Subject: Re: Merging data files

Posted by Janet-DHS on Thu, 16 Nov 2023 19:31:40 GMT

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Following is a response from DHS staff member, Tom Pullum:

As with the earlier merge, I will give the code in Stata. I hope you can follow the logic and convert to SPSS. Note that the AR file is used twice, once to merge with the men in the CR file and once to merge with the women. The CR file is sorted differently for each merge

- \* Stata lines to merge the CR and AR files for the Cameroon 2018 DHS survey
- \* Note that the cases in the CR file are couples
- \* The cases in the AR file are individuals, with sex not specified
- \* In the CR file, v001=mv001 and v002=mv002
- \* Specify a workspace cd e:\DHS\DHS\_data\scratch
- \* Prepare the AR file use "...CMAR71FL.DTA", clear

gen cluster=hivclust gen hh=hivnumb

- \* w for woman, m for man gen wline=hivline gen mline=hivline sort cluster hh wline save CMARtemp.dta, replace
- \* Prepare the CR file
  use "...CMCR71FL.DTA"
  gen cluster=v001
  gen hh=v002
  gen wline= v003
  gen mline=mv003
  gen in\_CR=1
  sort cluster hh wline
  save CMCRtemp.dta, replace
- \* Merge the women merge 1:1 cluster hh wline using CMARtemp.dta keep if in\_CR==1 rename hiv\* w hiv\*
- \* women who are in both files have \_wmerge=3 rename \_merge \_wmerge

\* Must re-sort both files save CMCRtemp.dta, replace

use CMARtemp.dta, clear sort cluster hh mline save CMARtemp.dta, replace

- \* Go back to the CR file that has been merged for women use CMCRtemp.dta sort cluster hh mline
- \* Merge the men
- \* This is not a 1:1 merge because husbands can appear more than once merge m:1 cluster hh mline using CMARtemp.dta keep if in\_CR==1 rename hiv\* m\_hiv\*

rename \_merge \_mmerge tab \_wmerge \_mmerge,m tab \_wmerge \_mmerge if in\_CR==1,m tab w\_hiv03 m\_hiv03,m drop in\_CR \*merge

label list HIV03
replace w\_hiv03=10 if w\_hiv03==.
replace m\_hiv03=10 if m\_hiv03==.
label define HIV03 10 "Not tested", modify tab w\_hiv03 m\_hiv03,m

\* Save the merged cases with a new file name