```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.1.1
     Notes:
                                       Re-programming sample tables as
part of QC.
*/
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
/*Create macro variables of dataset variables to access later*/
%let vartestcd="BMI","WEIGHT","HEIGHT";
%let varlist=usubjid age sex race ethnic;
%let varlist2=usubjid vstestcd vsstresn visit;
/*Create subset of dm dataset with variables needed for table*/
data dm1;
set r.dm(keep=&varlist)
r.vs(keep=&varlist2 where=(vstestcd in (&vartestcd) and
visit="SCREENING"));
by usubjid;
run;
/*Transpose dataset*/
proc transpose data=dm1 out=dm2;
by usubjid;
 id vstestcd;
 var vsstresn;
 where vstestcd^='';
/*Merge back with Transposed dataset*/
data dm2;
merge dm1(drop=vstestcd vsstresn) dm2;
by usubjid;
drop name;
run;
/*Sort to keep unique records per subject*/
proc sort data=dm2 nodupkey;
by usubjid;
run;
/*Prepare data for variables age weight height and bmi */
%macro prep;
```

```
%let var1=age;
%let var2=weight;
%let var3=height;
%let var4=bmi;
%do i=1 %to 4;
proc univariate data=dm2 noprint;
   var &&var&i;
   output out=&&var&i n=N mean=MEAN std=SD min=MIN median=MEDIAN
max=MAX;
run;
%end;
%do i=1 %to 4;
data &&var&i;
length varname $10.;
set &&var&i;
varname=upcase("&&var&i");
run;
proc transpose data=&&var&i out=&&var&i;
id varname;
var numeric;
run;
data &&var&i;
length order1 8.;
set &&var&i;
order1=&i;
run;
data &&var&i;
length statual $30.;
format statval $30.;
length dmvar $30.;
set &&var&i;
if order1=1 and NAME ^='SD' then statval=put(&&var&i,14.);
if NAME ='N' then statval=put(&&var&i,14.);
if NAME = 'SD' and order1=1 then statval=put(&&var&i,14.1);
if NAME = 'SD' and order1^=1 then statval=put(&&var&i,14.2);
if order1^=1 and NAME not in ('N','SD') then
statval=trim(put(round(&&var&i,.1),14.1));
if NAME ='N'
                     then order2=1;
if NAME = 'MEAN' then order2=2;
if NAME = 'SD'
                     then order2=3;
if NAME = 'MIN'
                     then order2=4;
if _NAME_='MEDIAN' then order2=5;
if NAME = 'MAX'
                     then order2=6;
if order1=1 then dmvar="Age (years)";
if order1=2 then dmvar="Weight (kg)";
if order1=3 then dmvar="Height (cm)";
```

```
if order1=4 then dmvar="Body Mass Index (kg/m2)";
drop &&var&i;
run;
%end;
%mend prep;
%prep;
/*Prepare data for variables sex, ethnicity, and race*/
%macro prep2;
%let var1=sex;
%let var2=ethnic;
%let var3=race;
%do i=1 %to 3;
proc freq data=dm2 noprint;
   tables &&var&i / out=&&var&i missing;
run;
data &&var&i;
length statual $30.;
length dmvar $30.;
length dmvar2 $30.;
set &&var&i;
statval=put(count, 8.) | | " (" | | trim(right(put(percent, 5.1))) | | "%) ";
if &&var&i=" " then &&var&i="Unknown";
if &i=1 then do
dmvar="Gender:";
end:
else if &i=2 then do
dmvar="Ethnicity:";
end:
else if \&i=3 then do
dmvar="Race:";
end;
dmvar2=propcase(&&var&i);
NAME ="n (%)";
drop &&var&i percent;
run;
%end;
/*proc sql;
insert into sex set dmvar="Gender";
insert into ethnic set dmvar="Ethnicity";
insert into race set dmvar="Race";
quit;
```

```
*/
%do i=1 %to 3;
data &&var&i;
set &&var&i;
order1=&i+4;
%end;
%mend prep2;
%prep2;
/*Format for Demographic variables*/
proc format;
  value $dmvar
                'F'='Female'
                      'M'='Male'
                      'Hispanic Or Latino'='Hispanic or Latino'
                      'Not Hispanic Or Latino'='Not Hispanic or Latino'
                      'Black Or African American'='Black or African
American'
                      'Unknown'='Unknown'
                      'Body Mass Index (kg/m2)'='Body Mass Index
(kg/m^{super 2})'
                      'Other'='Other';
run;
/*Final Report dataset creation*/
Data fnlreport(drop=count);
Set age(in=i1) weight(in=i2) height(in=i3) bmi(in=i4) sex(in=i5)
ethnic(in=i6) race(in=i7);
If trim(upcase(dmvar2)) =: trim(upcase('m'))
                                                                    then
order2=1;
If trim(upcase(dmvar2)) =: trim(upcase('f'))
                                                                    then
order2=2;
If trim(upcase(dmvar2))=:upcase('hispanic or latino')
                                                             then
order2=1;
If trim(upcase(dmvar2))=:upcase('not hispanic or latino') then
order2=2;
If trim(upcase(dmvar2)) = :upcase('unknown')
                                                                    then
order2=9;
If trim(upcase(dmvar2)) = : upcase('white')
     then order2=1;
If trim(upcase(dmvar2))=:upcase('black or african american')
                                                                    then
order2=2;
If trim(upcase(dmvar2)) = : upcase('other')
     then order2=3;
If trim(upcase(dmvar2))=:upcase('unknown')
     then order2=9;
If order1 < 5 and order2^=1 then dmvar='';</pre>
```

```
/*page counter*/
counter = n ;
mypage = 1;
if counter > 24 then do;
counter+1;
mypage+1;
end;
Run;
options nodate nonumber;
proc sort data=fnlreport;
by order1 order2;
run;
/*Create internal labels for report variables*/
data fnlreport;
set fnlreport;
label dmvar="Demographic";
label name ="Statistic";
label statval="Summary Result";
if order2^=1 then dmvar='';
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull',today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
proc sort data=fnlreport;
by order1 order2 mypage dmvar;
run;
/*Create blank rows after each demographic variable data*/
data fnlreport;
set fnlreport;
output;
by order1;
if last.order1;
array allnums [*] numeric ;
array allchar [*] character ;
do i=1 to dim(allnums); allnums(i)=.; end;
do i=1 to dim(allchar); allchar{i}=' '; end;
```

```
output; /* Output blank observation */
run;
/* Proc Report section */
/* Run two proc reports simultaneously to get desired output */
/* Manual post processing done to the output by removing an empty page
and a empty row */
ods escapechar='^';
ods listing close;
ods rtf
file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
1-1.rtf" bodytitle style=Styles.custom;
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.152in bordercolor=black}
     style(header) = {font face=arial font size=2.5 cellheight=.152in
foreground=black bordercolor=black}
     columns (order1 dmvar name statval mypage);
     where mypage=1;
     define order1 /order order = internal noprint;
     define dmvar /display "Demographic" format=$dmvar.
style={cellwidth=20% just=1};
     define name /display center "Statistic" style={cellwidth=35%
just=c};
     define statual /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=22.5%
just=dec);
     define mypage /order group style(column)={cellwidth=22%
foreground=white background=white} " ";
     compute after _page_;
     line @1 " ";
     endcomp;
     compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.1-1
```

```
DATE: &dtnull2
```

```
TIME: & systime";
     title2 j=c "Descriptive Statistics of Screening Demographics:
Enrolled Population";
     title3 j=c "(Page 1 of 2)";
     *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Enrolled
";
     footnote j=1 "Source: Listing 16.2.4-4a";
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.2in bordercolor=black}
     style(header)={font face=arial font size=2.5 cellheight=.15in
foreground=black bordercolor=black}
     columns (order1 order2 dmvar dmvar2 name statval mypage );
     where mypage=2;
     define order1 /order order = internal noprint;
     define order2 /order = internal noprint;
define dmvar /display "Demographic" format=$dmvar.
style={cellwidth=10% just=1};
     define dmvar2 /display left " " format=$dmvar.
style={cellwidth=25% just=1};
     define name /display center "Number (%) of Subjects"
style={cellwidth=32% just=c};
     define statval /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=17.% just=dec};
     define mypage /order group style(column)={cellwidth=15%
foreground=white background=white} " ";
     compute after page;
     line @1 " ";
     endcomp;
     compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                 Table 14.1-1
```

DATE: &dtnull2

```
TIME:&systime";
    title2 j=c "Descriptive Statistics of Screening Demographics:
Enrolled Population";
    title3 j=c "(Page 2 of 2)";
    *title3 j=c "Page ^{thispage} of ^{lastpage}";
    title4;
    title5 j=l "Population: Enrolled
";
    footnote j=l "Source: Listing 16.2.4-4a";
run;quit;
ods listing;
ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
     Date:
                                       06-18-**
     Study:
                                       OTR1021
                                       14.1.2
     Table:
     Notes:
                                       Re-programming sample tables as
part of QC.
