
PURPOSE:
PROGRAMMER:
DATE:
OUTPUT:
MACROS:

MODIFICATION HISTORY:

*****;

libname sasraw '/home/u64395997/SASSTUDY/Raw data';

```
proc copy in=sasraw out=work;  
  select demog demog1 ex;  
run;
```

```
options validvarname=upcase;
```

```
/* SUBJID */
```

```
data dm1;  
  set demog;  
  subjid1=strip(put(subjid,best.));  
  studyid='S-CDSK-12445';  
  domain='DM';  
  drop subjid;  
run;
```

```
/* RFXSTDTC */
```

```
data dm2;  
  set ex;
```

```
dt1=input(exstdat_raw,date11.);
rfstdtc=strip(put(dt1,is8601da.));
dt2=input(exendat_raw,date11.);
rfendtc=strip(put(dt2,is8601da.));
rfxstdtc=rfstdtc;
rfxendtc=rfendtc;
run;
```

```
proc sort data=dm1;
  by usubjid;
run;
```

```
proc sort data=dm2;
  by usubjid;
run;
```

```
data dm3;
  merge dm1(in=a) dm2(in=b);
  by usubjid;
run;
```

```
/* RFICDTC */
```

```
data dm4;
  set demog1;
  dt=input(icdat_raw,date11.);
  rficdtc=strip(put(dt,is8601da.));
  drop subjid dt enddat_raw;
run;
```

```
data dm5;
  merge dm3(in=a) dm4(in=b);
  by usubjid;
  if a;
run;
```

```
/* RFPENDTC */
```

```
data dm6;
```

```
set demog1(rename=(icdat_raw=date)) demog1(rename=(enddat_raw=date)) ex(rename=(exstdat_raw=date))  
  ex(rename=(exendat_raw=date));  
dtx=input(date,date11.);  
keep usubjid date dtx;  
run;
```

```
proc sort data=dm6;  
  by usubjid dtx;  
run;
```

```
data dm7;  
  set dm6;  
  by usubjid dtx;  
  if last.usubjid;  
run;
```

```
data dm8;  
  merge dm5(in=a) dm7(in=b);  
  by usubjid;  
  if a;  
  dthdtc='';  
  dthfl='';  
  brthdtc0=strip(put(brthdtc,is8601da.));  
  actarm=arm;  
  actarmcd=armcd;  
  drop brthdtc subjid;  
run;
```

```
data dm (label='Demographics');  
  retain STUDYID DOMAIN USUBJID SUBJID1 RFSTDTC RFENDTC RFXSTDTC RFXENDTC RFICDTC RFPENDTC  
  DTHDTC DTHFL SITEID BRTHDTC0 AGE AGEU SEX RACE ETHNIC ARMCD ARM ACTARMCD ACTARM COUNTRY;  
  set dm8;  
  label STUDYID='Study Identifier' DOMAIN='Domain Abbreviation'  
  USUBJID='Unique Subject Identifier' SUBJID='Subject Identifier for the Study'  
  RFSTDTC='Subject Reference Start Date/Time' RFENDTC='Subject Reference End Date/Time'  
  RFXSTDTC='Date/Time of First Study Treatment' RFXENDTC='Date/Time of Last Study Treatment'  
  RFICDTC='Date/Time of Informed Consent' RFPENDTC='Date/Time of End of Participation'  
  DTHDTC='Date/Time of Death' DTHFL='Subject Death Flag'  
  SITEID='Study Site Identifier' BRTHDTC='Study Site Identifier'
```

```
AGE='Age' AGEU='Age Units' SEX='Sex' RACE='Race' ETHNIC='Ethnic'
ARMCD='Planned Arm Code' ARM='Description of Planned Arm' ACTARMCD='Actual Arm Code'
ACTARM='Description of Actual Arm' COUNTRY='Country';
rename SUBJID1=SUBJID BRTHDTC0=BRTHDTC;
keep STUDYID DOMAIN USUBJID SUBJID1 RFSTDTC RFENDTC RFXSTDTC RFXENDTC RFICDTC RFPENDTC
DTHDTC DTHFL SITEID BRTHDTC0 AGE AGEU SEX RACE ETHNIC ARMCD ARM ACTARMCD ACTARM COUNTRY;
run;
```

