALL YXMULT(M,MM,N,NN,NDAT,YY,RTOT,IROW,IADDR,
    IROWWGT,COLWGT,IDAT)          TWI 1766
  CALL YXMULT(M,MM,N,NN,NDAT,Y,XX,IIRCW,IADDR,ROWWGT,COLWGT,IDAT)
  DO 50 I=1,M                         TWI 1767
50 X(I)=X(I)+XX(I)/RTOT(I)          TWI 1768
  RETURN                               TWI 1769
  END                                  TWI 1770
C                                     TWI 1771
C                                     TWI 1772
C                                     TWI 1773
C                                     SUBROUTINE ZONEUP(X,AXMIN,CRMIN,CRMAX,AXMAX,M,MZ,MZOUT,
  1MZCRIT,IZONE)                   TWI 1774
C INPUT IS ORDINATION IN VECTOR X, WITH MAXIMUM AXMAX, MINIMUM      TWI 1775
C AXMIN. SETS UP ZONED ORDINATION, STORED IN VECTOR IZONE.          TWI 1776
C CRITICAL ZONE LIES BETWEEN CRMIN,CRMAX. TOTAL NUMBER OF           TWI 1777
C ZONES IS MZ.                                         TWI 1778
C                                         TWI 1779
  REAL X(M)                           TWI 1780
  INTEGER IZONE(M)                  TWI 1781
  MZCRIT=MZ-2*MZOUT                TWI 1782
  SMALL=1.0E-10                     TWI 1783
  SEG1=(CRMIN-AXMIN)/(FLOAT(MZOUT)+SMALL)+SMALL        TWI 1784
  SEG2=(CRMAX-CRMIN)/(FLOAT(MZCRIT)+SMALL)+SMALL       TWI 1785
  SEG3=(AXMAX-CRMAX)/(FLOAT(MZOUT)+SMALL)+SMALL        TWI 1786
  DO 10 I=1,M                        TWI 1787
    AX=X(I)
    IF(AX.GE.CRMIN) GOTO 3          TWI 1788
    IZ=IFIX((AX-AXMIN)/SEG1)+1     TWI 1789
    IF(IZ.GT.MZOUT) IZ=MZOUT       TWI 1790
    IF(IZ.LT.1) IZ=1               TWI 1791
    GOTO 8                          TWI 1792
3  IF(AX.LE.CRMAX) GOTO 6          TWI 1793
  IZ=IFIX((AX-CRMAX)/SEG3)+1      TWI 1794
  IF(IZ.LT.1) IZ=1               TWI 1795
  IF(IZ.GT.MZOUT) IZ=MZOUT       TWI 1796
  IZ=IZ+MZ-MZOUT                 TWI 1797
  GOTO 8                          TWI 1798
6  IZ=IFIX((AX-CRMIN)/SEG2)+1      TWI 1799
  IF(IZ.LT.1) IZ=1               TWI 1800
  IF(IZ.GT.MZCRIT) IZ=MZCRIT    TWI 1801
  IZ=IZ+MZOUT                    TWI 1802
8  IZONE(I)=IZ                     TWI 1803
10 CCNTINUE                         TWI 1804
  RETURN                             TWI 1805
  END                                TWI 1806
C                                     TWI 1807
C                                     TWI 1808
C                                     SUBROUTINE TOPIND(N,NN,MIND,FEEBLE,Y,INDORD,INDSIG,
  1JJCOL,INDPOT,JNAM,NIND,LIND)   TWI 1809
C INPUT IS INDICATOR VALUES STORED IN Y. SETS UP NIND BEST      TWI 1810
C INDICATORS IN INDORD, WITH THEIR SIGN IN INDSIG. IY IS WORKSPACE. TWI 1811
C NIND IS SET TO MIND UNLESS THERE IS SHORTAGE OF GOOD INDICATORS. TWI 1812
C                                         TWI 1813
  REAL Y(N)                           TWI 1814
  INTEGER INDORD(N),INDSIG(MIND),JJCOL(N),INDPOT(NN)          TWI 1815
```

```
INTEGER JNAM(NN)          TWI 1816
INTEGER LIND(10)           TWI 1817
IBIG=10000                 TWI 1818
DO 10 J=1,N                 TWI 1819
10 INDORD(J)=-IBIG*IFIX(ABS(Y(J)*500.0))-IBIG+J   TWI 1820
CALL ISORT(INDORD,N)       TWI 1821
DO 20 J=1,N                 TWI 1822
IV=-INDORD(J)              TWI 1823
20 INDORD(J)=(IV/IBIG)*IBIG-IV+IBIG                TWI 1824
C INDORD NOW CONTAINS ORDERING BUT MUST BE DOCTORED TO TRUNCATE OFF    TWI 1825
