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?