
Subject: Re: HH head education level merge
Posted by [kate2](#) on Mon, 03 Nov 2014 08:09:55 GMT
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Thanks for the link and the suggestions. Using m:1 gives me the error about not uniquely IDing observations in the 'using' data; and using 1:m gives me the error about not uniquely IDing observations in the master data....

So clearly I should be able to uniquely ID observation in the dataset that I want to merge onto my master data set other wise its not useful. I thought that I had then worked out the problem - that I need to use cluster as well as household id to uniquely identify observations in my HH head education dataset - because HH IDs are not sequential through the whole dataset, rather they (I think) restart for each new cluster. Bingo...thought I had it. So I ran these commands below...but then got the same error result...

```
use "HH member recode"  
keep if hv101 = 1  
gen HH_ed = hv106  
keep v001 v002 HH_ed  
sort v001 v002  
save "HH_member_recode_HHedonly"  
clear  
use "child merged dataset"  
sort v001 v002  
merge m:1 v001 v002 using "HH_member_recode_HHedonly"
```

```
>>> variables v001 v002 do not uniquely identify observations in the using data
```

So I am clearly missing something as to how I uniquely identify households. Unless some households have more than one HH head recorded -- will look into that possibility.

Any other ideas? Thank for your help!
Kate