C POOR AND DISALLOWED INDICATORS                         TWI 1826
NIND=0                  TWI 1827
DO 60 IND=1,N             TWI 1828
IF(NIND.EQ.MIND) GOTO 60  TWI 1829
J=INDORD(IND)             TWI 1830
JJ=JJCOL(J)               TWI 1831
JJP=INDPOT(JJ)             TWI 1832
IF(JJP.LT.1) GOTO 60      TWI 1833
IP=INDPOT(JJP)             TWI 1834
IF(IP.LT.1) GOTO 60       TWI 1835
IF(ABS(Y(J)).LT.FEEBLE) GOTO 60  TWI 1836
IL=JNAM(JJ)               TWI 1837
IF(LIND(IL).EQ.0) GOTO 60  TWI 1838
NIND=NIND+1               TWI 1839
INDORD(NIND)=J             TWI 1840
INDPOT(JJP)=-INDPOT(JJP)  TWI 1841
INDSIG(NIND)=1              TWI 1842
IF(Y(J).LT.0) INDSIG(NIND)=-1  TWI 1843
60 CCNTINUE               TWI 1844
IF(NIND.EQ.0) GOTO 80      TWI 1845
DO 70 IND=1,NIND           TWI 1846
J=INDORD(IND)             TWI 1847
JJ=JJCOL(J)               TWI 1848
JJP=IABS(INDPOT(JJ))      TWI 1849
INDPOT(JJP)=JJP            TWI 1850
70 CONTINUE                 TWI 1851
80 RETURN                   TWI 1852
END                        TWI 1853
C
SUBROUTINE CODESC(M,MM,N,NN,NDAT,NIND,INDORD,IX,IY,
IIROW,IADDR,IDAT)        TWI 1854
C STARTING WITH NIND BEST INDICATORS, STORES INDICATOR-COMPOSITION      TWI 1855
C OF SAMPLES IN CODED FORM IN IX. THE FIRST (BEST) INDICATOR IS GIVEN     TWI 1856
C A CODED SCORE CF 1, THE NEXT A SCORE OF 2, THE NTH A SCORE 2**{N-1}.    TWI 1857
C CODE NUMBERS ARE ADDED TO OBTAIN CODED INDICATOR-COMPOSITION.          TWI 1858
INTEGER INDORD(N),IX(M),IY(N),IIROW(M),IADDR(MM),IDAT(NDAT)           TWI 1859
DO 10 J=1,N                 TWI 1860
10 IY(J)=0                  TWI 1861
IIY=1                      TWI 1862
DO 20 IND=1,NIND             TWI 1863
J=INDORD(IND)               TWI 1864
IY(J)=IIY                  TWI 1865
20 IIY=IIY*2                 TWI 1866
DO 30 I=1,M                  TWI 1867
II=IIROW(I)                 TWI 1868
30 CONTINUE                  TWI 1869
END                        TWI 1870
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IIX=0          TWI 1871
ID=IADDR(II)   TWI 1872
25 J=IDAT(ID)   TWI 1873
    IF(J.EQ.-1) GOTO 26   TWI 1874
    IIX=IIX+IY(J)   TWI 1875
    ID=ID+1   TWI 1876
    GOTO 25   TWI 1877
26 IX(I)=IIX   TWI 1878
30 CCNTINUE   TWI 1879
    RETURN   TWI 1880
    END   TWI 1881
C   TWI 1882
    SUBROUTINE TABLE(M,MMZ,MMS,MIND,IX,IZONE,INDSIG,IPICT,
1INDTRU,IIND)   TWI 1883
C CODED INDICATOR-COMPOSITION IS IN IX. CALCULATES MATRIX IPICT,
C RELATING ZONES TO INDICATOR-SCORES. ONLY IIND TOP INDICATORS ARE   TWI 1884
C USED. ISHIFT IS ADDED TO INDICATOR-SCORES TO KEEP THEM POSITIVE.   TWI 1885
C IF THE SWITCH INDTRU=1, SUBROUTINE ALSO REPLACES CODED INDICATOR-   TWI 1886
C COMPOSITION BY THE TRUE INDICATOR-SCORES.   TWI 1887
    INTEGER IX(M),IZONE(M),INDSIG(MIND),IPICT(MMZ,MMS)
    COMMON/PICT/MZ,MS,IZD,IIZD,ISD,ISHIFT,MZIND,MZCRIT,MZOUT,NIND   TWI 1888
    MS=IIND+1   TWI 1889
    ISHIFT=1   TWI 1890
    DO 10 IND=1,IIND   TWI 1891
    IF(INDSIG(IND).LT.0) ISHIFT=ISHIFT+1   TWI 1892
10 CONTINUE   TWI 1893
    DO 20 IZ=1,MZ   TWI 1894
    DO 20 IS=1,MS   TWI 1895
20 IPICT(IZ,IS)=0   TWI 1896
    DO 30 I=1,M   TWI 1897
    IZ=IZONE(I)   TWI 1898
    IS=ISHIFT   TWI 1899
    IIX=IX(I)   TWI 1900
    DO 25 IND=1,IIND   TWI 1901
    IIIX=IIX/2   TWI 1902
    IS=IS+INDSIG(IND)*(IIX-IIIIX*2)   TWI 1903