* /
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
/*Create macro variables of dataset variables to access later*/
%let vartestcd="BMI","WEIGHT","HEIGHT";
%let varlist=usubjid age sex race ethnic;
%let varlist2=usubjid vstestcd vsstresn visit;
/*Create subset of dm dataset with variables needed for table*/
data dm1;
set r.dm(keep=&varlist)
r.vs(keep=&varlist2 where=(vstestcd in (&vartestcd) and
visit="SCREENING"));
by usubjid;
run;
proc sql noprint;
delete from dm1
where usubjid not in (select distinct usubjid from r.dm where armcd ^=
"SCRNFAIL"
and usubjid in (select usubjid from r.ex where usubjid in
(select distinct usubjid from r.vs where vstptnum > 6)));
quit;
/*Transpose dataset*/
proc transpose data=dm1 out=dm2;
by usubjid;
id vstestcd;
var vsstresn;
where vstestcd^='';
run;
/*Merge back with Transposed dataset*/
data dm2;
merge dm1(drop=vstestcd vsstresn) dm2;
by usubjid;
drop name;
run;
```

```
/*Sort to keep unique records per subject*/
proc sort data=dm2 nodupkey;
by usubjid;
run;
/*Prepare data for variables age weight height and bmi */
%macro prep;
%let var1=age;
%let var2=weight;
%let var3=height;
%let var4=bmi;
%do i=1 %to 4;
proc univariate data=dm2 noprint;
   var &&var&i;
   output out=&&var&i n=N mean=MEAN std=SD min=MIN median=MEDIAN
max=MAX;
run;
%end;
%do i=1 %to 4;
data &&var&i;
length varname $10.;
set &&var&i;
varname=upcase("&&var&i");
/* Transpose numeric variables */
proc transpose data=&&var&i out=&&var&i;
id varname;
var numeric;
run;
data &&var&i;
length order1 8.;
set &&var&i;
order1=&i;
run;
/* Create char variable statval to store stat values with appropriate
formats */
data &&var&i;
length statval $30.;
format statual $30.;
length dmvar $30.;
set &&var&i;
if order1=1 and NAME ^='SD' then statval=put(&&var&i,14.);
if NAME ='N' then statval=put(&&var&i,14.);
if NAME = 'SD' and order1=1 then statval=put(&&var&i,14.1);
if NAME = 'SD' and order1^=1 then statval=put(&&var&i,14.2);
```

```
if order1^=1 and NAME not in ('N','SD') then
statval=trim(put(round(&&var&i,.1),14.1));
if NAME = 'N'
                     then order2=1;
if NAME = 'MEAN' then order2=2;
if NAME = 'SD'
                     then order2=3;
if NAME = 'MIN'
                     then order2=4;
if NAME = 'MEDIAN'
                    then order2=5;
if NAME = 'MAX'
                     then order2=6;
if order1=1 then dmvar="Age (years)";
if order1=2 then dmvar="Weight (kg)";
if order1=3 then dmvar="Height (cm)";
if order1=4 then dmvar="Body Mass Index (kg/m2)";
drop &&var&i;
run;
%end;
%mend prep;
%prep;
/* Macro to prepare data for variables sex, ethnicity, and race*/
%macro prep2;
%let var1=sex;
%let var2=ethnic;
%let var3=race;
%do i=1 %to 3;
proc freq data=dm2 noprint;
   tables &&var&i / out=&&var&i missing;
/* Create char variable statval to store stat values with appropriate
formats */
data &&var&i;
length statual $30.;
length dmvar $30.;
length dmvar2 $30.;
set &&var&i;
statval=put(count, 8.) | | " ("||trim(right(put(percent, 5.1))) | | "%) ";
if &&var&i=" " then &&var&i="Unknown";
if &i=1 then do
dmvar="Gender:";
end:
else if &i=2 then do
dmvar="Ethnicity:";
end;
else if &i=3 then do
```

```
dmvar="Race:";
end:
dmvar2=propcase(&&var&i);
NAME ="n (%)";
drop &&var&i percent;
run;
%end;
/*proc sql;
insert into sex set dmvar="Gender";
insert into ethnic set dmvar="Ethnicity";
insert into race set dmvar="Race";
quit;
*/
%do i=1 %to 3;
data &&var&i;
set &&var&i;
order1=&i+4;
run;
%end;
%mend prep2;
%prep2;
/*Format for Demographic variables*/
proc format;
 value $dmvar 'F'='Female'
                      'M'='Male'
                      'Hispanic Or Latino'='Hispanic or Latino'
                      'Not Hispanic Or Latino'='Not Hispanic or Latino'
                      'Black Or African American'='Black or African
American'
                      'Unknown'='Unknown'
                      'Body Mass Index (kg/m2)'='Body Mass Index
(kg/m^{super 2})'
                      'Other'='Other';
run;
/*Final Report dataset creation*/
Data fnlreport(drop=count);
Set age(in=i1) weight(in=i2) height(in=i3) bmi(in=i4) sex(in=i5)
ethnic(in=i6) race(in=i7);
If trim(upcase(dmvar2)) =: trim(upcase('m'))
                                                                   then
order2=1;
If trim(upcase(dmvar2))=:trim(upcase('f'))
                                                                   then
order2=2;
If trim(upcase(dmvar2))=:upcase('hispanic or latino') then
order2=1;
```

```
If trim(upcase(dmvar2))=:upcase('not hispanic or latino') then
order2=2;
If trim(upcase(dmvar2)) =: upcase('unknown')
                                                                    then
order2=9;
If trim(upcase(dmvar2)) =: upcase('white')
     then order2=1;
If trim(upcase(dmvar2))=:upcase('black or african american')
                                                                    then
order2=2;
If trim(upcase(dmvar2))=:upcase('other')
     then order2=3;
If trim(upcase(dmvar2)) =: upcase('unknown')
     then order2=9;
If order1 < 5 and order2^=1 then dmvar='';</pre>
/*page counter*/
counter = n ;
mypage = 1;
if counter > 24 then do;
counter+1;
mypage+1;
end;
Run;
options nodate nonumber;
proc sort data=fnlreport;
by order1 order2;
run;
/*Create internal labels for report variables*/
data fnlreport;
set fnlreport;
label dmvar="Demographic";
label name ="Statistic";
label statval="Summary Result";
if order2^=1 then dmvar='';
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
proc sort data=fnlreport;
by order1 order2 mypage dmvar;
run;
```

```
/*Create blank rows after each demographic variable data*/
data fnlreport;
set fnlreport;
output;
by order1;
if last.order1;
array allnums [*] numeric;
array allchar [*] character ;
drop i;
do i=1 to dim(allnums); allnums(i)=.; end;
do i=1 to dim(allchar); allchar{i}=' '; end;
output; /* Output blank observation */
run;
/* Proc Report section */
/* Run two proc reports simultaneously to get desired output */
/* Manual post processing done to the output by removing an empty page
and a empty row */
ods escapechar='^';
options orientation=landscape;
ods listing close;
ods rtf
file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
1-2.rtf" bodytitle style=Styles.custom;
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.152in bordercolor=black}
     style(header) = {font face=arial font size=2.5 cellheight=.152in
foreground=black bordercolor=black}
     ;
     columns (order1 dmvar name statval mypage);
     where mypage=1;
     define order1 /order order = internal noprint;
                    /display "Demographic" format=$dmvar.
     define dmvar
style={cellwidth=20% just=1};
     define name /display center "Statistic" style={cellwidth=35%
just=c};
     define statual /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=22.5%
just=dec};
```

```
define mypage /order group style(column)={cellwidth=22%
foreground=white background=white} " ";
     compute after page;
     line @1 " ";
     endcomp;
     compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.1-2
                                                   DATE: &dtnull2
TIME: &systime";
     title2 j=c "Descriptive Statistics of Screening Demographics:
Randomized Safety Population";
     title3 j=c "(Page 1 of 2)";
    *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety
";
     footnote j=1 "Source: Listing 16.2.4-4b";
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.2in bordercolor=black}
     style(header)={font face=arial font size=2.5 cellheight=.15in
foreground=black bordercolor=black}
     columns (order1 order2 dmvar dmvar2 name statval mypage );
     where mypage=2;
     define order1 /order order = internal noprint;
     define order2 /order = internal noprint;
     define dmvar /display "Demographic" format=$dmvar.
style={cellwidth=10% just=1};
     define dmvar2 /display left " " format=$dmvar.
style={cellwidth=25% just=1};
     define name /display center "Number (%) of Subjects"
style={cellwidth=32% just=c};
```

```
define statual /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=17.% just=dec};
     define mypage /order group style(column)={cellwidth=15%
foreground=white background=white} " ";
     compute after _page_;
     line @1 " ";
     endcomp;
     compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.1-2
                                                    DATE: &dtnull2
TIME: & systime";
     title2 j=c "Descriptive Statistics of Screening Demographics:
Randomized Safety Population";
     title3 j=c "(Page 2 of 2)";
     *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety
";
     footnote j=1 "Source: Listing 16.2.4-4b";
run; quit;
ods listing;
ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.1.3
     Notes:
                                       Re-programming sample tables as
part of QC.
*/
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
/*Create macro variables of dataset variables to access later*/
%let vartestcd="BMI","WEIGHT","HEIGHT";
%let varlist=usubjid age sex race ethnic;
%let varlist2=usubjid vstestcd vsstresn visit;
/*Create subset of dm dataset with variables needed for table*/
data dm1;
set r.dm(keep=&varlist)
r.vs(keep=&varlist2 where=(vstestcd in (&vartestcd) and
visit="SCREENING"));
by usubjid;
run;
proc sql noprint;
delete from dm1
where usubjid not in (select distinct usubjid from r.dm where armcd ^=
"SCRNFAIL"
and usubjid in (select usubjid from r.ex where usubjid in
(select distinct usubjid from r.pp)));
quit;
/*Transpose dataset*/
proc transpose data=dm1 out=dm2;
by usubjid;
id vstestcd;
var vsstresn;
where vstestcd^='';
run;
/*Merge back with Transposed dataset*/
data dm2;
merge dm1(drop=vstestcd vsstresn) dm2;
by usubjid;
drop name;
run;
```

```
/*Sort to keep unique records per subject*/
proc sort data=dm2 nodupkey;
by usubjid;
run;
/*Prepare data for variables age weight height and bmi */
%macro prep;
%let var1=age;
%let var2=weight;
%let var3=height;
%let var4=bmi;
%do i=1 %to 4;
proc univariate data=dm2 noprint;
   var &&var&i;
   output out=&&var&i n=N mean=MEAN std=SD min=MIN median=MEDIAN
max=MAX;
run;
%end;
%do i=1 %to 4;
data &&var&i;
length varname $10.;
set &&var&i;
varname=upcase("&&var&i");
run;
proc transpose data=&&var&i out=&&var&i;
id varname;
var numeric;
run;
data &&var&i;
length order1 8.;
set &&var&i;
order1=&i;
run;
data &&var&i;
length statval $30.;
format statval $30.;
length dmvar $30.;
set &&var&i;
if order1=1 and NAME ^='SD' then statval=put(&&var&i,14.);
if NAME ='N' then statval=put(&&var&i,14.);
if NAME = 'SD' and order1=1 then statval=put(&&var&i,14.1);
if NAME = 'SD' and order1^=1 then statval=put(&&var&i,14.2);
if order1^=1 and NAME not in ('N','SD') then
statval=trim(put(round(&&var&i,.1),14.1));
if NAME ='N'
                      then order2=1;
if NAME = 'MEAN' then order2=2;
```

```
if NAME = 'SD'
                     then order2=3;
if _NAME ='MIN'
                     then order2=4;
if NAME = 'MEDIAN'
                     then order2=5;
if NAME = 'MAX'
                      then order2=6;
if order1=1 then dmvar="Age (years)";
if order1=2 then dmvar="Weight (kg)";
if order1=3 then dmvar="Height (cm)";
if order1=4 then dmvar="Body Mass Index (kg/m2)";
drop &&var&i;
run;
%end;
%mend prep;
%prep;
/*Prepare data for variables sex, ethnicity, and race*/
%macro prep2;
%let var1=sex;
%let var2=ethnic;
%let var3=race;
%do i=1 %to 3;
proc freq data=dm2 noprint;
   tables &&var&i / out=&&var&i missing;
run;
data &&var&i;
length statval $30.;
length dmvar $30.;
length dmvar2 $30.;
set &&var&i;
statval=put(count, 8.) || " ("||trim(right(put(percent, 5.1))) || "%) ";
if &&var&i=" " then &&var&i="Unknown";
if &i=1 then do
dmvar="Gender:";
end;
else if &i=2 then do
dmvar="Ethnicity:";
else if &i=3 then do
dmvar="Race:";
end;
dmvar2=propcase(&&var&i);
NAME ="n (%)";
drop &&var&i percent;
run;
```

```
%end;
/*proc sql;
insert into sex set dmvar="Gender";
insert into ethnic set dmvar="Ethnicity";
insert into race set dmvar="Race";
quit;
*/
%do i=1 %to 3;
data &&var&i;
set &&var&i;
order1=&i+4;
run;
%end;
%mend prep2;
%prep2;
/*Format for Demographic variables*/
proc format;
  value $dmvar 'F'='Female'
                      'M'='Male'
                      'Hispanic Or Latino'='Hispanic or Latino'
                      'Not Hispanic Or Latino'='Not Hispanic or Latino'
                      'Black Or African American'='Black or African
American'
                      'Unknown'='Unknown'
                      'Body Mass Index (kg/m2)'='Body Mass Index
(kq/m^{super 2})'
                      'Other'='Other';
run;
/*Final Report dataset creation*/
Data fnlreport(drop=count);
Set age(in=i1) weight(in=i2) height(in=i3) bmi(in=i4) sex(in=i5)
ethnic(in=i6) race(in=i7);
If trim(upcase(dmvar2))=:trim(upcase('m'))
                                                                    then
order2=1;
If trim(upcase(dmvar2)) =: trim(upcase('f'))
                                                                    then
order2=2;
If trim(upcase(dmvar2))=:upcase('hispanic or latino')
                                                             then
order2=1;
If trim(upcase(dmvar2))=:upcase('not hispanic or latino') then
order2=2;
If trim(upcase(dmvar2)) = :upcase('unknown')
                                                                    then
order2=9;
If trim(upcase(dmvar2))=:upcase('white')
     then order2=1;
```

```
If trim(upcase(dmvar2)) =: upcase('black or african american')
order2=2;
If trim(upcase(dmvar2)) = : upcase('other')
     then order2=3;
If trim(upcase(dmvar2)) =: upcase('unknown')
     then order2=9;
If order1 < 5 and order2^=1 then dmvar='';</pre>
/*page counter*/
counter = _n_;
mypage = 1;
if counter > 24 then do;
counter+1;
mypage+1;
end;
Run;
options nodate nonumber;
proc sort data=fnlreport;
by order1 order2;
run;
/*Create internal labels for report variables*/
data fnlreport;
set fnlreport;
label dmvar="Demographic";
label name ="Statistic";
label statval="Summary Result";
if order2^=1 then dmvar='';
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull',today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
proc sort data=fnlreport;
by order1 order2 mypage dmvar;
run;
/*Create blank rows after each demographic variable data*/
data fnlreport;
set fnlreport;
```

```
output;
by order1;
if last.order1;
array allnums [*] _numeric_;
array allchar [*] character ;
drop i;
do i=1 to dim(allnums); allnums(i)=.; end;
do i=1 to dim(allchar); allchar{i}=' '; end;
output; /* Output blank observation */
run;
/* Proc Report section */
/* Run two proc reports simultaneously to get desired output */
/* Manual post processing done to the output by removing an empty page
and a empty row */
ods escapechar='^';
options orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
1-3.rtf" bodytitle style=Styles.custom;
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.152in bordercolor=black}
     style(header) = {font face=arial font size=2.5 cellheight=.152in
foreground=black bordercolor=black}
     columns (order1 dmvar name statval mypage);
     where mypage=1;
     define order1 /order order = internal noprint;
                   /display "Demographic" format=$dmvar.
