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Subject: merging IR file with KR

Posted by [mmbah](#) on Thu, 07 Mar 2019 00:09:59 GMT

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Dear Sir/ madam,

I wanted to calculate the total number of csection using the IR file for Malawi using the following commands:

use MWIR7HFL.dta

. gen csection=0

. replace csection=1 if m17\_1==1 | m17\_2==1 | m17\_3==1 | m17\_4==1 | m17\_5==1 | m17\_6==1  
(976 real changes made)

tab csection

csection	Freq.	Percent	Cum.
0	23,586	96.03	96.03
1	976	3.97	100.00
Total	24,562	100.00	

However this is less than the total csection in the KR files

tab m17

delivery by   caesarean   section	Freq.	Percent	Cum.
no	16,122	93.53	93.53
yes	1,116	6.47	100.00
Total	17,238	100.00	

I tried to merge the IR and and the KR file so that I can use the m17 variable but loose lots of data in the merging process. I wanted someone to kindly help me calculated the total csection using the IR file or how to merge the m17 to the IR file

secondly.....

i calculate the number of stillbirths using the following commands:

```
use MWIR7HFL  
gen stillbirths = regexm(vcal_1,"TPPPPPP")  
label var stillbirths "stillbirths"
```

i obtained 213 stillbirths and the DHS final report indicated 236. Could someone help explained why and how to get the right values.

thanks for your kind help and understanding in advance  
regards  
Mamadou

Subject: Re: merging IR file with KR  
Posted by [Mu](#) on Thu, 07 Mar 2019 07:36:03 GMT  
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Hello Mamadou,

This is how I calculated the c-section births. I did not use the IR file, but instead used the BR file  
FOR C-SECTION

```
clear all  
set matsize 800  
set maxvar 10000  
set mem 1g  
cd "..."  
use "MWBR7HFL", clear  
set more off
```

\*\* WEIGHT VARIABLE  
gen weight = v005/1000000

\*\* SURVEY SET





```
ssc install hplot, replace //Need internet connection
ssc install catplot, replace //Need internet connection

catplot c_section wealth [iw=weight], percent(wealth) stack asyvars ///
bar(1, bcolor(gold*.6)) bar(2, bcolor(ltblue*.9)) bar(3, bcolor(red*.2)) ///
blabel(bar, format(%9.1f) pos(center)) bar(4, bcolor(gs14)) ytitle(%) ///
title(Percentage of births delivered by C-section) subtitle(Malawi DHS 2015-16)

catplot delivery_place wealth [iw=weight], percent(wealth) stack asyvars ///
bar(1, bcolor(gold*.6)) bar(2, bcolor(ltblue*.9)) bar(3, bcolor(red*.2)) ///
blabel(bar, format(%9.1f) pos(center) size(small) orientation(vert)) ///
bar(4, bcolor(gs14)) ytitle(%) ///
title(Percent distribution of births by place of delivery) ///
subtitle(Malawi DHS 2015-16)

catplot facility_delivery wealth [iw=weight], percent(wealth) stack asyvars ///
bar(1, bcolor(gold*.6)) bar(2, bcolor(ltblue*.9)) bar(3, bcolor(red*.2)) ///
blabel(bar, format(%9.1f) pos(center)) bar(4, bcolor(gs14)) ytitle(%) ///
title(Percentage of births by type of health facility) ///
subtitle(Malawi DHS 2015-16)

catplot skilled_birth wealth [iw=weight], percent(wealth) stack asyvars ///
bar(1, bcolor(gold*.6)) bar(2, bcolor(ltblue*.9)) bar(3, bcolor(red*.2)) ///
blabel(bar, format(%9.1f) pos(center)) bar(4, bcolor(gs14)) ytitle(%) ///
title(Percentage of births by type of provider) ///
subtitle(Malawi DHS 2015-16)
```

exit

Subject: Re: merging IR file with KR  
Posted by [Mlue](#) on Thu, 07 Mar 2019 07:54:00 GMT  
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This one might give you the total for stillbirths = 236

```
clear all  
set matsize 800  
set maxvar 10000  
set mem 1g  
cd "..."
```



Subject: Re: merging IR file with KR  
Posted by [mmbah](#) on Thu, 07 Mar 2019 14:06:00 GMT

Dear MLUE,

Thank you very much for the wonder assistance which was very very helpful. I want to also calculated prenatal mortality and I used the following command. However, i am not sure I have calculated it right.

```
gen early_neonatal_deaths = 0
label var early_neonatal_deaths "earlyneo"
gen infant_deaths = 0

gen child_deaths = 0
label var child_deaths "child mortality"
label var infant_deaths "infant mortality"
* rename birth variables for use in forvalues loop
rename b*_0* b*_*

* Any early neonatal deaths, infant deaths, child deaths
forvalues b = 1/80 {
    * early neonatal - days 0-6
    replace early_neonatal_deaths = 1 if inrange(b6_`b',100,106)
}
gen wt = v005/1000000
replace wt = stillbirths*v005/1000000
replace wt = early_neonatal_deaths *v005/1000000
* perinatal mortality includes stillbirths and early neonatal deaths
gen perinatal_deaths = (stillbirths | early_neonatal_deaths)
label var perinatal_deaths "perinatal"
```

secondly I wants to run a logistic regression model to look at association between csection and perinatal mortality adjusting for demographics variables, morbidity, and ANC visits. could you help with graphs

Thank you very much for you kind assistance which is very well appreciated.

regards  
Mamadou

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Subject: Re: merging IR file with KR  
Posted by [mmbah](#) on Thu, 11 Apr 2019 01:41:29 GMT  
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Dear Mlue,

Thanks you very much for the codes as they are very helpful. I wanted to calculate the number of ealy neonatal mortality and perinatal mortality. However, when I reshape the data using the codes you provide i have over 150000 observations. If I calculate the early neonatal and perinatal

deaths, bmi, parity using the below commands before or after the reshaping i get wrong estimation due to the high number of observation.I really need help.

```
replace end = v008
replace beg = v008-59

* rename b3 and b6 variables to facilitate use in the for loop
rename b3_0* b3_*
rename b6_0* b6_*

* Loop through birth history summing births and early neonatal deaths
* in the five years preceding the survey
forvalues i = 1/20 {
    * restrict to 60 months preceding survey
    replace births2 = births2+1 if inrange(b3_`i',beg,end)
    replace earlyneo = earlyneo+1 if inrange(b3_`i',beg,end) & inrange(b6_`i',100,106)
}
gen perinatal = earlyneo+stillbirths
label variable perinatal "Perinatal mortality"

label define bmic 1 "Underweight" 2 "Normal" 3 "Overweight/obese"
gen bmi= v445/100
label var bmi "bmi"
gen bmic= 1 if bmi<18.5
replace bmic=2 if bmi>=18.5 & bmi<25
replace bmic=3 if bmi>=25 & bmi!=.
label values bmic bmic
label var bmic "bmic"
tab bmic
rename v201 parity
recode parity 0/1=1 2=2 3=3 4=4 5/max=5
tab parity
label define paritylab 5 "5+"
label var parity paritylab
label value parity paritylab
tab parity
```

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