25 IIX=IIIIX   TWI 1904
    IF(INDTRU.EQ.1)IX(I)=IS   TWI 1905
30 IPICT(IZ,IS)=IPICT(IZ,IS)+1   TWI 1906
    RETURN   TWI 1907
    END   TWI 1908
C   TWI 1909
    SUBROUTINE FIND(MMZ,MMS,IPICT,IZD1,ISD1,MISCL)   TWI 1910
C INPUT IS MATRIX IPICT, RELATING INDICATOR SCORES TO ZONES.   TWI 1911
C SUBROUTINE FINDS CLASSIFICATION CENTER IZD1,ISD1 THAT MINIMIZES   TWI 1912
C NUMBER OF MISCLASSIFICATIONS MISCL.   TWI 1913
    INTEGER IPICT(MMZ,MMS)
    COMMON/PICT/MZ,MS,IZD,IIZD,ISD,ISHIFT,MZIND,MZCRIT,MZOUT,NIND   TWI 1914
    MINZ=MZOUT+MZIND   TWI 1915
    MAXZ=MZOUT+MZCRIT   TWI 1916
C WE FORM THE TOTALS IN IPICT OF THE ELEMENTS THAT LIE BELOW AND TO   TWI 1917
C THE LEFT OF A GIVEN ONE. I.E. THE NEW VALUE OF IPICT(IZ,IS) IS THE   TWI 1918
C SUM OF ALL THE OLD VALUES IF IPICT(IIZ,IIS) FOR WHICH IIZ IS LESS   TWI 1919
C THAN OR EQUAL TO IZ AND IIS IS LESS THAN OR EQUAL TO IS.   TWI 1920
    DC 10 IS=2,MS   TWI 1921
    TWI 1922
    TWI 1923
    TWI 1924
    TWI 1925
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10 IPICT(1,IS)=IPICT(1,IS-1)+IPICT(1,IS)           TWI 1926
    DO 20 IZ=2,MZ                                     TWI 1927
20 IPICT(IZ,1)=IPICT(IZ-1,1)+IPICT(IZ,1)          TWI 1928
    DO 30 IS=2,MS                                     TWI 1929
    DO 30 IZ=2,MZ                                     TWI 1930
    IIS=IS-1                                         TWI 1931
    IIIZ=IZ-1                                         TWI 1932
30 IPICT(IZ,IS)=IPICT(IIIZ,IS)+IPICT(IZ,IIS)-     TWI 1933
    1IPICT(IIIZ,IIS)+IPICT(IZ,IS)                  TWI 1934
    MISCL=10000                                       TWI 1935
    DO 40 IZ=MINZ,MAXZ                               TWI 1936
    DO 40 IS=1,MS                                     TWI 1937
    IIIZ=IZ-MZIND                                    TWI 1938
    MISS=IPICT(IIIZ,MS)-IPICT(IIIZ,IS)+IPICT(MZ,IS)-IPICT(IZ,IS)
    A=FLOAT(IPICT(IIIZ,MS))                         TWI 1940
    B=FLOAT(IPICT(MZ,MS)-IPICT(IZ,MS))             TWI 1941
    C=ABS((A-B)/(A+B))                             TWI 1942
    IA=IABS(IS-ISHIFT)                            TWI 1943
    IB=IABS(1+MZ-IZ-IIIZ)                          TWI 1944
    IF(MISS-MISCL) 35,36,40                         TWI 1945
35 MISCL=MISS                                       TWI 1946
    IZD1=IZ                                         TWI 1947
    ISD1=IS                                         TWI 1948
    IIIA=IA                                         TWI 1949
    IIIB=IB                                         TWI 1950
    CC=C                                           TWI 1951
    GOTO 40                                         TWI 1952
36 IF(C-CC) 35,38,40                               TWI 1953
38 IF(IA.LT.IIA) GOTO 35                           TWI 1954
    IF(IB.LT.IIB) GOTO 35                           TWI 1955
40 CONTINUE                                         TWI 1956
    RETURN                                           TWI 1957
    END                                              TWI 1958
C
C      SUBROUTINE REPORT(M,MM,N,NN,NDAT,MMZ,MMS,IC,X,XX,ROWWGT,
C      1Y,YY,COLWGT,IX,IZONE,IIROW,IY,JJCOL,IADDR,IDAT,IPICT,
C      2INAME1,INAME2,JNAME1,JNAME2,JNAM)
C REPORTS ON DICHOTOMY. TRUE INDICATOR SCORES ARE NOW IN IX.