     define dmvar
style={cellwidth=20% just=1};
     define name /display center "Statistic" style={cellwidth=35%
just=c};
     define statual /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=22.5%
just=dec};
     define mypage /order group style(column)={cellwidth=22%
foreground=white background=white} " ";
     compute after page;
     line @1 " ";
     endcomp;
```

```
compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.1-3
                                                   DATE: &dtnull2
TIME: & systime";
     title2 j=c "Descriptive Statistics of Screening Demographics:
Full Analysis Population";
     title3 j=c "(Page 1 of 2)";
    *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Full Analysis
";
     footnote j=1 "Source: Listing 16.2.4-4c";
proc report
     data = fnlreport
     nowd
     ls=122
     ps=30
     headline
     headskip
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.2in bordercolor=black}
     style(header)={font face=arial font size=2.5 cellheight=.15in
foreground=black bordercolor=black}
     columns (order1 order2 dmvar dmvar2 name statval mypage );
     where mypage=2;
     define order1 /order order = internal noprint;
     define order2 /order = internal noprint;
     define dmvar /display "Demographic" format=$dmvar.
style={cellwidth=10% just=1};
     define dmvar2 /display left " " format=$dmvar.
style={cellwidth=25% just=1};
     define name /display center "Number (%) of Subjects"
style={cellwidth=32% just=c};
     define statual /display "Summary Result" center
style={protectspecialchars=off cellpadding=4 cellwidth=17.% just=dec};
     define mypage /order group style(column)={cellwidth=15%
foreground=white background=white} " ";
     compute after page;
     line @1 " ";
```

```
endcomp;
     compute before mypage;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.1-3
                                                    DATE: &dtnull2
TIME: & systime";
     title2 j=c "Descriptive Statistics of Screening Demographics:
Full Analysis Population";
     title3 j=c "(Page 2 of 2)";
     *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=l "Population: Full Analysis
";
     footnote j=1 "Source: Listing 16.2.4-4c";
 run; quit;
ods listing;
 ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
     Date:
                                       06-18-**
     Study:
                                       OTR1021
     Table:
                                       14.1.7
     Notes:
                                       Re-programming sample tables as
part of QC.
* /
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
data ex;
length trtc $1;
set r.ex;
if EXDOSTXT="FINELY CRUSHED 10 MG OTR" then trtc="A";
if EXDOSTXT="COARSELY CRUSHED 10 MG OTR" then trtc="B";
if EXDOSTXT="FINELY CRUSHED 10 MG OC" then trtc="C";
run;
proc sql;
create table trta as select epoch, count(trtc) as trta from ex
where trtc="A"
group by 1
order by 1,2;
create table trtb as select epoch, count(trtc) as trtb from ex
where trtc="B"
group by 1
order by 1,2;
create table trtc as select epoch, count(trtc) as trtc from ex
where trtc="C"
group by 1
order by 1,2;
quit;
data overall;
length epoch2 8;
merge trta trtb trtc;
by epoch;
epoch2=put(substr(epoch, 8, 1), 8.);
drop epoch;
rename epoch2=epoch;
run;
proc transpose data=overall out=overall;
var _all_;
run;
```

```
data overall;
length col4 8;
set overall;
col4=sum(col1,col2,col3);
proc transpose data=overall out=overall2;
id name;
var _numeric_;
run;
proc sort data=overall2;
by epoch;
run;
data overall3:
length overall $25;
length stat1-stat3 $30;
set overall2;
by epoch;
if epoch^=6 then do;
Overall=trim(put(sum(trta,trtb,trtc),8.))||'
('||put(trim(put(sum(trta,trtb,trtc),5.2))/trim(put(sum(trta,trtb,trtc
),5.2))*100,5.2)||'%)';
end;
if epoch^=6 then do;
stat1=trta||' ('||put(trta/sum(trta,trtb,trtc)*100,5.1)||'%)';
stat2=trtb||' ('||put(trtb/sum(trta,trtb,trtc)*100,5.1)||'%)';
stat3=trtc||' ('||put(trtc/sum(trta,trtb,trtc)*100,5.1)||'%)';
end;
if epoch=6 then do;
stat1=trta||' ('||put(trta/30*100,5.1)||'%)';
stat2=trtb||' ('||put(trtb/30*100,5.1)||'%)';
stat3=trtc||' ('||put(trtc/30*100,5.1)||'%)';
end;
run;
options nonumber;
data _null ;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
proc format;
     value epoch
           6="Overall";
run;
```

```
ods escapechar='^';
ods listing close;
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
1-7.rtf" bodytitle style=Styles.custom;
proc report
     data = overall3
     nowd
     ls=122
     ps=30
     headline
     headskip
     split='*'
     style(report)={just=left font face=arial font size=2.5
bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.275in bordercolor=black}
     style(header) = {font face=arial font size=2.5 cellheight=.3in
foreground=black bordercolor=black}
     columns (epoch ('Number (%) of
Subjects*
                                 stat1 stat2 stat3
overall));
     define epoch /order order = internal "Study Period" center
format=epoch. style={cellwidth=19% just=c};;
     define stat1 /display "Treatment A" center style={cellwidth=20%
just=c};
     define stat2 /display "Treatment B" center style={cellwidth=20%
just=c};
     define stat3 /display "Treatment C" center style={cellwidth=20%
just=c};
     define overall /display "Overall" center style={cellwidth=20%
just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after ;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.1-7
                                                DATE: &dtnull2
TIME: & systime";
     title2 j=c "Study Treatment Administration by Study Period:
Randomized Safety Population";
     title3 j=c "(Page 1 of 1)";
```

```
*title3 j=c "Page ^{thispage} of ^{lastpage}";
    title4;
    title5 j=l "Population: Randomized Safety
";
    footnote j=l "NOTE: TreatmentA = Finely crushed 10mg OTR;
TreatmentB = Coarsely crushed 10mg OTR; TreatmentC = Finely crushed 10mg OC.";
    footnote2 j=l "Source: Listing 16.2.5-2c";
run;
ods listing;
ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
     Date:
                                       06-18-**
     Study:
                                       OTR1021
                                       14.2.2-1
     Table:
     Notes:
                                       Re-programming sample tables as
part of QC.
                                       Output table format from this
program is not completely per specs
* /
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
/* Copy EX dataset to work folder and assign single letter treatment
codes */
data ex;
length treatc $1;
set r.ex(keep=studyid usubjid exseq exdostxt exstdtc exendtc epoch
visitnum);
if EXDOSTXT="FINELY CRUSHED 10 MG OTR" then treatc="A";
if EXDOSTXT="COARSELY CRUSHED 10 MG OTR" then treatc="B";
if EXDOSTXT="FINELY CRUSHED 10 MG OC" then treatc="C";
drop exdostxt;
run;
data pp;
set r.pp;
run;
proc transpose data=pp out=transpp;
by usubjid ppgrpid;
id pptestcd;
var ppstresn;
run;
data transpp2(drop=llambda ulambda nlambda name sccat);
merge transpp(in=a) r.sc(in=b keep=usubjid scorres SCCAT
where=(sccat=:"RANDOMIZATION NUMBER"));
by usubjid;
if a and b;
run;
data transpp2;
length ppgrpid2 8;
set transpp2;
ppgrpid2=put(ppgrpid,8.);
```

```
drop ppgrpid;
rename ppgrpid2=ppgrpid;
run;
data ppex;
merge transpp2 ex(keep=usubjid treatc exseq rename=(exseq=ppgrpid));
by usubjid ppgrpid;
run;
proc sort data=ppex;
by treatc;
run;
proc univariate data=ppex;
by treatc;
var AUC0 T;
output n=N mean=amean std=SD min=MIN median=med max=MAX cv=cvar;
run;
proc univariate data=ppex;
by treatc;
var cmax;
output out=cmax n=N mean=amean std=SD min=MIN median=med max=MAX
cv=cvar;
run;
proc transpose data=cmax out=cmaxtrans(rename=(col1=cmax));
by treatc;
var numeric;
run;
option nolabel spool;
proc sql flow number;
create table final2 as select treatc,
trim(trim(scorres)||'/'||trim(substr(usubjid,15,5))) as randsubject
length=15,
ppgrpid, AUCO T, AUCO INF, CMAX, TMAX, LAMBDAZ, TERMHALF, TLAG from ppex
order by 1,2;
quit;
%macro prep 14221;
%let stat1=AUC0 T;
%let stat2=AUC0 INF;
```

```
%let stat3=CMAX;
%let stat4=TMAX;
%let stat5=LAMBDAZ;
%let stat6=TERMHALF;
%let stat7=TLAG;
%DO I=1 %TO 7;
proc univariate data=ppex;
by treatc;
var &&stat&i;
output out=&&stat&i n= N mean= MEAN std= SD min= MIN median= MEDIAN
max= MAX cv= CV;
run;
proc transpose data=&&stat&i out=&&stat&i(rename=(col1=&&stat&i));
by treatc;
var _numeric_;
run;
proc sort data=&&stat&i;
by treatc name;
run;
data &&stat&i;
set &&stat&i;
*&&stat&i=round(&&stat&i,.5);
format &&stat&i best12.;
%end;
data final;
merge &stat1 &stat2 &stat3 &stat4 &stat5 &stat6 &stat7;
by treatc name;
run;
%mend prep 14221;
%prep 14221;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
```

```
/* Proc Report Step */
dm "odsresults; clear;";
options nonumber nodate orientation=landscape;
ods listing close;
ods escapechar='^';
ods rtf
file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
2-2-1.rtf" style=Styles.custom;
proc report
     data = final2
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.4in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns treatc randsubject ppgrpid AUC0 T AUC0 INF CMAX TMAX
LAMBDAZ TERMHALF TLAG;
     where treatc="A";
     define treatc / group "Treatment" style={cellwidth=10% just=c};
     define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define ppgrpid / display "Period" style={cellwidth=8% just=c};
     define AUCO T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUC0 INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
```

```
title1 j=l "OTR1021
                Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=l "Population: Full Analysis
";
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote4;
     footnote5 j=1 "Source: Listing 16.2.6-4";
run; quit;
proc report
     data = final
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column) = { just=center font face=arial font size=2.5
cellheight=.15in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns treatc name AUCO T AUCO INF CMAX TMAX LAMBDAZ TERMHALF
TLAG;
     where treatc="A";
     define treatc / group "Treatment" style={cellwidth=25% just=c};
     *define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define name / display "Period" style={cellwidth=10% just=c};
     define AUC0 T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUCO INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
```

```
define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Full Analysis
";
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote4;
     footnote5 j=1 "Source: Listing 16.2.6-4";
run; quit;
proc report
     data = final2
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.4in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
```

```
where treatc="B";
     define treatc / group "Treatment" style={cellwidth=10% just=c};
     define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define ppgrpid / display "Period" style={cellwidth=8% just=c};
     define AUC0 T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUCO INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=l "Population: Full Analysis
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote4:
```

```
footnote5 j=1 "Source: Listing 16.2.6-4";
 run; quit;
proc report
     data = final
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.15in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns treatc name AUCO T AUCO INF CMAX TMAX LAMBDAZ TERMHALF
TLAG;
     where treatc="B";
     define treatc / group "Treatment" style={cellwidth=25% just=c};
     *define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define _name_ / display "Period" style={cellwidth=10% just=c};
     define AUC0 T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUCO INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
```

```
title1 j=l "OTR1021
                Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Full Analysis
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote4;
     footnote5 j=1 "Source: Listing 16.2.6-4";
 run; quit;
proc report
     data = final2
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.4in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns treatc randsubject ppgrpid AUCO T AUCO INF CMAX TMAX
LAMBDAZ TERMHALF TLAG;
     where treatc="C";
     define treatc / group "Treatment" style={cellwidth=10% just=c};
     define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define ppgrpid / display "Period" style={cellwidth=8% just=c};
     define AUC0 T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUCO_INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
```

```
define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title5 j=l "Population: Full Analysis
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote5 j=1 "Source: Listing 16.2.6-4";
run; quit;
proc report
     data = final
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report) = {font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.15in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     ;
```

```
columns treatc name AUCO T AUCO INF CMAX TMAX LAMBDAZ TERMHALF
TLAG;
     where treatc="C";
     define treatc / group "Treatment" style={cellwidth=25% just=c};
     *define randsubject / display "Randomization Number/#Subject
Number" style={cellwidth=20% just=c};
     define _name_ / display "Period" style={cellwidth=10% just=c};
define AUCO_T / display "AUCt#(ng*h/mL)" format=best6.
