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Subject: Re: Height and Weight Files

Posted by [rejone4@emory.edu](mailto:rejone4@emory.edu) on Wed, 19 Nov 2014 19:36:31 GMT

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thanks that worked well for most datasets. However there are a few HW files that have hwhhid as the variable for case identification.

For example, Armenia 2005 has this type of setup. When I used the following code:

```
*rename variables and sort data for merging
```

```
ren hwhhid caseid
```

```
ren hwline midx
```

```
sort caseid midx
```

```
save "AMHW01_temp.dta", replace
```

```
*sort the children's recode data and then merge
```

```
use "H:\Thesis\Armenia_2010\Armenia_2005(V)\AMKR42FL.dta", clear
```

```
sort caseid midx
```

```
merge 1:1 caseid midx using "H:\Thesis\Armenia_2010\Armenia_2005(V)\AMHW01_temp.dta"
```

It did not match the data anymore. Is there something that has to be done differently with these files?