C MATRIX IPICT CONTAINS DIAGRAM OF DICHOTOMY, RELATING ZONES
C TO INDICATOR SCORES.
    REAL X(M),XX(M),ROWWGT(M),Y(N),YY(N),COLWGT(N)
    INTEGER ITEM(200)
    INTEGER INAME1(MM),INAME2(MM),JNAME1(NN),JNAME2(NN),JNAM(NN)
    INTEGER IX(M),IZONE(M),IIROW(M),IY(N),JJCOL(N),
    1IADDR(MM),IDAT(NDAT),IPICT(MMZ,MMS),NCAT(6)
    COMMON/PICT/MZ,MS,IZD,IIZD,ISD,ISHIFT,MZIND,MZCRIT,MZOUT,NIND
    COMMON/LIMS/RARE,FEEBLE,FRQLIM,TOL,RATLIM,REPLIM
    COMMON/SWITCH/IFAIL,IDIAGR,ISTOP,IREWT
    COMMON/IARBS/ICWEXP,IEND,MMIN,IPREXP,LEVMAX
    COMMON/WORK/ITEM
    COMMON/SECOND/ISEC
    DO 10 ICAT=1,6
10 NCAT(ICAT)=0
    DO 20 I=1,M
    IZ=IZONE(I)
    TWI 1959
    TWI 1960
    TWI 1961
    TWI 1962
    TWI 1963
    TWI 1964
    TWI 1965
    TWI 1966
    TWI 1967
    TWI 1968
    TWI 1969
    TWI 1970
    TWI 1971
    TWI 1972
    TWI 1973
    TWI 1974
    TWI 1975
    TWI 1976
    TWI 1977
    TWI 1978
    TWI 1979
    TWI 1980
```

```
IS=IX(1)          TWI 1981
IF(IZ.GT.IIZD) GOTO 13  TWI 1982
IF(IS.GT.ISD) GOTO 12  TWI 1983
ICAT=1            TWI 1984
GOTO 18            TWI 1985
12 ICAT=3          TWI 1986
GOTO 18            TWI 1987
13 IF(IZ.GT.IZD) GOTO 16  TWI 1988
IF(IS.GT.ISD) GOTO 15  TWI 1989
ICAT=2            TWI 1990
GOTO 18            TWI 1991
15 ICAT=5          TWI 1992
GOTO 18            TWI 1993
16 IF(IS.GT.ISD) GOTO 17  TWI 1994
ICAT=6            TWI 1995
GOTO 18            TWI 1996
17 ICAT=4          TWI 1997
18 X(I)=FLOAT(ICAT)  TWI 1998
NCAT(ICAT)=NCAT(ICAT)+1  TWI 1999
20 CONTINUE        TWI 2000
NNEG=NCAT(1)+NCAT(2)+NCAT(3)  TWI 2001
NPOS=NCAT(4)+NCAT(5)+NCAT(6)  TWI 2002
CALL INDSCG(M,M4,N,NN,NDAT,X,AXPOS,AXNEG,3.5,3.5,
1Y,YY,IROW,ROWWGT,IADDR,IDAT)  TWI 2003
PRLIM=(REPLIM-1.0)/(REPLIM+1.0)-0.0001  TWI 2004
DO 30 J=1,N        TWI 2005
AY=Y(J)/AXNEG      TWI 2006
AYY=YY(J)/AXPOS    TWI 2007
IIY=0              TWI 2008
IF(AY.GE.RARE) IIY=1  TWI 2009
IF(AYY.GE.RARE) IIY=1  TWI 2010
PREF=(AYY-AY)/(AYY+AY)  TWI 2011
IF(PREF.GT.PRLIM) GOTO 25  TWI 2012
IF(PREF.LT.-PRLIM) GOTO 23  TWI 2013
IY(J)=IIY          TWI 2014
GOTO 30            TWI 2015
23 IY(J)=2*IIY      TWI 2016
GOTO 30            TWI 2017
25 IY(J)=3*IIY      TWI 2018
30 CONTINUE        TWI 2019
C WE NOW HAVE A VARIABLE IY THAT IS 0 FOR RARITIES, 3 FOR POSITIVE
C PREFERENTIALS, 2 FOR NEGATIVE PREFERENTIALS, 1 FOR INDIFFERENTS.  TWI 2020
IF(NIND.EQ.0) GOTO 60  TWI 2021
ISD1=ISD-ISHIFT     TWI 2022
ISD2=ISD1+1          TWI 2023
PRINT 1050,ISD1,ISD2  TWI 2024
1050 FORMAT(1X,'MAXIMUM INDICATOR SCORE FOR NEGATIVE GROUP ',I4,
1'           MINIMUM INDICATOR SCORE FOR POSITIVE GROUP ',I4)
IF(IDIAGR.EQ.1) CALL DIAGR(IPICT,MMZ,MMS)  TWI 2025
60 IIC=IC*2          TWI 2026
CALL BIN(ITEM,IIC)   TWI 2027
PRINT 1002,IIC,NNEG,(ITEM(IT),IT=1,30)  TWI 2028
1002 FORMAT('0ITEMS IN NEGATIVE GROUP',I4,' (N=',I5,'),',9X,'I.E. GRCUP ',30A1)
CALL XLIST(M,MM,X,IROW,1,2,3,INAME1,INAME2)  TWI 2029
                                         TWI 2030
                                         TWI 2031
                                         TWI 2032
                                         TWI 2033
                                         TWI 2034
                                         TWI 2035