style={cellwidth=9% just=c};
     define AUCO INF / display "AUCinf#(ng*h/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define CMAX / display "Cmax#(ng/mL)" format=best6.
style={cellwidth=8.5% just=c};
     define TMAX / display "tmax#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define LAMBDAZ / display "Lz#(1/h)" format=best6.
style={cellwidth=8.5% just=c};
     define TERMHALF / display "t1/2z#(h)" format=best6.
style={cellwidth=8.5% just=c};
     define TLAG / display "tlag#(h)" format=best6.
style={cellwidth=9% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                 Table 14.2.2-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Individual Subject and Mean Plasma Oxycodone
Pharmacokinetic Metrics";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=l "Population: Full Analysis
     footnote1 j=1 "NOTE: NA = Not applicable, NC = Not calculated";
     footnote2 j=1 '09'x "All Randomized Safety subjects are included
in the data listing. Only Full Analysis subjects are used in the
calculation of the summary statistics.";
     footnote3 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
```

```
footnote4;
     footnote5 j=1 "Source: Listing 16.2.6-4";
run; quit;
ods listing;
ods rtf close;
options nolabel;
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
/* Copy EX dataset to work folder and assign single letter treatment
codes */
data ex;
length treatc $1;
set r.ex(keep=studyid usubjid exseq exdostxt exstdtc exendtc epoch
visitnum);
if EXDOSTXT="FINELY CRUSHED 10 MG OTR" then treatc="A";
if EXDOSTXT="COARSELY CRUSHED 10 MG OTR" then treatc="B";
if EXDOSTXT="FINELY CRUSHED 10 MG OC" then treatc="C";
drop exdostxt;
run;
data qs;
set r.qs;
run;
proc sort data=qs;
by usubjid visitnum;
run;
proc sort data=ex;
by usubjid visitnum;
run;
data exqs;
merge qs ex(keep=usubjid epoch visitnum treatc);
by usubjid visitnum;
run;
```

```
proc sort data=exqs;
by usubjid treatc qstestcd qstptnum;
run;
data cfp;
     do until (last.usubjid);
           set exqs (where=(qsstresn^=. and visitnum=int(visitnum) and
qstestcd^="CALLMADE"));
           by usubjid treatc qstestcd qstptnum;
           if qstpt="PREDOSE" then pd=qsstresn;
                else do;
                      chgpd = qsstresn - pd;
                      output;
                end;
     end;
run;
proc sort data=cfp;
by treatc qstestcd qstest qstptnum qstpt;
run;
proc univariate data=cfp noprint round=0;
by treatc qstestcd qstest qstptnum qstpt;
var chqpd;
output out=cfpstat n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc sort data=exqs;
by treatc qstestcd qstest qstptnum qstpt;
run;
proc univariate data=exqs noprint round=0;
where qstestcd^="CALLMADE";
by treatc qstestcd qstest qstptnum qstpt;
var qsstresn;
output out=pdstat n=a N mean=b mean std=c sd min=d min median=e median
max=f max;
run;
proc sort data=pdstat;
by treatc qstestcd qstest qstptnum qstpt;
run;
```

```
proc transpose data=pdstat out=pdstat;
by treatc qstestcd qstest qstptnum qstpt;
var a N b mean c sd d min e median f max;
run;
data pdstat;
length qstpt $50;
set pdstat;
qstpt=propcase(qstpt);
proc sort data=cfpstat;
by treatc qstestcd qstest qstptnum qstpt;
run;
proc transpose data=cfpstat out=cfpstat;
by treatc qstestcd qstest qstptnum qstpt;
var a N b mean c sd d min e median f max;
run;
data cfpstat;
length qstpt $50;
set cfpstat;
if qstpt^="PREDOSE" then qstpt=propcase(trim(qstpt)||trim(", Change
from Predose"));
qstpt=propcase(qstpt);
run;
proc sort data=cfpstat;
by treatc qstestcd qstest qstptnum qstpt;
run;
proc sort data=pdstat;
by treatc qstestcd qstest qstptnum qstpt;
run;
data final;
set pdstat cfpstat;
by treatc qstestcd qstptnum;
run;
proc sort data=final;
by treatc qstestcd qstest qstptnum qstpt;
run;
proc format;
```

```
value $statf 'a N'='N'
                       'b mean'='MEAN'
                      'c sd'='SD'
                      'd min'='MIN'
                      'e median'='MEDIAN'
                      'f max'='MAX';
run;
data final;
length statual $30.;
format statual $30.;
set final;
if NAME ='b mean' then statval=put(round(col1,.5),5.0);
if NAME ='c sd' then statval=put(round(col1,0.01),5.1);
if NAME not in ('b mean','c sd') then
statval=put (round (col1, 1.0), \overline{5.0});
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull, 6, 4);
/* Proc Report Step */
dm "odsresults; clear;";
options nonumber nodate orientation=landscape;
ods listing close;
ods escapechar='^';
ods rtf file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\14-
2-4-1.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=39
     headline
     headskip
     split='#'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.15in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     ;
```

```
columns treatc qstestcd qstest qstptnum qstpt name statval;
     *where treatc="A";
     *where also qstestcd="NASAL01";
     define treatc / group order=internal "Treatment"
style={cellwidth=15% just=c};
     define qstestcd / group order=internal noprint;
     define qstest / group order=internal "Assessment"
style={cellwidth=15% just=c};
     define qstptnum / group order=internal noprint;
     define qstpt / group order=internal "Timepoint"
style={cellwidth=37% just=c};
     define name / display "Statistic" format=$statf.
style={cellwidth=15% just=c};
     define statval / display "Result" style={protectspecialchars=off
cellwidth=17% just=dec};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     compute after qstpt;
     line @1 " ";
     endcomp;
     break after qstptnum / suppress page;
     title1 j=l "OTR1021
                Table 14.2.4-1
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Descriptive Statistics and Change from Predose on
Intranasal Tolerability Rating Scale Data by Treatment";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety
";
     footnote1 j=1 "NOTE: Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote2 j=1 "Source: Listing 16.2.8-13.";
run; quit;
ods listing;
ods rtf close;
```

```
/*
     Title:
                                       OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.3.1-1a
     Notes:
                                       Re-programming sample tables as
part of QC.
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
/* Copy EX dataset to work folder and assign single letter treatment
codes */
data ex;
length treatc $1;
set r.ex(keep=studyid usubjid exseq exdostxt exstdtc exendtc epoch
visitnum);
if EXDOSTXT="FINELY CRUSHED 10 MG OTR" then treatc="A";
if EXDOSTXT="COARSELY CRUSHED 10 MG OTR" then treatc="B";
if EXDOSTXT="FINELY CRUSHED 10 MG OC" then treatc="C";
drop exdostxt;
run;
/* Population counts by treatment assigned to macro variables */
proc sql noprint;
select count(distinct usubjid) into :trta from ex where treatc="A";
select count(distinct usubjid) into :trtb from ex where treatc="B";
select count(distinct usubjid) into :trtc from ex where treatc="C";
select count(distinct usubjid) into :overall from ex where treatc in
("A", "B", "C");
quit;
/* Re-assign Population count macro variables to remove blanks */
%let trta=&trta;
%let trtb=&trtb;
%let trtc=&trtc;
%let overall=&overall;
/* Transpose ex dataset to get start and end dates of exposure by
treatment */
proc transpose data=ex out=trex(drop= name );
by usubjid;
id treatc ;
var exstdtc;
run;
/* Transpose ex dataset to get start and end dates of exposure by
treatment */
```

```
proc transpose data=ex out=trex2(drop= name rename=(a=paendt b=pbendt
c=pcendt));
by usubjid;
id treatc ;
var exendtc;
run;
/* Add :00 to the ae start and end dates to make use of the is8601
function later on the variables */
data ae;
set r.ae;
aestdtc2=trim(trim(aestdtc)||trim(":00"));
aeendtc2=trim(trim(aeendtc)||trim(":00"));
drop aestdtc aeendtc;
rename aestdtc2=aestdtc aeendtc2=aeendtc;
run;
/* Merge ae and transposed ex datasets to get start and stop dates of
exposure for all treatments for every subject ae record */
data aeexst;
merge ae trex trex2;
by usubjid;
run;
/* Create numeric dates using is8601 function and datetime20 format
for later use */
data aeexst;
set aeexst;
array dates(8) aestdtc aeendtc a b c paendt pbendt pcendt;
array numdates(8) aestdtc aeendtc a b c a b c;
do i = 1 to 8;
if dates(i) \^='' then
numdates(i) = input(dates(i), IS8601DT.);
end;
drop i;
format aestdtc aeendtc a b c a b c datetime20.;
run;
/* Create numeric dates using is8601 function and datetime20 format
for later use */
data dm;
set r.dm;
if rfstdtc ^='' then do;
rfstdt=input(rfstdtc, IS8601DT.);
rfendt=input(rfendtc, IS8601DT.);
end;
format rfstdt rfendt datetime20.;
run;
/* Merge dm data with rfstdt and rfendt variables with merged ae and
ex dataset from above */
data aed;
```

```
merge aeexst dm(keep=usubjid rfstdt rfendt);
by usubjid;
run;
/* Delete non ae records from the merged dataset */
proc sql;
delete from aed
where usubjid not in (select usubjid from r.ae);
quit;
/* Assign treatment groups to each AE record depending on when AE
occured - used ae start dates in calculation */
proc sal;
create table aed2 as select *,
case when aestdtc<=rfstdt then "Z"</pre>
      when ( aestdtc between a and a+48*60*60) or ( aestdtc between a
and a+48*60*60) then "A"
      when ( aestdtc between b and b+48*60*60) or ( aestdtc between b
and b+48*60*60) then "B"
     when (aestdtc between c and c+48*60*60)or(aestdtc between c
and c+48*60*60) then "C"
      else "" end as treatc
from aed;
quit;
options mprint mlogic symbolgen;
/* Macro to prepare final report dataset by creating horizontal and
vertical datasets and merging */
%macro aeprep;
%let trt1=%upcase(a);
%let trt2=%upcase(b);
%let trt3=%upcase(c);
%let trt4=%upcase(z);
%let trt5=overall;
%let str=%nrstr(put(count(*),3.0) ||compbl(" (N="||put(count(distinct
usubjid),3.0)||")"));
/* Vertical dataset "overall" to get overall numbers - no grouping */
proc sql ;
     create table &trt5 as
     (select "Any Adverse Event" as AE GRP, %unquote(&str) as Overall,
"1" as order from aed2)
union all
     (select distinct(propcase(aesev)) as AE GRP, %unquote(&str) as
Overall, "2" as _order from aed2
     group by 1)
union all
```

```
(select "Possible, Probably, or Definitely Related Adverse Event"
as AE GRP,
     %unquote(&str) as Overall, "3" as order from aed2
     where aerel in ("POSSIBLY", "PROBABLY", "DEFINITELY"))
union all
     (select "Adverse Event Leading to Study Discontinuation" as
AE GRP,
     %unquote(&str) as Overall, "4" as order from aed2
     where aeacnoth in ("WITHDRAWN FROM STUDY", "TREATMENT GIVEN AND
WITHDRAWN FROM STUDY"))
union all
     (select "Serious Adverse Event" as AE GRP, %unquote(&str) as
Overall, "5" as order from aed2
     where aeser="Y");
quit;
/* Vertical datasets "Pre-Study Drug", "A", "B", "C" - grouping by
treatment and pre-study */
%do i=1 %to 4;
proc sql ;
     create table &&trt&i as
     (select "Any Adverse Event" as AE GRP, %unquote(&str) as &&trt&i,
"1" as order from aed2
     where treatc="&&trt&i")
union all
     (select distinct(propcase(aesev)) as AE GRP, %unquote(&str) as
&&trt&i, "2" as order from aed2
     where treatc="&&trt&i"
     group by 1)
union all
     (select "Possible, Probably, or Definitely Related Adverse Event"
as AE GRP,
     %unquote(&str) as &&trt&i, "3" as order from aed2
     where treatc="&&trt&i"
     and aerel in ("POSSIBLY", "PROBABLY", "DEFINITELY"))
union all
     (select "Adverse Event Leading to Study Discontinuation" as
AE GRP,
     %unquote(&str) as &&trt&i, "4" as order from aed2
     where treatc="&&trt&i"
     and aeacnoth in ("WITHDRAWN FROM STUDY", "TREATMENT GIVEN AND
WITHDRAWN FROM STUDY"))
union all
     (select "Serious Adverse Event" as AE GRP, %unquote(&str) as
&&trt&i, "5" as order from aed2
     where treatc="&&trt&i"
     and aeser="Y");
quit;
%end;
/* Create row for dataset Z where proc sql did not create a row for
SEVERE ae category */
```

```
data z;
set z;
if n = 3 then do;
output;
ae grp="SEVERE";
order="2";
z="---";
end;
output;
run;
/* Massage data to include ae group labels and order variable values
%do i=1 %to 5;
data &&trt&i;
set &&trt&i;
if n =6 and ae grp='' then do;
ae grp="Adverse Event Leading to Study Discontinuation";
order="4";
end;
if n = 7 and ae grp='' then do;
ae grp="Serious Adverse Event";
order="5";
end;
if order="2" then do;
ae_grp=" "||ae_grp;
end;
proc sort data = &&trt&i;
by order;
run;
%end;
/* Create final report dataset by merging all vertical datasets
created above */
data final;
merge z a b c overall;
by order;
run;
proc datasets library=work;
save final;
run; quit;
%mend aeprep;
%aeprep; /* Macro execution step to create vertical datasets */
/* Create blank rows after each ae group category */
data final;
set final;
output;
```

```
by order;
if order^=1 and last. order;
array allchar [*] character ;
drop i;
do i=1 to dim(allchar); allchar{i}=' '; end;
output; /* Output blank observation */
run;
/* Format values per SAP specifications */
proc format;
     value $col 30. ' 0(N=0)'=' --- '
                           ' 0 (N= 0)'=' --- '
run;
/*Create macro variable for sysdate system macro variable in format
specified in shells */
data null;
today = put(date(), date9.);
call symput('dtnull',today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull, 6, 4);
/* Report procedure to export rtf file */
dm "odsresults; clear;";
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-1-1a_new.rtf" bodytitle style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=30
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black}
     columns (ae grp('Incidences (Number of Distinct Subjects)*
```

```
define ae grp /display " " style={protectspecialchars=off
cellwidth=30% just=l asis=on} flow;
     define z /display center "Pre-Study*Drug" format=$col.
