Subject: Merge Other Biomarkers with HH Posted by mer99 on Thu, 07 Jul 2022 10:15:21 GMT View Forum Message <> Reply to Message

Good evening,

I am trying to merge all Other Biomarkers dataset (from men, children and women) from Malawi 2015-2016 into the Household Dataset. I have used append to unify all four Other Biomarkers datasets, and then merged them into the HH dataset by using the following code:

gen mcluster = hv001 gen mnumber = hv002 gen m01 = hv003

sort mcluster mnumber m01

* merge mother's characteristics into PR file merge m:1 mcluster mnumber m01 using "`secondary'", keep(master match)

However, all I get are duplicates and I am not sure at all of what I am doing wrong. If anyone could help me with this I would be quite grateful. Thank you!