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IF(NIND.EQ.0) GOTO 70           TWI 2036
NOUT=NCAT(2)                   TWI 2037
IF(NOUT.EQ.0) GOTO 65          TWI 2038
PRINT 1003,NOUT                TWI 2039
1003 FORMAT('OBORDERLINE NEGATIVES   (N=',I5,')')
CALL XLIST(M,MM,X,IIRW,2,2,2,INAME1,INAME2)    TWI 2040
65 NCUT=NCAT(3)                 TWI 2041
IF(NOUT.EQ.0) GOTO 70          TWI 2042
PRINT 1004,NOUT                TWI 2043
1004 FORMAT('OMISCLASSIFIED NEGATIVES (N=',I5,')')
CALL XLIST(M,MM,X,IIRW,3,3,3,INAME1,INAME2)    TWI 2044
70 IIC=IC*2+1                  TWI 2045
CALL BIN(ITEM,IIC)             TWI 2046
PRINT 1005,IIC,NPOS,(ITEM(IT),IT=1,30)        TWI 2047
1005 FORMAT('OITEMS IN POSITIVE GROUP',I4,' (N=',
1I5,)',9X,'I.E. GROUP ',30A1)    TWI 2048
CALL XLIST(M,MM,X,IIRW,4,5,6,INAME1,INAME2)    TWI 2049
IF(NIND.EQ.0) GOTO 80          TWI 2050
NOUT=NCAT(5)                  TWI 2051
IF(NOUT.EQ.0) GOTO 75          TWI 2052
PRINT 1006,NOUT                TWI 2053
1006 FORMAT('OBORDERLINE POSITIVES  (N=',I5,')')
CALL XLIST(M,MM,X,IIRW,5,5,5,INAME1,INAME2)    TWI 2054
75 NOUT=NCAT(6)                 TWI 2055
IF(NOUT.EQ.0) GOTO 80          TWI 2056
PRINT 1007,NOJT                TWI 2057
1007 FORMAT('OMISCLASSIFIED POSITIVES (N=',I5,')')
CALL XLIST(M,MM,X,IIRW,6,6,6,INAME1,INAME2)    TWI 2058
C WE NOW LIST OUT THE COMMONER SPECIES ACCORDING AS THEY ARE
C PREFERENTIAL OR NEUTRAL      TWI 2059
80 IF(ISEC.EQ.2) GOTO 100       TWI 2060
CALL INDSCO(M,MM,N,NN,NDAT,X,AXPOS,AXNEG,3.5,3.5,
1Y,YY,IIRW,ROWWGT,IADDR,IDAT)    TWI 2061
PRINT 1010                      TWI 2062
1010 FORMAT('ONEGATIVE PREFERENTIALS')
CALL YLIST(N,NN,Y,YY,IY,JCOL,2,JNAME1,JNAME2,JNAM)    TWI 2063
PRINT 1011                      TWI 2064
1011 FORMAT('OPOSITIVE PREFERENTIALS')
CALL YLIST(N,NN,Y,YY,IY,JCOL,3,JNAME1,JNAME2,JNAM)    TWI 2065
PRINT 1012                      TWI 2066
1012 FORMAT('ONON-PREFERENTIALS')
CALL YLIST(N,NN,Y,YY,IY,JCOL,1,JNAME1,JNAME2,JNAM)    TWI 2067
100 RETURN                     TWI 2068
END                           TWI 2069
C
SUBROUTINE XYMULT(M,MM,N,NN,NDAT,X,Y,IIRW,IADDR,ROWWGT,
1COLWGT,IDAT)                 TWI 2070
C FORMS MATRIX PRODUCT Y=AX, APPLYING ROW AND COLUMN WEIGHTS
C ROWWGT,COLWGT.                TWI 2071
REAL X(M),Y(N),ROWWGT(M),COLWGT(N)    TWI 2072
INTEGER IIRW(M),IADDR(MM),IDAT(NDAT)    TWI 2073
DO 10 J=1,N                      TWI 2074
10 Y(J)=0.0                       TWI 2075
DO 20 I=1,M                      TWI 2076
II=IIRW(I)                      TWI 2077
20
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AX=X(I)*ROWWGT(I)          TWI 2091
ID=IADDR(II)                TWI 2092
15 J=IDAT(ID)               TWI 2093
  IF (J.EQ.-1) GOTO 20       TWI 2094
  Y(J)=Y(J)+AX               TWI 2095
  ID=ID+1                   TWI 2096
  GOTO 15                   TWI 2097
20 CONTINUE                  TWI 2098
  DO 30 J=1,N                TWI 2099
30 Y(J)=Y(J)*COLWGT(J)      TWI 2100
  RETURN                     TWI 2101
  END                        TWI 2102
C
C   SUBROUTINE YXMULT(M,MM,N,NN,NDAT,Y,X,IIROW,IADDR,
1 ROWWGT,COLWGT,IDAT)      TWI 2103
C FORMS MATRIX PRODUCT X=AY, WEIGHTED AS SUBROUTINE XYMULT.    TWI 2104