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*****;

```
libname sasraw '/home/u64395997/SASSTUDY/Raw data';
```

```
proc copy in=sasraw out=work;
  select ae2 ae1 aecode aedt;
run;
```

```
options validvarname=upcase;
```

```
data A1;
  set AE1;
  studyid='S-CDSK-01';
  domain='AE';
  AESEQ=_n_;
  AESEPID=recordno;
```

```
AETERM=UPCASE(aeterm1);  
if aeterm1='AGITATED' then AEMODIFY='AGITATION';  
if aeterm1='VOMIT' then AEMODIFY='VOMITING';  
by usubjid;  
drop recordno;  
run;
```

```
proc sort data=A1;  
  by aeterm1;  
run;
```

```
proc sort data=aecode;  
  by aeterm1;  
run;
```

```
data A2;  
  merge A1(in=a) aecode(in=b);  
  by aeterm1;  
  if a;  
  drop aeterm1;  
run;
```

```
data A3;  
  set AE2;  
  by usubjid;  
  length aesev $20;  
  if ae_sev=1 then aesev='Mild';  
  if ae_sev=2 then aesev='Moderate';  
  if ae_sev=3 then aesev='Severe';  
  if ae_sev le 2 then aeser='N';  
  else aeser='Y';  
  AEACN=AE_ACN;  
  AEREL=AE_REL;  
  AET=input(ae_endat,date11.);  
  AEENDTC=strip(put(aet,is8601da.));  
  drop aet ae_acn ae_rel ae_sev ae_endat;  
run;
```

```
proc sort data=A2;  
  by usubjid;  
run;
```

```
proc sort data=A3;  
  by usubjid;  
run;
```

```
data A4;  
  merge A2(in=a) A3(in=b);  
  by usubjid;  
  if a;  
run;
```

```
/* AESTDTC */
```

```
data A5;  
  set aedt;  
  aest=input(aestdat_raw,date11.);  
  AESTDTC=strip(put(aest,is8601da.));  
  drop aest aestdat_raw;  
run;
```

```
data A6;  
  merge A4 A5;  
  by usubjid;  
  if AEENDTC='.' then do;  
    AEENRF='AFTER';  
  END;  
run;
```

```
data A7;  
  merge A6(in=a) DM(in=b);  
  by usubjid;  
  if a;  
  aestdtc1=input(aestdtc,is8601da.);  
  aeendtc1=input(aeendtc,is8601da.);  
  rfstdtc1=input(rfstdtc,is8601da.);  
  if aestdtc1 ge rfstdtc1 then do;
```

```
aestdy1=(aestdtc1-rfstdtc1)+1;  
if aestdtc1 lt rfstdtc1 then do;  
aestdy1=(aestdtc1-rfstdtc1);  
end;  
end;  
if aeendtc1 ge rfstdtc1 then aeendy1=(aeendtc1-rfstdtc1)+1;  
else aeendy1=(aeendtc1-rfstdtc1);  
aestdy=day(aestdy1);  
aeendy=day(aeendy1);  
keep usubjid aestdy aeendy;  
run;
```

```
data A8;  
merge A6(in=a) A7(in=b);  
by usubjid;  
if a;  
run;
```

```
data AE (label='Adverse Events');  
retain STUDYID DOMAIN USUBJID AESEQ AESPID AETERM AEMODIFY AEDECOD AEBODSYS AESEV AESER  
AEACN AEREL AESTDTC AEENDTC AESTDY AEENDY AEENRF;  
set A8;  
label STUDYID='Study Identifier' DOMAIN='Domain Abbreviation'  
USUBJID='Unique Subject Identifier' AESEQ='Sequence Number'  
AESPID='Sponcer Dsfind Identifier' AETERM='Reported Term for the Adverse Evevnts'  
AEMODIFY='Modified Reported Term' AEDECOD='Dictionary Derived Term'  
AEBODSYS='Body System Organ Class' AESEV='Severity/Intensity'  
AESER='Serious Event' AEACN='Action Taken with Study Treatment'  
AEREL='Causality' AESTDTC='Start Date/Time of Adverse Event'  
AEENDTC='End Date/Time of Adverse Event' AESTDY='Study Day of Start of Adverse Event'  
AEENDY='Study Day of End of Adverse Event' AEENRF='End Relative to Reference Period';  
keep STUDYID DOMAIN USUBJID AESEQ AESPID AETERM AEMODIFY AEDECOD AEBODSYS AESEV AESER  
AEACN AEREL AESTDTC AEENDTC AESTDY AEENDY AEENRF;  
run;
```