style={protectspecialchars=off cellwidth=13% just=c};
     define a /display center "A* (N=&trta)" format=$col.
style={protectspecialchars=off cellwidth=13% just=c};
     define b /display center "B*(N=&trtb)" format=$col.
style={protectspecialchars=off cellwidth=13% just=c};
     define c /display center "C*(N=&trtc)" format=$col.
style={protectspecialchars=off cellwidth=13% just=c};
     define overall /display center "Overall* (N=&overall)"
format=$col. style={protectspecialchars=off cellwidth=17% just=c};
     define order /order=internal noprint;
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.3.1-1a
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Summary of All Adverse Events";
     title3 j=c "(Page 1 of 1)";
     *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Enrolled
                          Treatment A = Finely crushed 10 mg OTR;
     footnote j=1 "NOTE:
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote2 j=1 "Source: Listing 16.2.7-1a";
run; quit;
ods listing;
ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.3.1.1c
     Notes:
                                       Re-programming sample tables as
part of QC.
* /
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
/* Copy ex dataset to get treatment information */
data ex;
length treatc $1;
set r.ex(keep=studyid usubjid exseq exdostxt exstdtc exendtc epoch
visitnum);
if EXDOSTXT="FINELY CRUSHED 10 MG OTR" then treatc="A";
if EXDOSTXT="COARSELY CRUSHED 10 MG OTR" then treatc="B";
if EXDOSTXT="FINELY CRUSHED 10 MG OC" then treatc="C";
drop exdostxt;
run;
/* Get unique subject counts by treatment groups - Not used for this
report */
proc sql noprint;
select count(distinct usubjid) into :trta from ex where treatc="A";
select count(distinct usubjid) into :trtb from ex where treatc="B";
select count(distinct usubjid) into :trtc from ex where treatc="C";
select count(distinct usubjid) into :overall from ex where treatc in
("A", "B", "C");
quit;
/* Not used for this report */
%let trta=&trta;
%let trtb=&trtb;
%let trtc=&trtc;
%let overall=&overall;
/* Transpose ex dose start and end dates */
proc transpose data=ex out=trex(drop= name );
by usubjid;
id treatc ;
var exstdtc;
/* Transpose ex dose end dates */
proc transpose data=ex out=trex2(drop=_name__rename=(a=paendt b=pbendt
c=pcendt));
by usubjid;
id treatc ;
```

```
var exendtc;
run:
/* Add :00 to the ae start and end dates to make use of the is8601
function later on the variables */
data ae;
set r.ae;
aestdtc2=trim(trim(aestdtc)||trim(":00"));
aeendtc2=trim(trim(aeendtc)||trim(":00"));
drop aestdtc aeendtc;
rename aestdtc2=aestdtc aeendtc2=aeendtc;
run;
/* Merge ae and transposed ex datasets to get start and stop dates of
exposure for all treatments for every subject ae record */
data aeexst;
merge ae trex trex2;
by usubjid;
run;
/* Create numeric dates using is8601 function and datetime20 format
for later use */
data aeexst;
set aeexst;
array dates(8) aestdtc aeendtc a b c paendt pbendt pcendt;
array numdates(8) aestdtc aeendtc a b c a b c;
do i = 1 to 8;
if dates(i) \(^=''\) then
numdates(i) = input(dates(i), IS8601DT.);
end;
drop i;
format aestdtc aeendtc a b c a b c datetime20.;
/* Create numeric dates using is8601 function and datetime20 format
for later use */
data dm;
set r.dm;
if rfstdtc ^='' then do;
rfstdt=input(rfstdtc, IS8601DT.);
rfendt=input(rfendtc, IS8601DT.);
format rfstdt rfendt datetime20.;
run;
/* Merge dm data with rfstdt and rfendt variables with merged ae and
ex dataset from above */
data aed;
merge aeexst dm(keep=usubjid rfstdt rfendt);
by usubjid;
run;
```

```
/* Delete non ae records from the merged dataset */
proc sql;
delete from aed
where usubjid not in (select usubjid from r.ae);
quit;
/* Assign treatment groups to each AE record depending on when AE
occured - used ae start dates in calculation */
proc sql;
create table aed2 as select *,
case when aestdtc<=rfstdt then "Z"</pre>
      when ( aestdtc between a and a+48*60*60) or ( aestdtc between a
and _a+48*60*60) then "A"
     when (aestdtc between b and b+48*60*60)or(aestdtc between b
and b+48*60*60) then "B"
      when (aestdtc between c and c+48*60*60) or (aestdtc between c
and c+48*60*60) then "C"
      else "" end as treatc
from aed;
quit;
/* Create vertical tables left to right per shell specification */
/* Step to create N subject counts by treatment group */
%let str=%nrstr(select distinct treatc, count(distinct usubjid) as
ncount from ex where treatc in);
proc sql noprint;
create table ncount as
%unquote(&str)("A")
union all
%unquote(&str)("B")
union all
%unquote(&str)("C")
union all
select "Total" as treatc, count(distinct usubjid) as ncount from ex
where treatc in ("A", "B", "C");
quit;
/* Step to create TEAE counts by treatment groups */
proc sql;
create table nteae as select distinct treatc, count (aesev) as nteae
from aed2
where aestdtc>rfstdt
group by 1
union all
select "Total" as treatc, count(aesev) as nteae from aed2
where aestdtc>rfstdt;
quit;
/* Macro to prepare datasets mild, moderate, severe - ae severity
groups by treatment groups */
```

```
%macro prep;
%let sev1=MILD;
%let sev2=MODERATE;
%let sev3=SEVERE;
%do i=1 %to 3;
proc sql;
create table &&sev&i as select distinct treatc, count(*) as &&sev&i
from aed2
where aesev="&&sev&i"
and aestdtc>rfstdt
group by treatc
union all
select "Total" as treatc, count(*) as &&sev&i from aed2
where aesev="&&sev&i"
and aestdtc>rfstdt;
quit;
%end;
%mend prep;
/* Macro execution step */
% prep;
/* Second prep macro for aerel sub categories and corresponding
vertical datasets */
%macro prep2;
%let aerel1=NOT RELATED;
%let aerel2=UNLIKELY;
%let aerel3=POSSIBLY;
%let aerel4=PROBABLY;
%let aerel5=DEFINITELY;
%do i=1 %to 5;
proc sql;
create table &&aerel&i as select distinct treatc, count(*) as
&&aerel&i from aed2
where aerel like ("&&aerel&i")
and aestdtc>rfstdt
group by treatc
union all
select "Total" as treatc, count(*) as &&aerel&i from aed2
where aerel like ("&&aerel&i")
and aestdtc>rfstdt;
quit;
%end;
%mend prep2;
/* Macro execution */
% prep2;
```

```
/\star Create final dataset for report by merging all vertical datasets by
treatment groups and total row */
data final;
merge ncount nteae mild moderate severe not related unlikely possibly
probably definitely;
by treatc;
run;
/* Massage final dataset - assign value 0 where value is missing */
data final;
set final;
by treatc;
array allnum [*] numeric ;
do i=1 to dim(allnum);
if allnum{i}=. then allnum{i}=0;
end;
drop i;
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
dm "odsresults; clear;";
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-1-1c.rtf" bodytitle style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=30
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
cellheight=.25in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
```

```
columns treatc ncount nteae
                             ' mild moderate severe)
('Severity*
            ('Relationship to
Drug*
                                                   ' not related
unlikely possibly probably definitely);
     define treatc / display "Treatment" style={cellwidth=10% just=c};
     define ncount / display "N" style={cellwidth=8% just=c};
     define nteae / display "Total Number of*Treatment-Emergent AEs"
style={cellwidth=17% just=c};
     define mild / display "Mild" style={cellwidth=8% just=c};
     define moderate / display "Moderate" style={cellwidth=8% just=c};
     define severe / display "Severe" style={cellwidth=8% just=c};
     define not related / display "NR" style={cellwidth=8% just=c};
     define unlikely / display "Unlikely" style={cellwidth=8% just=c};
     define possibly / display "Possibly" style={cellwidth=8% just=c};
     define probably / display "Probably" style={cellwidth=8% just=c};
     define definitely / display "Definitely" style={cellwidth=8%
just=c};
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.3.1-1c
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Summary of Treatment-Emergent Adverse Events Based on
Severity and Relationship to Drug";
     title3 j=c "(Page 1 of 1)";
     *title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety
";
     footnote1 j=1 "NOTE: N=Sample Size, NR=Not Related";
     footnote2 j=1 '09'x "Treatment A = Finely crushed 10 mg OTR;
Treatment B = Coarsely crushed 10 mg OTR; Treatment C = Finely crushed
10 mg OC.";
     footnote3;
     footnote4 j=1 "Source: Listing 16.2.7-2";
run; quit;
 ods listing;
 ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.3.5-1a
     Notes:
                                       Re-programming sample tables as
part of QC.