  REAL Y(N),X(M),ROWWGT(M),COLWGT(N)                         TWI 2105
  INTEGER IIROW(M),IADDR(MM),IDAT(NDAT)                      TWI 2106
  DO 10 J=1,N                                         TWI 2107
10 Y(J)=Y(J)*COLWGT(J)                                     TWI 2108
  DO 20 I=1,M                                         TWI 2109
  II=IIROW(I)                                         TWI 2110
  AX=0.0                                              TWI 2111
  ID=IADDR(II)                                         TWI 2112
15 J=IDAT(ID)                                           TWI 2113
  IF (J.EQ.-1) GOTO 16                                     TWI 2114
  AX=AX+Y(J)                                         TWI 2115
  ID=ID+1                                         TWI 2116
  GOTO 15                                         TWI 2117
16 X(I)=AX*ROWWGT(I)                                     TWI 2118
20 CCNTINUE                                         TWI 2119
  DO 30 J=1,N                                         TWI 2120
30 Y(J)=Y(J)/COLWGT(J)                                    TWI 2121
  RETURN                                         TWI 2122
  END                                         TWI 2123
C
C   SUBROUTINE INDSO(M,MM,N,NN,NDAT,X,AXPOS,AXNEG,
1 CUT1,CUT2,Y,YY,IIRCW,ROWWGT,IADDR,IDAT)      TWI 2124
C INPUT IS ORDINATION IN X. OUTPUT IS NEGATIVE INDICATOR SCORES IN Y,    TWI 2125
C POSITIVE INDICATOR SCORES IN YY. AXPOS,AXNEG ARE ROW-WEIGHTED SUMS    TWI 2126
C OF POSITIVE AND NEGATIVE ELEMENTS. CUT1,CUT2 ARE CUT-POINTS FOR    TWI 2127
C ZONE OF INTERMEDIATENESS.                                         TWI 2128
  INTEGER IIROW(M),IADDR(MM),IDAT(NDAT)      TWI 2129
  REAL X(M),Y(N),YY(N),ROWWGT(M)                 TWI 2130
  CUTMID=(CUT1+CUT2)/2.0                         TWI 2131
  CUTHLF=(CUT2-CUT1)/2.0+1.0E-10                TWI 2132
  AXPOS=0.0                                         TWI 2133
  AXNEG=0.0                                         TWI 2134
  DO 10 J=1,N                                         TWI 2135
  Y(J)=0.0                                         TWI 2136
10 YY(J)=0.0                                         TWI 2137
  DO 30 I=1,M                                         TWI 2138
  II=IIROW(I)                                         TWI 2139
  ID=IADDR(II)                                         TWI 2140
  AX=(X(I)-CUTMID)/CUTHLF                         TWI 2141
  DO 30 I=1,M                                         TWI 2142
  II=IIROW(I)                                         TWI 2143
  ID=IADDR(II)                                         TWI 2144
  AX=(X(I)-CUTMID)/CUTHLF                         TWI 2145
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IF(AX.GT.1.0)AX=1.0          TWI 2146
IF(AX.LT.-1.0)AX=-1.0        TWI 2147
IF(AX.GT.0.0) GOTO 26        TWI 2148
AX=AX*ROWWGT(I)
AXNEG=AXNEG-AX
25 J=IDAT(ID)
IF(J.EQ.-1) GOTO 30
Y(J)=Y(J)-AX
ID=ID+1
GOTO 25
26 AX=AX*ROWWGT(I)
AXPOS=AXPOS+AX
27 J=IDAT(ID)
IF(J.EQ.-1) GOTO 30
YY(J)=YY(J)+AX
ID=ID+1
GOTO 27
30 CONTINUE
RETURN
END

C
SUBROUTINE DIAGR(IPICT,MMZ,MMS)
C PRINTS OUT DIAGRAM OF RELATION BETWEEN ZONES AND INDICATOR-SCORES
C STORED IN MATRIX IPICT.  *'S ARE USED AS PUNCTUATION.
