

```
*****
```

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;
  subj1=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 RFSTDTCT RFENDTC RFXSTDTC RFXENDTC RFICDTCT RFPENDTC
DTHDTCT 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'
RFSTDTCT='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'
RFICDTCT='Date/Time of Informed Consent' RFPENDTC='Date/Time of End of Participation'
DTHDTCT='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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PURPOSE:

PROGRAMMER:

DATE:

OUTPUT:

MACROS:

\*\*\*\*\*

MODIFICATION HISTORY:

\*\*\*\*\*;

```
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 aelev $20;
if ae_sev=1 then aelev='Mild';
if ae_sev=2 then aelev='Moderate';
if ae_sev=3 then aelev='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 AESTDTA AEENDTC AESTDY AEENDY AEENRF;
set A8;
label STUDYID='Study Identifier' DOMAIN='Domain Abbreviation'
USUBJID='Unique Subject Identifier' AESEQ='Sequence Number'
AESPID='Spencer Dsdefined 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' AESTDTA='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 AESTDTA AEENDTC AESTDY AEENDY AEENRF;
run;
```

