Session 3 - SPSS¹

Saving a SPSS data file. Reading a SPSS data file. Another way to read a hierarchical file. Matching (merging) SPSS data files. More on errors and error detection.

Saving and Accessing SPSS data files (AKA SPSS systems file, SPSS file)

So far we have used raw data – numbers that you can look at. An SPSS data file is different. You cannot look at it on the disk and read it. It is stored in binary format, and has special information that only SPSS (and other programs trained to understand SPSS). It has all of the information that was given to SPSS about variables before the file was saved. For example, if variable names were provided, or new variables were created, this information is stored in the file.

To save an SPSS data file, the following code is used.

save outfile = 'd:/fullpath/filename.sav'.

The file can be accessed with the command:

get file = 'd:/fullpath/filename.sav'.

More about Hierarchical Files, and Other Ways to Read Them.

Reading a hierarchical file is sometimes called "rectangularizing" it. In the small example we have been using, we have three household records, and seven person records. We end up with seven records, but more variables. For a tiny data set like this, storage and processing is not an issue. But, with real census data sets, storage can become an important issue. For example, with one of the 1990 1% IPUMS files (Integrated Public Use Microdata Series, see http://www.ipums.umn.edu/), there are 1,105,583 household records with 105 variables, and 2,500,052 person records with 149 variables. It takes much less space to store the 105 household variables 1.1 million times than it does to store them 2.5 million times. If these files were rectangularized, they would take far more space to store. Another options is to store the person and household records separately, and merge them when you need both household and person records. The following program reads the data, and saves two systems files.

```
* prog4.sps.
title 'prog4.sps' .
* Read the data and save separate files for Household and Person Records.
** The data (as a comment) follow:
h011360010
p0101125
p0102222
p0103302
p0104301
h023130010
p0201145
p0202325
h034530010
p0301166
** READ and save HOUSEHOLD cases and VARIABLES .
input program .
DATA LIST file = 'c:\all\spssclass\progs\prog3.data' /RECTYPE 1 (a).
```

¹This document has been prepared by Patty Glynn, University of Washington.

```
do if (RECTYPE eq 'h').
 reread.
    DATA LIST file = 'c:\all\spssclass\progs\prog3.data'
      / serial 2-3 region 4 statefip 5-6 county 7-10 stateco 5-10 .
end case.
end if.
** keep and save only household records and variables .
select if (RECTYPE eq 'h').
end input program.
variable label
rectype = 'Record Type'
 serial = 'Number unique to HH'
region = 'Geographic Region'
statefip= 'State Fips Code'
county = 'County ICPSR Code'
 stateco = 'Fips State + ICPSR County' .
save outfile = 'c:\all\spssclass\progs\prog4hh.sav' .
** READ and save PERSON cases and VARIABLES .
input program .
DATA LIST file = 'c:\all\spssclass\progs\prog3.data' /RECTYPE 1 (a).
do if (RECTYPE eq 'p').
 reread .
    DATA LIST file = 'c:\all\spssclass\progs\prog3.data'
        / serial 2-3 pernum 4-5 relate 6 age 7-8.
end case.
end if.
end input program.
select if (RECTYPE eq 'p').
variable label
rectype = 'Record Type'
pernum = 'Person number in HH'
relate = 'Relationship to head'
       = 'Age in years' .
age
save outfile = 'c:\all\spssclass\progs\prog4pers.sav' .
** program ends here .
```

Note that two "save outfile" commands are issued, saving two different data sets. The first saves the household cases and variables, and the second saves the person records and variables.

Variations on Merging - Wrong and Right.

It is common in social science to have to put to data sets together so that each case has more variables. For example, putting household and person variables for PUMS files, if they are stored separately would require merging.

In SPSS, the command to accomplish this is "MATCH FILES". The syntax varies somewhat depending on the situation, as I will show later.

If both files have one case for each "variable in common" then the syntax for matching files is :

```
match files file = 'c:\directory\fileone.sav' /
file = 'c:\directory\fileone.sav' /
by {variable(s) in common}.
```

While a BY statement is not always required, it is best to include one if possible. The likelihood of putting data together incorrectly is greater if you don't use a "by" statement.

Examples of successful and unsuccessful merges, and explanations follow.

This a simple merge where there are the same observations in both data sets, and the observations are in the same order.