INTEGER IPICT(MMZ,MMS)
CCCOMMON/PICT/MZ,MS,IZD,IIZD,ISD,ISHIFT,MZIND,MZCRIT,MZOUT,NIND
IBIG=1000000
MZZ=MZ+2
DO 10 IZ=1,MZZ
IPICT(IZ,MS+1)=IBIG
10 IPICT(IZ,MS+2)=IZ
MSS=MS+2
DO 20 IS=1,MSS
IPICT(MZ+1,IS)=IBIG
20 IPICT(MZ+2,IS)=IS-ISHIFT
IPICT(MZZ,MSS)=IBIG
IPICT(MZ+2,MS+1)=IBIG
DO 30 IS=1,MSS
IIZ=MZ-IZD+2
IZ=MZ+4
DO 21 IIZ=1,IIZ
IPICT(IZ,IS)=IPICT(IZ-2,IS)
21 IZ=IZ-1
IPICT(IZ,IS)=IBIG
IZ=IZ-1
IIZ=IZD-IIZD
DO 22 IIZ=1,IIZ
IPICT(IZ,IS)=IPICT(IZ-1,IS)
22 IZ=IZ-1
IPICT(IZ,IS)=IBIG
30 CCNTINUE
MZZ=MZ+4
DO 40 IZ=1,MZZ
IS=MS+3
IIS=MS-ISD+2
TWI 2151
TWI 2152
TWI 2153
TWI 2154
TWI 2155
TWI 2156
TWI 2157
TWI 2158
TWI 2159
TWI 2160
TWI 2161
TWI 2162
TWI 2163
TWI 2164
TWI 2165
TWI 2166
TWI 2167
TWI 2168
TWI 2169
TWI 2170
TWI 2171
TWI 2172
TWI 2173
TWI 2174
TWI 2175
TWI 2176
TWI 2177
TWI 2178
TWI 2179
TWI 2180
TWI 2181
TWI 2182
TWI 2183
TWI 2184
TWI 2185
TWI 2186
TWI 2187
TWI 2188
TWI 2189
TWI 2190
TWI 2191
TWI 2192
TWI 2193
TWI 2194
TWI 2195
TWI 2196
TWI 2197
TWI 2198
TWI 2199
TWI 2200
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DO 31 IIIS=1,IIS          TWI 2201
  IPICT(IZ,IS)=IPICT(IZ,IS-1)
31 IS=IS-1                TWI 2202
  IPICT(IZ,IS)=IBIG       TWI 2203
40 CCNTINUE               TWI 2204
C WE NOW HAVE THE DESIRED INFORMATION IN IPICT; ALL WE NOW NEED TO    TWI 2205
C DO IS TO PRINT IT OUT
  MSS=MSS+3              TWI 2206
  IS=MSS                  TWI 2207
  DO 50 IIIS=1,MSS        TWI 2208
    PRINT 1000,(IPICT(IZ,IS),IZ=1,MZZ)      TWI 2209
1000 FORMAT(1X,60I2)        TWI 2210
  IS=IS-1                TWI 2211
50 CCNTINUE               TWI 2212
  RETURN                  TWI 2213
  END                     TWI 2214
C
C   SUBROUTINE XLIST(M,MM,X,IIROW,ICAT1,ICAT2,           TWI 2215
  1ICAT3,INAME1,INAME2)      TWI 2216
C PRINTS OUT LIST OF INDIVIDUALS (ROWS) FOR WHICH X=ICAT1,2,3.      TWI 2217
  REAL X(M)                TWI 2218
  INTEGER IIROW(M),ITEM(200)  TWI 2219
  INTEGER INAME1(MM),INAME2(MM)  TWI 2220
  COMMON/WORK/ITEM          TWI 2221
  COMMON/SECOND/ISEC         TWI 2222
  NITEM=12                 TWI 2223
  IF(ISEC.EQ.2) NITEM=11      TWI 2224
  ITEND=2*(NITEM-1)          TWI 2225
  IT=0                      TWI 2226
  I=0                      TWI 2227
3  I=I+1                   TWI 2228
  IIX=IFIX(X(I)+0.01)        TWI 2229
  IF((IIX-ICAT1)*(IIX-ICAT2)*(IIX-ICAT3).NE.0) GOTO 5      TWI 2230
  II=IIROW(I)                TWI 2231
  IT=IT+1                   TWI 2232
  ITEM(IT)=INAME1(II)        TWI 2233
  IT=IT+1                   TWI 2234
