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clear
capture log close
log using "c:\usr\rino\projects\ovarian_new\results\cr.embla.doc", text replace

version 8.2
set mem 50M
/*
*****
Date           : 24/01/2005
Update        : 01/02/2005
Authors      : Rino Bellocco

Name of file   : cr.embla.do

Input Data file : Files from questioannaire
Output Data file: embla, and files from questionnarire with _1

Content :
          1: Generate new pnr
          2: Read file and merge them consecutively
          3: Save variables of interest
          4: file
*****
*/
use "c:\usr\rino\projects\ovarian_new\data\embla\personalia.dta", clear
capture gen var1=substr(pnr,1,6)
capture gen var2=substr(pnr,8,4)
capture gen newpnr="19" + var1+ var2
sort lopnr
drop var1 var2
save "c:\usr\rino\projects\ovarian_new\data\embla\personalia_1.dta",replace

use "c:\usr\rino\projects\ovarian_new\data\embla\diagnosis.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\diagnosis_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\cause.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\cause_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\status.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\status_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\operations.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\operations_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\properties.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\properties_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\chemo.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\chemo_1.dta", replace

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use "c:\usr\rino\projects\ovarian_new\data\embla\radiation.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\radiation_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\hormons.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\hormons_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\progress.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\progress_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\newtumor.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\newtumor_1.dta", replace

use "c:\usr\rino\projects\ovarian_new\data\embla\deceased.dta", clear
capture rename sub_lopnr lopnr
sort lopnr
save "c:\usr\rino\projects\ovarian_new\data\embla\deceased_1.dta", replace

cd c:\usr\rino\projects\ovarian_new\data\embla\

merge lopnr using deceased_1 newtumor_1 progress_1 hormons_1 radiation_1 ///
                chemo_1 properties_1 operations_1 status_1 cause_1
diagnosis_1 personalia_1

drop pnr
rename newpnr pnr
move pnr person
keep lopnr pnr exclude ///
    diagdat fupdat ///
    na0301 gyninv symptom other ///
    notop opy1 opdat1 na0601_1 oprad1 opnrad1 na0603_1 rescm1 restxt1 resnm1
na0602_1 ///
    hyst1 subhyst1 soeun1 soebill1 oment1 lgllbuk1 ///
    opother1 opy2 opdat2 na0601_2 oprad2 opnrad2 na0603_2 rescm2 restxt2 ///
    resnm2 na0603_3 rescm3 restxt3 resnm3 sekopn sekopy1 sekopd1 sekopy2
sekopd2 sekopy3 sekopd3 ///
    na0702 maxdia maxtxt na0703 ///
    figoia figoib figoic figoix figoiaa figoibb figoicc figoixx figoiaa ///
    figoibb figoibc figoibx figoivx figofr figofo figoni ///
    na0704 malbor malmal tisepi tisnepi ///
    na0706 whohi whome whole ///
    na08011 chemoy1 chemon1 typlal1 typcar1 ///
    typtax1 admcnt1 cycles1 startd1 lastd1 na08012 chemoy2 chemon2 ///
    startd2 lastd2 na08013 chemoy3 chemon3 na08014 chemoy4 chemon4 na08015
chemoy5 chemon5 na08016 chemoy6 chemon6 chemonow ///
    na0901 rady radn startd na1001 enn eny enstd enendd ennow na1002 hrtm hrty
hrtstd hrtnow hrtendd ///
    prfn prfy prfdat metn mety metdat npcy npcn ///
    npcnd npgova npgend npgcrx decn decy decdat causeca causeot causetxt

sort pnr

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save "c:\usr\rino\projects\ovarian_new\data\embla\embla.dta",replace
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log close
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