*** PROGRAM BEGINS HERE *** ; * testmrq1.SPS. title 'testmrg1.SPS - no by statement' . ** Read and save first data set. DATA LIST FREE / Name (a9) g1 (A2) g2 (A2) g3 (A2) . BEGIN DATA Alice al a2 a3 Brian b1 b2 b3 Charles c1 c2 c3 Dianne d1 d2 d3 END DATA. save outfile = 'c:\all\spssclass\trash\f1.sav' . ** Read and save second data set. DATA LIST FREE / Name (a9) q4(A2) q5(A2) q6(A2). BEGIN DATA Alice a4 a5 a6 Brian b4 b5 b6 Charles c4 c5 c6 Dianne d4 d5 d6 END DATA. save outfile = 'c:\all\spssclass\trash\f2.sav' . match files file = 'c:\all\spssclass\trash\f1.sav' / file = 'c:\all\spssclass\trash\f2.sav' . list var = all . NAME G1 G2 G3 G4 G5 G6 al a2 a3 a4 a5 a6 b1 b2 b3 b4 b5 b6 Alice Brian Charles c1 c2 c3 c4 c5 c6 Dianne d1 d2 d3 d4 d5 d6 Number of cases read: 4 Number of cases listed: 4 Everything went well in this case because the cases matched in both data sets. But, suppose the data were not in the same order in both data sets. *** PROGRAM BEGINS HERE *** ; * testmrg2.SPS. title 'testmrg2.SPS, no by statement, wrong - cases not match' . ** Read and save first data set. DATA LIST FREE / Name (a9) g1 (A2) g2 (A2) g3 (A2) . BEGIN DATA Alice a1 a2 a3 Brian b1 b2 b3 Charles c1 c2 c3 Dianne d1 d2 d3 END DATA. save outfile = 'c:\all\spssclass\trash\f1.sav' . ** Read and save second data set. DATA LIST FREE / Name (a9) g4(A2) g5 (A2) g6 (A2) . BEGIN DATA Brian b4 b5 b6

Charles c4 c5 c6 Dianne d4 d5 d6 Alice a4 a5 a6 END DATA. save outfile = 'c:\all\spssclass\trash\f2.sav' . match files file = 'c:\all\spssclass\trash\f1.sav' file = 'c:\all\spssclass\trash\f2.sav' list var = all . G1 G2 G3 G4 G5 G6 NAME al a2 a3 b4 b5 b6 Alice b1 b2 b3 c4 c5 c6 Brian Charles c1 c2 c3 d4 d5 d6 Dianne d1 d2 d3 a4 a5 a6 Number of cases read: Number of cases listed: 4 4

Notice that the data for Brian and Alice are mixed up. The match did not go well, and our data are very wrong.

Now consider the following example. The data are not in the right order, and, data for Brian is missing from the second data set. Examine the names. Each case can have only one value for each variable, and the values for the variable "name" were different in the two data sets. One data set will overwrite the values of another. Note that the values from data set two overwrote the values for name in data set one.

*** PROGRAM BEGINS HERE *** ; * testmrg3.SPS. title 'testmrg3.SPS, no by statement, wrong - cases not match' . ** Read and save first data set. DATA LIST FREE / Name (a9) g1 (A2) g2 (A2) g3 (A2) . BEGIN DATA Alice al a2 a3 b1 b2 b3 Brian Charles c1 c2 c3 Dianne d1 d2 d3 END DATA. save outfile = 'c:\all\spssclass\trash\f1.sav' . ** Read and save second data set. DATA LIST FREE / Name (a9) q4(A2) q5(A2) q6(A2). BEGIN DATA Charles c4 c5 c6 Dianne d4 d5 d6 а4 а5 аб Alice END DATA. save outfile = 'c:\all\spssclass\trash\f2.sav' . match files file = 'c:\all\spssclass\trash\f1.sav' / file = 'c:\all\spssclass\trash\f2.sav' . list var = all . NAME G1 G2 G3 G4 G5 G6 a1 a2 a3 c4 c5 c6 Alice b1 b2 b3 d4 d5 d6 Brian Charles c1 c2 c3 a4 a5 a6 d1 d2 d3 Dianne

Number of cases read: 4 Number of cases listed: 4 Now we will add a by statement, and examine the results. And, this time we must look at the log to understand what happens. (Of course, we should always look at the log!)