*/
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
%macro chqbase;
/* Change Information here for other lab final datasets */
%let lab1=ALBM;
%let lab2=GLUC;
%let lab3=PROTT;
/** ---- **/
%DO I=1 %TO 3;
data &&lab&i;
set r.lb (where = (lbcat="CHEMISTRY" and lbtestcd="&&lab&i"));
           by usubjid visitnum;
run;
proc univariate data=&&lab&i(where=(lbtptnum=1)) noprint round=0;
var lbstresn;
output out=scr_&&lab&i n=a N mean=b mean std=c_sd min=d min
median=e median max=f max;
run;
proc transpose data=scr &&lab&i out=scr &&lab&i;
var _all_;
run;
data scr &&lab&i;
set scr &&lab&i;
lbtest="&&lab&i";
period="a Screening";
run;
proc sort data=scr &&lab&i;
by name;
run;
proc univariate data=&&lab&i(where=(lbtptnum=2)) noprint round=0;
```

```
var lbstresn;
output out=base &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=base &&lab&i out=base &&lab&i;
var _all_;
run;
data base &&lab&i;
set base &&lab&i;
lbtest="&&lab&i";
period="b Baseline";
run;
proc sort data=base &&lab&i;
by name;
run;
proc univariate data=&&lab&i(where=(lbtptnum>2 and
visitnum=int(visitnum))) noprint round=0;
var lbstresn;
output out=c EOS &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=c EOS &&lab&i out=c EOS &&lab&i;
var all;
run;
data c EOS &&lab&i;
set c EOS &&lab&i;
lbtest="&&lab&i";
period="c_EOS";
run;
proc sort data=c EOS &&lab&i;
by name;
run;
data chgbl &&lab&i;
     do until (last.usubjid);
           set &&lab&i (where = (lbstresn^= . and
visitnum=int(visitnum)));
           by usubjid visitnum;
           if visitnum <= 2 then bl=lbstresn;</pre>
                 else do;
                      chgbl = lbstresn - bl;
                      output;
                 end;
     end;
run;
```

```
proc univariate data=chgbl &&lab&i noprint round=0;
var chgbl;
output out=chqbl &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=chgbl &&lab&i out=chgbl &&lab&i;
var _all_;
run;
data chgbl &&lab&i;
set chqbl &&lab&i;
lbtest="&&lab&i";
period="d CFB";
run;
proc sort data=chgbl &&lab&i;
by name;
run;
data lab &&lab&i;
set scr &&lab&i base &&lab&i c EOS &&lab&i chgbl &&lab&i;
/* Report Decimal places that correspond to the particular lab value
/* Report SD to one decimal place more than mean */
/* Labs Specific */
%if lab &&lab&i^="lab gluc" %then %do;
data lab &&lab&i;
length col2 $20;
set lab &&lab&i;
if _name_ not in ("b_mean","c_sd") then col2=put(col1,3.);
if name = "b mean" then col2=put(round(col1,1.0),3.);
if name = "c sd" then col2=put(round(col1,.1),5.1);
run;
%end;
%if lab &&lab&i="lab gluc" %then %do;
data lab gluc;
length col2 $20;
set lab gluc;
if name ="N" then col2=put(col1,3.);
if name = "c sd" then col2=put(round(col1,.001),7.3);
if name in ("e median", "f max", "d min", "b mean") then
col2=put(round(col1,.001),7.2);
run;
%end;
%end;
```

```
%mend chgbase;
%chgbase;
/** Change Information here for other lab final datasets **/
data final;
set lab albm lab gluc lab prott;
if lbtest="ALBM" then order1=1; /* Labs Specific */
if lbtest="GLUC" then order1=2; /* Labs Specific */
if lbtest="PROT" then order1=3; /* Labs Specific */
if period="a Screening" then order2=1;
if period="b Baseline" then order2=2;
if period="c EOS" then order2=3;
if period="d CFB" then order2=4;
run;
proc format;
                            = 'N'
     value $reportf 'a N'
                         'b mean'
                                      = 'MEAN'
                         __sα'
'd_min'
'e_~
                                    = 'SD'
= 'MIN'
                         'e median' = 'MEDIAN'
                         'f max' = 'MAX'
                         'a Screening'= 'Screening'
                         'b Baseline' = 'Baseline, Day -1'
                                 = 'End of Study'
= 'Change from Baseline'
                         'c EOS'
                         'd CFB'
                         'ALBM'
                                     = 'Albumin (q/L)'
     /* Labs Specific */
                         'GLUC' = 'Glucose (mmol/L)'
                                                                /*
Labs Specific */
                         'PROT' = 'Total Protein (g/L)'; /*
Labs Specific */
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull',today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
```

```
/* Proc Report Step */
/* Make changes here for additional lab reports */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14.
3.5-la.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font_face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.15in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns order1 lbtest period name col2 col1;
     define order1 / order order=internal noprint ;
     *define order2 / order order=internal noprint;
     define lbtest / group "Laboratory Parameter" format=$reportf.
style={cellwidth=20% just=c};
     define period / group order=internal "Timepoint" format=$reportf.
style={cellwidth=25% just=c};
     define name / display "Statistic" format=$reportf.
style={cellwidth=25% just=c};
     define col2 / display "Result" center format=$reportf.
style={cellwidth=15% just=dec};
     define col1 / center style={cellwidth=14% just=c
foreground=white};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     compute after order1;
     line @1 ' ';
     endcomp;
     compute after period;
```

```
line @1 ' ';
     endcomp;
     title1 j=1 "OTR1021
                Table 14.3.5-1a
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Descriptive Statistics and Change from b Baseline of
Chemistry Laboratory Data: Metabolic Substrates";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: Change from b Baseline is the change from
b Baseline, Day -1 compared to End of Study.";
     footnote2 j=1 '09'x "Lab units are presented as SI units.";
     footnote3;
     footnote4 j=1 "Source: Listing 16.2.8-2a";
 run; quit;
 ods listing;
ods rtf close;
/*
                                       OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.3.5-1b
                                       Re-programming sample tables as
     Notes:
part of QC.
*/
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
options mprint mlogic symbolgen;
/* Make changes here for additional labs */
%let str=%nrstr("ALBM", "GLUC", "PROTT");
%let str2=%nrstr(ALBM,GLUC,PROTT);
%let str3=%nrstr(ALBM||GLUC||PROTT);
```

```
/*----Generic re-usable code starts here-----
____*/
proc sql;
create table lbsub as select usubjid, lbdtc, lbtpt, lbcat, lbtestcd,
lbstresn, lbnrind,
case when substr(lbnrind,1,1)^='' then
trim(put(lbstresn,best12.)||' '||trim(substr(lbnrind,1,1))||', N')
else ' ' end as lbstresn2 from r.lb
where lbtestcd in (%unquote(&str));
quit;
proc transpose data=lbsub out=lbsub;
by usubjid lbdtc lbtpt;
id lbtestcd;
var lbstresn2;
run;
data lbsub;
length sex $10;
merge lbsub(in=a)
        r.sc(in=b keep=usubjid scorres where=(scorres like ("RN%")))
        r.dm(in=c keep=usubjid age sex race);
by usubjid;
if a;
if sex='F' then sex="Female";
if sex='M' then sex="Male";
data lbsub;
length rsub $25;
length asr $45;
length lbdtc2 8;
set lbsub;
rsub=trim(trim(scorres)||'/'||substr(usubjid,15,5));
asr=propcase(trim(trim(age)||'/'||trim(sex)||'/'||trim(race)));
lbdtc2=input(substr(lbdtc,1,10),yymmdd10.);
format lbdtc2 date9.;
run;
proc sql;
create table final as select rsub, asr, propcase(lbtpt) as lbtpt,
lbdtc2, substr(lbdtc, 12, 5) as time, %unquote(&str2)
from lbsub
where index((%unquote(&str3)),',') ^=0;
quit;
/*----End Generic code-------
----*/
/* Make changes here for additional lab reports */
```

```
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-1b.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     1s=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.4in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns rsub asr lbtpt lbdtc2 time albm gluc prott;
     define rsub / display order order=internal center
"Randomization*No/Subject No" style={cellwidth=15% just=1};
     define asr / display center "Age/Gender/*Race" flow
style={cellwidth=12.5% just=c};
     define lbtpt / display center "Timepoint" style={cellwidth=13.5%
     define lbdtc2 / display "Date" style={cellwidth=12.5% just=c};
     define time / display "Actual*Time" center style={cellwidth=12.5%
just=c};
     define albm / display center "Albumin*(g/L)" style={cellwidth=11%
just=c};
     define gluc / display center "Glucose* (mmol/L)"
style={cellwidth=11% just=c};
     define prott / display center "Total Protein*(g/L)"
style={cellwidth=11% just=c};
     compute before;
     line @1 ' ';
```

```
endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-1b
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Subjects with Abnormal Chemistry Laboratory Data
Based on the Reference Ranges: Metabolic Substrates";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: H = laboratory value above the reference
laboratory range";
     footnote2 j=1 '09'x "L = laboratory value below the reference
laboratory range";
     footnote3 j=1 '09'x "A = laboratory value abnormal";
     footnote4 j=1 '09'x "C = laboratory value deemed clinically
significant by the investigator";
     footnote5 j=1 '09'x "N = laboratory value deemed not clinically
significant by the investigator.";
     footnote6 j=l '09'x "Values out of the normal range are
flagged.";
     footnote7 j=1 '09'x "Lab units are presented as SI units.";
     footnote8 j=1 '09'x "Source: Listing 16.2.8-2a";
     footnote9 j=1 '09'x "Reference Range Source: Listing 16.2.8-1";
run; quit;
ods listing;
 ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
     Date:
                                       06-18-**
     Study:
                                       OTR1021
                                       14.3.5-1c
     Table:
     Notes:
                                       Re-programming sample tables as
part of QC.
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
%macro chqbase;
/* Change Information here for other lab final datasets */
%let lab1=ALP;
%let lab2=AST;
%let lab3=ALT;
%let lab4=LDH;
%let lab5=BILID;
%let lab6=BILIT;
/** ---- **/
%do i=1 %to 6;
data &&lab&i;
set r.lb (where = (lbcat="CHEMISTRY" and lbtestcd="&&lab&i"));
           by usubjid visitnum;
proc univariate data=&&lab&i(where=(lbtptnum=1)) noprint round=0;
var lbstresn;
output out=scr &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=scr &&lab&i out=scr &&lab&i;
var _all_;
run;
data scr &&lab&i;
length lbtest $15;
set scr &&lab&i;
lbtest="&&lab&i";
period="a Screening";
run;
```

```
proc sort data=scr &&lab&i;
by _name_;
run;
proc univariate data=&&lab&i(where=(lbtptnum=2)) noprint round=0;
var lbstresn;
output out=base &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=base &&lab&i out=base &&lab&i;
var _all_;
run;
data base &&lab&i;
set base &&lab&i;
lbtest="&&lab&i";
period="b Baseline";
run;
proc sort data=base &&lab&i;
by name;
run;
proc univariate data=&&lab&i(where=(lbtptnum>2 and
visitnum=int(visitnum))) noprint round=0;
var lbstresn;
output out=c EOS &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=c EOS &&lab&i out=c EOS &&lab&i;
var _all_;
run;
data c EOS &&lab&i;
set c EOS &&lab&i;
lbtest="&&lab&i";
period="c EOS";
run;
proc sort data=c EOS &&lab&i;
by name;
run;
data chqbl &&lab&i;
     do until (last.usubjid);
           set &&lab&i (where = (lbstresn^= . and
visitnum=int(visitnum)));
           by usubjid visitnum;
           if visitnum <= 2 then bl=lbstresn;</pre>
```

```
else do;
                      chqbl = lbstresn - bl;
                      output;
                end;
     end;
run;
proc univariate data=chgbl &&lab&i noprint round=0;
var chqbl;
output out=chgbl &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=chgbl &&lab&i out=chgbl &&lab&i;
var all;
run;
data chgbl &&lab&i;
set chgbl &&lab&i;
lbtest="&&lab&i";
period="d CFB";
run;
proc sort data=chgbl &&lab&i;
by name;
run;
data lab &&lab&i;
length col2 $20;
set scr &&lab&i base &&lab&i c EOS &&lab&i chgbl &&lab&i;
if lbtest^="BILID" then do;
     if name ="a N" then col2=put(col1,8.);
     else if _name = "b_mean" then col2=put(round(col1,1.0),8.0);
     else if _name_= "c_sd" then col2=put(round(col1,.1),8.1);
     else if name not in ("a N", "b_mean", "c_sd") then
col2=put(round(col1,1.0),8.);
end;
if lbtest="BILID" then do;
     if name ="a N" then col2=put(col1,8.);
     else if _name ="c_sd" then col2=put(round(col1,.001),8.3);
     else if name not in ("a N", "c sd") then
col2=put(round(col1,.01),8.2);
end;
run;
%end;
data final;
set lab alp lab alt lab ast lab bilid lab bilit lab ldh;
run;
proc datasets lib=work nolist;
```

```
save final;
run; quit;
%mend chgbase;
%chgbase;
proc format;
     value portf 'a N' = 'N'
                          'b mean'
                                          = 'MEAN'
                          'c sd'
                                      = 'SD'
                          'd min' = 'MIN'
                          'e median' = 'MEDIAN'
                          'f max' = 'MAX'
                          'a Screening'= 'Screening'
                          'b Baseline' = 'Baseline, Day -1'
                          'c_EOS' = 'End of Study'
'd_CFB' = 'Change from Baseline'
                          'ALP'
                                      = 'Alkaline Phosphatase (U/L)'
                                      = 'AST (U/L)'
                          'AST'
                                      = 'ALT (U/L)'
                          'ALT'
                          'LDH'
                                      = 'LDH (U/L)'
                          'BILID' = 'Direct Bilirubin (\(\mu\text{mol/L}\)'
'BILIT' = 'Total Bilirubin (\(\mu\text{mol/L}\)';
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
/* Make changes here for additional lab reports */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-1c.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
```

```
ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.16in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns lbtest period name col2 col1;
     *define order1 / order order=internal noprint;
     *define order2 / order order=internal noprint ;
     define lbtest / group "Laboratory Parameter" format=$reportf.
