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Subject: Merging 2015 Angola HIV data with standard DHS

Posted by [Koray](#) on Fri, 03 May 2024 12:44:18 GMT

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Hello... Please I need help. Below are my steps in merging HIV data with the 2015 Angola standard DHS.

Prepare the AR file

use "...A0AR71FL.DTA", clear

gen cluster=hivclust

gen hh=hivnumb

gen line=hivline

sort cluster hh line

\* Prepare the IR file

use "...AOIR71FL.DTA", clear

gen cluster=v001

gen hh=v002

gen line=v003

sort cluster hh line

\* Do the merge

merge 1:1 cluster hh line using

Outcome

merge 1:1 cluster hh line using "

C:\Users\Gbang\Downloads\AO\_2015-16\_DHS\_05032024\_1157\_156908 \AO

> AR71DTVAOAR71FL.DTA"

Result	# of obs.
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not matched	13,403
from master	7,745 (_merge==1)
from using	5,658 (_merge==2)
matched	6,634 (_merge==3)

Almost 50% of the data were not matching. Where did I go wrong?

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Subject: Re: Merging 2015 Angola HIV data with standard DHS

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Posted by [Janet-DHS](#) on Fri, 10 May 2024 19:30:17 GMT

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

Nothing is wrong with what you did. The AR file is approximately evenly divided between women and men. You matched the women.

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