Dear Hassen,

We just recently uploaded the code for the FGM indicators on GitHub. You can find it here:
https://github.com/DHSProgram/DHS-Indicators-Stata/tree/mast er/Chap18_FG
Also you can read more about these indicators here:
https://www.dhsprogram.com/Data/Guide-to-DHS-Statistics/inde
x.htm#t=18_Female_Genital_Cutting.htm
The FGM indicators us the IR file.

As mentioned previously, before attempting to run any code from our GitHub code share site,
please read the readme file and the main file for the chapter.

Perinatal mortality is a rate and you cannot create a dichotomous variable from this.

For your question on community level variables, there are many ways to do this. One easy way is
to create a binary variable of the indicator you want at the community level and then take the
mean at the cluster level. For example in the code below I create a binary variable for education
that is secondary or more. Then I create a cluster or community level variable educlus as shown.

recode v106 (0/1=0 "noedu/primary") (2/3=1 "secondary+") (8/9=.), gen(edu)
bysort v001: egen educlus=mean(edu)

Hope this helps.

Best,
Shireen Assaf
The DHS Program