style={cellwidth=20% just=c};
     define period / group order=internal "Timepoint" format=$reportf.
style={cellwidth=25% just=c};
     define name / display "Statistic" format=$reportf.
style={cellwidth=25% just=c};
     define col2 / display "Result" center format=$reportf.
style={cellwidth=15% just=dec};
     define col1 / center style={cellwidth=14% just=c
foreground=white};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     compute after period;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-1c
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Descriptive Statistics and Change from Baseline of
Chemistry Laboratory Data: Hepatic Status Liver Function Tests";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: Change from Baseline is the change from
Baseline, Day -1 compared to End of Study.";
     footnote2 j=1 '09'x "Lab units are presented in SI units.";
     footnote3;
     footnote4 j=1 "Source: Listing 16.2.8-2b";
```

```
run; quit;
 ods listing;
 ods rtf close;
/*
     Title:
                                      OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                      06-18-**
     Date:
     Study:
                                      OTR1021
     Table:
                                      14.3.5-1d
     Notes:
                                      Re-programming sample tables as
part of QC.
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
options mprint mlogic symbolgen;
/* Make changes here for additional labs */
%let str=%nrstr("ALP","AST","ALT","LDH","BILID","BILIT");
%let str2=%nrstr(ALP, AST, ALT, LDH, BILID, BILIT);
%let str3=%nrstr(ALP||AST||ALT||LDH||BILID||BILIT);
/*----Generic re-usable code starts here-----
----*/
proc sql;
create table lbsub as select usubjid, lbdtc, lbtpt, lbcat, lbtestcd,
lbstresn, lbnrind,
case when substr(lbnrind,1,1)^='' then
trim(put(lbstresn,best12.)||' '||trim(substr(lbnrind,1,1))||', N')
else ' ' end as lbstresn2 from r.lb
where lbtestcd in (%unquote(&str));
quit;
proc transpose data=lbsub out=lbsub;
by usubjid lbdtc lbtpt;
id lbtestcd;
var lbstresn2;
run;
data lbsub;
length sex $10;
```

```
merge lbsub(in=a)
        r.sc(in=b keep=usubjid scorres where=(scorres like ("RN%")))
        r.dm(in=c keep=usubjid age sex race);
by usubjid;
if a;
if sex='F' then sex="Female";
if sex='M' then sex="Male";
run;
data lbsub;
length rsub $25;
length asr $45;
length lbdtc2 8;
set lbsub;
rsub=trim(trim(scorres)||'/'||substr(usubjid, 15, 5));
asr=propcase(trim(trim(age)||'/'||trim(sex)||'/'||trim(race)));
lbdtc2=input(substr(lbdtc,1,10),yymmdd10.);
format lbdtc2 date9.;
run;
proc sql;
create table final as select rsub, asr, propcase(lbtpt) as lbtpt,
lbdtc2, substr(lbdtc, 12, 5) as time, %unquote(&str2)
from lbsub
where index((%unquote(&str3)),',') ^=0;
quit;
/*----End Generic code------
____*/
/* Make changes here for additional lab reports */
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nac103\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-1d.rtf" style=Styles.custom;
proc report
     data = final
     nowd
```

```
ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.5in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns rsub asr lbtpt lbdtc2 time ALP AST ALT LDH BILID BILIT;
     define rsub / display group order order=internal center
"Randomization*No/Subject No" style={cellwidth=11% just=1};
     define asr / display group center "Age/Gender/*Race" flow
                                 style={cellwidth=11% just=c};
     define lbtpt / display center "Timepoint"
                                       style={cellwidth=11% just=c};
     define lbdtc2 / display "Date"
                                            style={cellwidth=8%
just=c};
     define time / display "Actual*Time" center
                                      style={cellwidth=5% just=c};
     define ALP / display center "Alkaline*Phosphatase* (U/L)"
                                 style={cellwidth=9% just=c};
     define AST / display center "Aspartate*Transferase*(U/L)"
                                 style={cellwidth=9% just=c};
     define ALT / display center "Alanine*Transferase* (U/L)"
                                 style={cellwidth=9% just=c};
     define LDH / display center "Lactic Dehydrogenase* (U/L)"
                                 style={cellwidth=11% just=c};
     define BILID / display center "Bilirubin Direct*(µmol/L)"
                                 style={cellwidth=7.5% just=c};
     define BILIT / display center "Total Bilirubin*(\u03c4mol/L)"
                                 style={cellwidth=7.5% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-1d
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Subjects with Abnormal Chemistry Laboratory Data
Based on the Reference Ranges: Hepatic Status Liver Function Tests";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
```

```
footnote1 j=1 "NOTE: H = laboratory value above the reference
laboratory range";
     footnote2 j=1 '09'x "L = laboratory value below the reference
laboratory range";
     footnote3 j=1 '09'x "A = laboratory value abnormal";
     footnote4 j=1 '09'x "C = laboratory value deemed clinically
significant by the investigator";
     footnote5 j=1 '09'x "N = laboratory value deemed not clinically
significant by the investigator.";
     footnote6 j=l '09'x "Values out of the normal range are
flagged.";
     footnote7 j=1 '09'x "Lab units are presented as SI units.";
     footnote8 j=1 '09'x "Source: Listing 16.2.8-2b";
     footnote9 j=1 '09'x "Reference Range Source: Listing 16.2.8-1";
run; quit;
ods listing;
ods rtf close;
/*
     Title:
                                      OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                      06-18-**
     Date:
     Study:
                                      OTR1021
     Table:
                                      14.3.5-2a
     Notes:
                                      Re-programming sample tables as
part of QC.
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
%macro chqbase;
/* Change Information here for other lab final datasets */
%let lab1=HCT;
%let lab2=HGB;
%let lab3=RBC;
%let lab4=PLT;
%let lab5=WBC;
/** ---- **/
```

```
%do i=1 %to 5;
data &&lab&i;
set r.lb (where = (lbcat="HEMATOLOGY" and lbtestcd="&&lab&i"));
           by usubjid visitnum;
run;
proc univariate data=&&lab&i(where=(lbtptnum=1)) noprint round=0;
var lbstresn;
output out=scr &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=scr &&lab&i out=scr &&lab&i;
var _all_;
run;
data scr &&lab&i;
length lbtest $15;
set scr &&lab&i;
lbtest="&&lab&i";
period="a Screening";
run;
proc sort data=scr &&lab&i;
by name;
run;
proc univariate data=&&lab&i(where=(lbtptnum=2)) noprint round=0;
var lbstresn;
output out=base &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=base &&lab&i out=base &&lab&i;
var _all_;
run;
data base &&lab&i;
set base &&lab&i;
lbtest="&&lab&i";
period="b Baseline";
run;
proc sort data=base &&lab&i;
by name;
run;
proc univariate data=&&lab&i(where=(lbtptnum>2 and
visitnum=int(visitnum))) noprint round=0;
var lbstresn;
```

```
output out=c EOS &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=c EOS &&lab&i out=c EOS &&lab&i;
var all;
run;
data c EOS &&lab&i;
set c EOS &&lab&i;
lbtest="&&lab&i";
period="c EOS";
run;
proc sort data=c EOS &&lab&i;
by name;
run;
data chgbl &&lab&i;
     do until (last.usubjid);
           set &&lab&i (where = (lbstresn^= . and
visitnum=int(visitnum)));
           by usubjid visitnum;
           if visitnum <= 2 then bl=lbstresn;
                 else do;
                      chgbl = lbstresn - bl;
                      output;
                 end;
     end;
run;
proc univariate data=chgbl &&lab&i noprint round=0;
var chgbl;
output out=chgbl &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=chqbl &&lab&i out=chqbl &&lab&i;
var _all_;
run;
data chqbl &&lab&i;
set chgbl &&lab&i;
lbtest="&&lab&i";
period="d CFB";
run;
proc sort data=chgbl &&lab&i;
by _name_;
run;
```

```
data lab &&lab&i;
length col2 $20;
set scr &&lab&i base &&lab&i c EOS &&lab&i chgbl &&lab&i;
if lbtest="WBC" then do;
     if name ="a N" then col2=put(col1,8.);
     else if name = "c sd" then col2=put(round(col1,.1),8.2);
     else if _name_ not in ("a_N","c_sd") then
col2=put(round(col1,1.0),8.1);
end;
else if lbtest in ("HCT", "RBC") then do;
     if name ="a N" then col2=put(col1,8.);
     else if name ="c sd" then col2=put(round(col1,.001),8.3);
     else if _name_ not in ("a_N","c_sd") then
col2=put(round(col1,.01),8.2);
end;
else if lbtest in ("PLT", "HGB") then do;
     if name = "a N" then col2=put(col1,8.);
     else if _name_="c_sd" then col2=put(round(col1,.001),8.1);
     else if name not in ("a N", "c sd") then
col2=put(round(col1,.01),8.);
end;
run;
%end;
data final;
set lab hct lab hgb lab rbc lab plt lab wbc;
run;
proc datasets lib=work nolist;
save final;
run; quit;
%mend chgbase;
%chqbase;
proc format;
     value $reportf 'a N'
                            = 'N'
                         'b mean'
                                         = 'MEAN'
                         'c sd'
                                     = 'SD'
                         'd min'
                                     = 'MIN'
                         'e median' = 'MEDIAN'
                         'f max'
                                    = 'MAX'
                         'a Screening'= 'Screening'
                         'b Baseline' = 'Baseline, Day -1'
                         'c EOS'
                                   = 'End of Study'
                         'd CFB'
                                     = 'Change from Baseline'
                                     = 'Hematocrit (fraction)'
                         'HCT'
                                    = 'Hemoglobin (g/L)'
= 'RBC (10^12/L)'
                         'HGB'
                         'RBC'
                                  = 'Platelet Count (10^9/L)'
                         'PLT'
```

```
= 'WBC (10^9/L)'
                         'WBC'
                         '0.00'
                                      = '0':
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
/* Make changes here for additional lab reports */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-2a.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report) = {font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.16in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns lbtest period name col2 col1;
     *define order1 / order order=internal noprint;
     *define order2 / order order=internal noprint ;
     define lbtest / group "Laboratory Parameter" format=$reportf.
style={cellwidth=25% just=c};
     define period / group order=internal "Timepoint" format=$reportf.
style={cellwidth=27.5% just=c};
     define name / display "Statistic" format=$reportf.
style={cellwidth=27.5% just=c};
```

```
define col2 / display "Result" center format=$reportf.
style={cellwidth=11% just=dec};
     define col1 / center style={cellwidth=8% just=c
foreground=white};
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
     endcomp;
     compute after period;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-2a
DATE:&dtnull2 TIME:&systime";
     title2 j=c "Descriptive Statistics and Change from Baseline of
Hematology Laboratory Data: CBC";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: Change from Baseline is the change from
Baseline, Day -1 compared to End of Study.";
     footnote2 j=1 '09'x "Lab units are presented in SI units.";
     footnote3;
     footnote4 j=1 "Source: Listing 16.2.8-3a";
run; quit;
ods listing;
ods rtf close;
/*
                                       OTR1021 Data Review
     Title:
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
     Study:
                                       OTR1021
     Table:
                                       14.3.5-2b
     Notes:
                                       Re-programming sample tables as
part of QC.