*** PROGRAM BEGINS HERE *** ; * testmrq4.SPS. title 'testmrg4.SPS, wrong - cases not sorted' . ** Read and save first data set. DATA LIST FREE / Name (a9) g1 (A2) g2 (A2) g3 (A2) . BEGIN DATA Alice al a2 a3 Brian b1 b2 b3 Charles c1 c2 c3 Dianne d1 d2 d3 END DATA. save outfile = 'c:\all\spssclass\trash\f1.sav' . ** Read and save second data set. DATA LIST FREE / Name (a9) g4(A2) g5 (A2) g6 (A2) . BEGIN DATA Charles c4 c5 c6 Dianne d4 d5 d6 Alice a4 a5 a6 END DATA. save outfile = 'c:\all\spssclass\trash\f2.sav' .

```
match files
file = 'c:\all\spssclass\trash\f1.sav' /
file = 'c:\all\spssclass\trash\f2.sav' / by name / map .
```

```
list var = all .
```

RESULTS:

NAME G1 G2 G3 G4 G5 G6 Alice al a2 a3 Brian b1 b2 b3

Charles c1 c2 c3 c4 c5 c6 File #2 KEY: Alice

>Error # 5130 >File out of order. All the files in MATCH FILES must be in non-descending >order on the BY variables. Use SORT CASES to sort the file. >This command not executed.

>Note # 35 >There is no working file to be restored. The state of the SPSS Processor >is as if a NEW FILE command had been executed.

The matched data are not correct, and we got an error message. Data to be matched must be sorted in the order of the variable(s) in the by statement. If the files are already in that order, then it is not necessary to sort them. Sorting is a resource intensive task.

Efficiency note: Use keep statements and SELECT IF statements to eliminate variables and cases not needed before sorting.

*** PROGRAM BEGINS HERE *** ;
* testmrg5.SPS.
title 'testmrg5.SPS, cases sorted, by statement - ok' .
** Read and save first data set.
DATA LIST FREE / Name (a9) g1 (A2) g2 (A2) g3 (A2) .

```
BEGIN DATA
Alice al a2 a3
Brian bl b2 b3
Charles c1 c2 c3
Dianne d1 d2 d3
END DATA.
save outfile = 'c:\all\spssclass\trash\f1.sav' .
** Read and save second data set.
DATA LIST FREE / Name (a9) g4(A2) g5 (A2) g6 (A2) .
BEGIN DATA
Charles c4 c5 c6
Dianne d4 d5 d6
Alice
       a4 a5 a6
END DATA.
** The cases in this file are not sorted in the correct order .
** The following command asks SPSS to sort them .
sort cases by name .
save outfile = 'c:\all\spssclass\trash\f2.sav' .
match files
   file = 'c:\all\spssclass\trash\f1.sav' /
   file = 'c:\all\spssclass\trash\f2.sav' / by name / map .
list var = all \cdot
match files
   file = 'c:\all\spssclass\trash\f1.sav' /
   file = 'c:\all\spssclass\trash\f2.sav' / by name / map .
```

Note the "/ map" in the above command line. This asks SPSS to tell you which file each variable came from in the resulting matched file. This is a very useful tool. The information this request provides is given below. Sometimes, variables might exist in two data set. Of course, each variable can only hold one value for each case. So, the value from one of the files will be lost in the resulting matched data set. The "MAP" option allows you to figure out which file "won", and to determine if any variables were over-witten.

Map of BY	variables				
Result	Input	l In	Input2		
NAME	NAME	- <u>-</u> NA	NAME		
Map of th Result 	e result file Input1 Input2				
NAME G1 G2 G3	NAME G1 G2 G3	NA	ME		
G4 G5 G6		G4 G5 G6			
list var NAME	= all G1 G2	G3 G4	G5	G6	
Alice Brian	a1 a2 b1 b2	a3 a4 b3	a5	аб	
Charles Dianne Number of	c1 c2 d1 d2 cases	c3 c4 d3 d4 read:	c5 d5 4	с6 d6	
Number of	cases	listed:		4	

. .

Homework for next week:

Write a SPSS program to read the following hierarchical data set. Read the data as an external file. Create TWO systems files - one with household data and one with person data, then merge them. Save the final merged data set as a permanent SPSS data set.

The variables for the Household data set are:

Record type household ID number of persons in the household region state county An indicator of whether it is in an MSA or not.

The variables for the Person data set are:

Record type household ID person number relationship to head age race gender highest grade completed