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*****
*****;
** Program Name : adxb.sas          **;
** Date Created : 17Nov2021          **;
** Programmer Name: (b) (4), (b)    **;
** Purpose      : Create adxb dataset **;
** Input data   : mb adsl adc19ef    **;
** Output data  : adxb.sas7bdat     **;
*****
*****;
%let
oprot=/Volumes/app/cdars/prod/sites/cdars4/prjC459/nda2_unblinded_esub/sbla1215_esub_sdtm/saseng/cdisc3_0/data/
sdtm;
%let
protori=/Volumes/app/cdars/prod/sites/cdars4/prjC459/nda2_unblinded_esub/sbla1215_esub_adam/saseng/cdisc3_0;
%let
prot=/Volumes/app/cdars/prod/sites/cdars4/prjC459/nda2_unblinded_esub/sbla1215_esub_adam/saseng/cdisc3_0/analy
sis/eSUB;
%let
expath=/Volumes/app/cdars/prod/sites/cdars4/prjC459/nda2_unblinded_esub/sbla1215_esub_adam/saseng/cdisc3_0/out
put/xpt;

libname dataprot "&oprot." access=readonly;
libname datvprot "&protori./data_vai" access=readonly;
libname datvout "&prot./data_vai";

*Insert the date of snapshot;
%let cutoff2=02SEP2021;

proc printto print="&prot./output/adxb.rpt"
    log="&prot./logs/adxb.log" new;
run;

* Clean *;
proc delete data=work._all_;
run;

*****Bring in SDTM MB dataset****/
proc sort data=dataprot.mb(where=(mbtestcd="WGSCLINE")) out=xb;
    by usbjid mbdtc visitnum visit mbtestcd MBSTREC;
run;

data xb;
    set xb;
Length PARAMCD $8. PARAM PARCAT1 $200. PARAMN AVAL AVISITN 8. AVALC $20.
    AVISIT $80. ADT 8.;
Format ADT date9.;

PARAM=strip(MBTEST);
PARAMCD=strip(MBTESTCD);
PARCAT1=strip(MBCAT);

IF upcase(MBTESTCD)="WGSCLINE" then
    PARAMN=1;

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AVAL=.;
AVALC=strip(MBSTRESC);
AVISITN=VISITNUM;
AVISIT=strip(VISIT);
ADT=input(mbdtc, ?? yymmdd10.);
Keep Usbjid PARAMCD PARAM PARCAT1 PARAMN AVAL AVALC AVISITN AVISIT ADT VISIT
VISITNUM;
run;

*****Bring in required Variable from ADSL dataset*****
proc sort data=datvprot.adsl(keep=studyid usbjid siteid subjid agegr1n agegr1
    sex race ethnic country armcd arm actarmcd actarm randdt trtsdt trtedt trt01a
    trt01an trt01p trt01pn vax101dt vax102dt dvstdt unblnddt randfl evaleffl
    aai1effl aai2effl /*mulenrfl*/ phasen hivfl) out=adsl;
    by usbjid;
run;

*****Bring in ADC19EF dataset to create required Flags***/
proc sort data=datvprot.adc19ef out=adc19ef;
    by usbjid;
run;

data eff1(keep=usbjid PDP27FL) eff2(keep=usbjid FC19D27 AVISIT);
    set adc19ef;
    by usbjid;

if first.usbjid then
    output eff1;

if paramcd in ("C19ONST") and FILOCRFL="Y" and AVALC="POS" and ILD27FL="Y"
    and ((not missing(DVSTD) and adt <=DVSTD) or missing(DVSTD)) and
    PDRMUPFL="N" and EVALEFFL='Y' then
        do;
        FC19D27="Y";
        output eff2;
    end;
run;

***get swab dates and visit from adc19ef dataset for first Occurence****
proc sort data=adc19ef(where=(paramcd in ("RTCOV2NS" "SARSCOV2") and avalc="POS")) out=posswab;
    by usbjid avisit;
run;

data eff21(keep=usbjid FC19D27 AVISIT visit adt);
    merge eff2(in=a) posswab;
    by usbjid avisit;
    if a;
run;

proc sort nodupkey;
    by usbjid visit adt;
run;

data feff(drop=avisit);

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set eff21;
by usubjid visit adt;
run;

/**Derive State from Pfizer Corporate Registry Database (based on Investigator ID)****/
data country;
length studyid $20. siteid $12. state $100.;
infile datalines delimiter=':';
input studyid $ siteid $ state $;
datalines;
C4591001: 1005: NEW YORK
C4591001: 1006: UTAH
C4591001: 1007: OHIO
C4591001: 1008: MISSOURI
C4591001: 1009: UTAH
C4591001: 1013: FLORIDA
C4591001: 1016: KENTUCKY
C4591001: 1039: TEXAS
C4591001: 1044: VIRGINIA
C4591001: 1057: FLORIDA
C4591001: 1084: TEXAS
C4591001: 1091: OHIO
C4591001: 1123: NEBRASKA
C4591001: 1124: RHODE ISLAND
C4591001: 1139: NORTH CAROLINA
C4591001: 1142: TEXAS
C4591001: 1150: OHIO
C4591001: 1152: CALIFORNIA
C4591001: 1156: FLORIDA
C4591001: 1235: LOUISIANA
C4591001: 1270: CALIFORNIA
;
run;

