Subject: Re: Child growth
Posted by mnisha on Tue, 26 Jul 2016 07:44:08 GMT
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Thank you so much. I was a bit confused about the merging. I tried 1:1_n merging before. It worked for me as well. But actually I have another problem. My base file is IR dataset. If I use "keep if _merge==3", the number of early neonatal deaths gets decreased from the actual number. So I would like to ask you if my base file is IR dataset and if I want to merge it with BR dataset in order to get the total number of perinatal deaths, is it necessary to drop the unmatched cases in the merged file?

Thanks in advance.