  ITEM(IT)=INAME2(II)        TWI 2235
5  IF(I.EQ.M) GOTO 6          TWI 2236
  IF(IT.LE.ITEND) GOTO 3      TWI 2237
  IF(ISEC.EQ.2) GOTO 10        TWI 2238
  PRINT 1000,(ITEM(IIT),IIT=1,IT)      TWI 2239
  IT=0                      TWI 2240
  GOTO 3                     TWI 2241
6  IF(ISEC.EQ.2) GOTO 15      TWI 2242
  IF(IT.NE.0) PRINT 1000,(ITEM(IIT),IIT=1,IT)      TWI 2243
1000 FORMAT(12(1X,2A4,1X))      TWI 2244
  RETURN                     TWI 2245
10  PRINT 1001,(ITEM(IIT),IIT=1,IT)      TWI 2246
  IT=0                      TWI 2247
  GOTO 3                     TWI 2248
15  IF(IT.NE.0) PRINT 1001,(ITEM(IIT),IIT=1,IT)      TWI 2249
1001 FORMAT(11(2X,A4,1X,A4))      TWI 2250
  RETURN                     TWI 2251
  END                       TWI 2252
                                TWI 2253
                                TWI 2254
                                TWI 2255
```

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C          SUBROUTINE YLIST(N,NN,Y1,Y2,IY,JJCOL,IIY,JNAME1,JNAME2,JNAM)      TWI 2256
C PRINTS OUT LIST OF ATTRIBUTES (COLUMNS) FOR WHICH IY(J)=IIY,           TWI 2257
C TOGETHER WITH NUMBERS OF OCCURRENCES ON 2 SIDES OF DICHOTOMY.           TWI 2258
REAL Y1(N),Y2(N)               TWI 2259
INTEGER IY(N),JJCOL(N),ITEM(200)           TWI 2260
INTEGER JNAME1(NN),JNAME2(NN),JNAM(NN)       TWI 2261
COMMON/WORK/ITEM              TWI 2262
NITEM=6                         TWI 2263
ITEND=5*(NITEM-1)             TWI 2264
IT=0                           TWI 2265
J=0                           TWI 2266
3 J=J+1                         TWI 2267
IF(IY(J).NE.IIY) GOTO 5          TWI 2268
JJ=JJCOL(J)                     TWI 2269
IT=IT+1                         TWI 2270
ITEM(IT)=JNAME1(JJ)            TWI 2271
IT=IT+1                         TWI 2272
ITEM(IT)=JNAME2(JJ)            TWI 2273
IT=IT+1                         TWI 2274
ITEM(IT)=JNAM(JJ)              TWI 2275
IT=IT+1                         TWI 2276
ITEM(IT)=IT+1                  TWI 2277
ITEM(IT)=IFIX(Y1(J)+0.001)     TWI 2278
IT=IT+1                         TWI 2279
ITEM(IT)=IFIX(Y2(J)+0.001)     TWI 2280
5 IF(J.EQ.N) GOTO 6            TWI 2281
IF(IT.LE.ITEND) GOTO 3          TWI 2282
PRINT 1000,(ITEM(IIT),IIT=1,IT)   TWI 2283
IT=0                           TWI 2284
GOTO 3                          TWI 2285
6 IF(IT.NE.0) PRINT 1000,(ITEM(IIT),IIT=1,IT)   TWI 2286
1000 FORMAT(6(1X,A4,1X,A4,I1,'(',I3,',',I3,')')) TWI 2287
RETURN                         TWI 2288
END                           TWI 2289
```