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proc sort data=adsl;
by studyid siteid usubjid;
run;

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proc sort data=country out=state(keep=studyid siteid state);
by studyid siteid;
run;

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data adsl;
merge adsl(in=a) state;
by studyid siteid;
if a;
run;

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```

proc sort data=adsl;
by usubjid;
run;

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*****merge all datasets to create ADXB dataset ****/
data adxb_;

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merge adsl(in=a) xb(in=b) eff1;
by usubjid;

if a and b;
length ADY 8.;

if trtsdt ^=. and adt ^=. then do;
    ady=adt - trtsdt;
    if adt >=trtsdt then ady=ady + 1;
end;
run;

proc sort;
    by usubjid visit adt;
run;

data adxb_1(drop=visit visitnum);
    merge adxb_(in=a) feff;
    by usubjid visit adt;

if a;

if FC19D27="" then FC19D27="N";
run;

*****Expand all the USA state Abbreviations as per Specifications****/
data adxb_2;
    set adxb_1;

if upcase(country)="USA" then do;

    if State="AK"      then state="ALASKA";
    else if State="AL"   then state="ALABAMA";
    else if State="AR"   then state="ARKANSAS";
    else if State="AZ"   then state="ARIZONA";
    else if State="CA"   then state="CALIFORNIA";
    else if State="CO"   then state="COLORADO";
    else if State="CT"   then state="CONNECTICUT";
    else if State="DE"   then state="DELAWARE";
    else if State="FL"   then state="FLORIDA";
    else if State="GA"   then state="GEORGIA";
    else if State="HI"   then state="HAWAII";
    else if State="IA"   then state="IOWA";
    else if State="ID"   then state="IDAHO";
    else if State="IL"   then state="ILLINOIS";
    else if State="IN"   then state="INDIANA";
    else if State="KS"   then state="KANSAS";
    else if State="KY"   then state="KENTUCKY";
    else if State="LA"   then state="LOUISIANA";
    else if State="MA"   then state="MASSACHUSETTS";
    else if State="MD"   then state="MARYLAND";
    else if State="ME"   then state="MAINE";
    else if State="MI"   then state="MICHIGAN";
    else if State="MN"   then state="MINNESOTA";

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else if State="MO" then state="MISSOURI";
else if State="MS" then state="MISSISSIPPI";
else if State="MT" then state="MONTANA";
else if State="NC" then state="NORTH CAROLINA";
else if State="ND" then state="NORTH DAKOTA";
else if State="NE" then state="NEBRASKA";
else if State="NH" then state="NEW HAMPSHIRE";
else if State="NJ" then state="NEW JERSEY";
else if State="NM" then state="NEW MEXICO";
else if State="NV" then state="NEVADA";
else if State="NY" then state="NEW YORK";
else if State="OH" then state="OHIO";
else if State="OK" then state="OKLAHOMA";
else if State="OR" then state="OREGON";
else if State="PA" then state="PENNSYLVANIA";
else if State="RI" then state="RHODE ISLAND";
else if State="SC" then state="SOUTH CAROLINA";
else if State="SD" then state="SOUTH DAKOTA";
else if State="TN" then state="TENNESSEE";
else if State="TX" then state="TEXAS";
else if State="UT" then state="UTAH";
else if State="VA" then state="VIRGINIA";
else if State="VT" then state="VERMONT";
else if State="WA" then state="WASHINGTON";
else if State="WI" then state="WISCONSIN";
else if State="WV" then state="WEST VIRGINIA";
else if State="WY" then state="WYOMING";

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end;

drop MULENRFL;

run;

*****Assign Labels****/

data adxb;

```

retain STUDYID USUBJID SITEID SUBJID AGEGR1N AGEGR1 SEX RACE ETHNIC COUNTRY STATE
ARMCD ARM ACTARMCD ACTARM RANDDT TRTSDT TRTEDT TRT01A TRT01AN TRT01P
TRT01PN

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VAX101DT VAX102DT DVSTDT UNBLNDDT RANDFL EVALEFFL AAI1EFFL AAI2EFFL
MULENRFL PHASE PHASEN HIVFL PARAMN PARAMCD PARAM PARCAT1 AVAL AVALC
AVISITN AVISIT

```

ADT ADY PDP27FL FC19D27 ;

set adxb_2;

label

PARAMN = "Parameter (N)"

PARAMCD = "Parameter Code"

PARAM = "Parameter"

PARCAT1 = "Parameter Category 1"

AVAL = "Analysis Value"

AVALC = "Analysis Value (C)"

AVISITN = "Analysis Visit (N)"

AVISIT = "Analysis Visit"

ADT = "Analysis Date"

ADY = "Analysis Relative Day"

FC19D27 = "First COVID Onset 7D Post D2 (PD)"

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STATE  ="State"
;
run;

*****
* Output datasets *;
*****
proc sort data=adxb out=datvout.adxb(label="Sequencing Analysis Dataset");
   by studyid usubjid paramn adt;
run;

proc printto;
run;
```