*/
```

```
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run:
options nolabel;
options mprint mlogic symbolgen;
/* Make changes here for additional labs */
%let str=%nrstr("HCT","HGB","RBC","PLT","WBC");
%let str2=%nrstr(HCT, HGB, RBC, PLT, WBC);
%let str3=%nrstr(HCT||HGB||RBC||PLT||WBC);
/*----Generic re-usable code starts here-----
----*/
proc sql;
create table lbsub as select usubjid, lbdtc, lbtpt, lbcat, lbtestcd,
lbstresn, lbnrind,
case when substr(lbnrind,1,1)^='' then
trim(put(lbstresn,best12.)||' '||trim(substr(lbnrind,1,1))||', N')
else ' ' end as lbstresn2 from r.lb
where lbtestcd in (%unquote(&str));
quit;
proc transpose data=lbsub out=lbsub;
by usubjid lbdtc lbtpt;
id lbtestcd;
var lbstresn2:
run;
data lbsub;
length sex $10;
merge lbsub(in=a)
        r.sc(in=b keep=usubjid scorres where=(scorres like ("RN%")))
        r.dm(in=c keep=usubjid age sex race);
by usubjid;
if a;
if sex='F' then sex="Female";
if sex='M' then sex="Male";
run;
data lbsub;
length rsub $25;
length asr $45;
length lbdtc2 8;
set lbsub;
rsub=trim(trim(scorres)||'/'||substr(usubjid, 15, 5));
asr=propcase(trim(trim(age)||'/'||trim(sex)||'/'||trim(race)));
lbdtc2=input(substr(lbdtc,1,10),yymmdd10.);
format lbdtc2 date9.;
run;
```

```
proc sql;
create table final as select rsub, asr, propcase(lbtpt) as lbtpt,
lbdtc2, substr(lbdtc, 12, 5) as time, %unquote(&str2)
from lbsub
where index((%unquote(&str3)),',') ^=0;
quit;
/*----End Generic code-----
____*/
/* Make changes here for additional lab reports */
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-2b.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.5in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns rsub asr lbtpt lbdtc2 time HCT HGB RBC PLT WBC;
     define rsub / display group order order=internal center
"Randomization*No/Subject No" style={cellwidth=12% just=1};
     define asr / display group center "Age/Gender/*Race" flow
                                style={cellwidth=12% just=c};
     define lbtpt / display center "Time-point"
                                     style={cellwidth=12% just=c};
```

```
define lbdtc2 / display "Date"
                                            style={cellwidth=10%
just=c};
     define time / display "Actual*Time" center
                                      style={cellwidth=9% just=c};
     define HCT / display center "Hematocrit*(fraction)"
                                 style={cellwidth=9% just=c};
     define HGB / display center "Hemoglobin*(g/L)"
                                      style={cellwidth=9% just=c};
     define RBC / display center "RBC*(10^12/L)"
                                      style={cellwidth=9% just=c};
     define PLT / display center "Platelet*Count*(10^9/L)"
                                 style={cellwidth=8.5% just=c};
     define WBC / display center "WBC*(10^9/L)"
                                      style={cellwidth=8.5% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-2b
DATE:&dtnull2 TIME:&systime";
     title2 j=c "Subjects with Abnormal Hematology Laboratory Data
Based on the Reference Ranges: CBC";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: H = laboratory value above the reference
laboratory range";
     footnote2 j=1 '09'x "L = laboratory value below the reference
laboratory range";
     footnote3 j=1 '09'x "A = laboratory value abnormal";
     footnote4 j=1 '09'x "C = laboratory value deemed clinically
significant by the investigator";
     footnote5 j=1 '09'x "N = laboratory value deemed not clinically
significant by the investigator.";
     footnote6 j=l '09'x "Values out of the normal range are
flagged.";
     footnote7 j=1 '09'x "Lab units are presented as SI units.";
     footnote8 j=1 '09'x "Source: Listing 16.2.8-3a";
     footnote9 j=1 '09'x "Reference Range Source: Listing 16.2.8-1";
run; quit;
ods listing;
 ods rtf close;
```

```
/*
                                       OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
                                       OTR1021
     Study:
     Table:
                                       14.3.5-2c
     Notes:
                                       Re-programming sample tables as
part of QC.
* /
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
%macro chgbase;
/* Change Information here for other lab final datasets */
%let lab1=BASOAB;
%let lab2=BASO;
%let lab3=EOSINAB;
%let lab4=EOSIN;
%let lab5=LYMPHAB;
%let lab6=LYMPH;
%let lab7=MONOAB;
%let lab8=MONO;
%let lab9=NEUTAB;
%let lab10=NEUT;
/** ---- **/
%do i=1 %to 10;
data &&lab&i;
set r.lb (where = (lbcat="HEMATOLOGY" and lbtestcd="&&lab&i"));
           by usubjid visitnum;
proc univariate data=&&lab&i(where=(lbtptnum=1)) noprint round=0;
var lbstresn;
output out=scr &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=scr &&lab&i out=scr &&lab&i;
```

```
var _all_;
run;
data scr &&lab&i;
length lbtest $15;
set scr &&lab&i;
lbtest="&&lab&i";
period="a Screening";
run;
proc sort data=scr &&lab&i;
by _name ;
run;
proc univariate data=&&lab&i(where=(lbtptnum=2)) noprint round=0;
var lbstresn;
output out=base &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=base &&lab&i out=base &&lab&i;
var all;
run;
data base &&lab&i;
set base &&lab&i;
lbtest="&&lab&i";
period="b Baseline";
run;
proc sort data=base &&lab&i;
by _name_;
run;
proc univariate data=&&lab&i(where=(lbtptnum>2 and
visitnum=int(visitnum))) noprint round=0;
var lbstresn;
output out=c EOS &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=c EOS &&lab&i out=c EOS &&lab&i;
var all;
run;
data c EOS &&lab&i;
set c EOS &&lab&i;
lbtest="&&lab&i";
period="c_EOS";
run;
proc sort data=c EOS &&lab&i;
```

```
by name;
run;
data chqbl &&lab&i;
     do until (last.usubjid);
           set &&lab&i (where = (lbstresn^= . and
visitnum=int(visitnum)));
           by usubjid visitnum;
           if visitnum = 2 then bl=lbstresn;
                else do;
                      chqbl = lbstresn - bl;
                      output;
                end;
     end;
run;
proc univariate data=chgbl &&lab&i noprint round=0;
var chgbl;
output out=chgbl &&lab&i n=a N mean=b mean std=c sd min=d min
median=e median max=f max;
run;
proc transpose data=chgbl &&lab&i out=chgbl &&lab&i;
var _all_;
run;
data chqbl &&lab&i;
set chgbl &&lab&i;
lbtest="&&lab&i";
period="d CFB";
run;
proc sort data=chgbl &&lab&i;
by name;
run;
data lab &&lab&i;
length col2 $20;
set scr &&lab&i base &&lab&i c EOS &&lab&i chgbl &&lab&i;
if lbtest not in ("BASO", "EOSIN", "LYMPH", "MONO", "NEUT") then do;
     if name ="a N" then col2=put(col1,8.);
     else if _name_= "c_sd" then col2=put(round(col1,.1),8.1);
     else if name not in ("a N", "c sd") then
col2=put(round(col1,1.0),8.);
end;
else if lbtest in ("BASO", "EOSIN", "LYMPH", "MONO", "NEUT") then do;
     if name ="a N" then col2=put(col1,8.);
     else if _name ="c_sd" then col2=put(round(col1,.001),8.3);
     else if name not in ("a N", "c sd") then
col2=put(round(col1,.01),8.2);
end;
```

```
run;
%end;
data final;
set lab baso lab basoab lab eosin lab eosinab lab lymph lab lymphab
lab mono lab monoab lab neut lab neutab;
run;
proc datasets lib=work nolist;
save final;
run; quit;
%mend chgbase;
%chqbase;
proc format;
     value $reportf 'a N' = 'N'
                       'b mean'
                                    = 'MEAN'
                                 = 'SD'
                       'c sd'
                       'd min' = 'MIN'
                       'e_median' = 'MEDIAN'
'f max' = 'MAX'
                       'a Screening'= 'Screening'
                       'b Baseline' = 'Baseline, Day -1'
                       = 'Basophils Absolute
(10^6/L)'
                       'BASO'
                                  = 'Basophils %(fraction)'
                       'EOSINAB' = 'Eosinophils Absolute
(10^6/L)'
                       'EOSIN'
                                    = 'Eosinophils %(fraction)'
                       'LYMPHAB'
                                   = 'Lymphocytes Absolute
(10^6/L)'
                       'LYMPH'
                                    = 'Lymphocytes % (fraction)'
                       'MONOAB'
                                         = 'Monocytes Absolute
(10^6/L)'
                       'MONO'
                                    = 'Monocytes %(fraction)'
                       'NEUTAB'
                                         = 'Neutrophils Absolute
(10^6/L)'
                       'NEUT'
                                  = 'Neutrophils % (fraction)'
                       '0.00'
                                   = '0'
                        0.00' = 0';
run;
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
```

```
call symput('dtnull', today);
run:
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull,6,4);
/* Proc Report Step */
/* Make changes here for additional lab reports */
dm "odsresults; clear;";
ods escapechar='~';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-2c.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.16in bordercolor=black}
     style(header) = {font face=arial font size=2.5 foreground=black
bordercolor=black flow}
     columns lbtest period name col2 col1;
     *define order1 / order order=internal noprint;
     *define order2 / order order=internal noprint;
     define lbtest / group "Laboratory Parameter" format=$reportf.
style={cellwidth=25% just=c};
     define period / group order=internal "Timepoint" format=$reportf.
style={cellwidth=27.5% just=c};
     define _name_ / display "Statistic" format=$reportf.
style={cellwidth=27.5% just=c};
     define col2 / display "Result" center format=$reportf.
style={cellwidth=11% just=dec};
     define col1 / center style={cellwidth=8% just=c
foreground=white);
     compute before;
     line @1 ' ';
     endcomp;
     compute after;
     line @1 ' ';
```

```
endcomp;
     compute after period;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-2c
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Descriptive Statistics and Change from Baseline of
Hematology Laboratory Data: Differential Counts";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ~{thispage} of ~{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: Change from Baseline is the change from
Baseline, Day -1 compared to End of Study.";
     footnote2 j=1 '09'x "Lab units are presented in SI units.";
     footnote3;
     footnote4 j=1 "Source: Listing 16.2.8-3b";
run; quit;
ods listing;
ods rtf close;
/*
                                       OTR1021 Data Review
     Reviewer/Programmer: R. Valaboju
                                       06-18-**
     Date:
                                       OTR1021
     Study:
                                       14.3.5-2d
     Table:
     Notes:
                                       Re-programming sample tables as
part of QC.
libname r "P:\Medical\SRSProd\otr\otr1021\core\forbsp";
run;
options nolabel;
options mprint mlogic symbolgen;
```

/\* Make changes here for additional labs \*/

```
%let
str=%nrstr("BASOAB", "BASO", "EOSINAB", "EOSIN", "LYMPHAB", "LYMPH", "MONOAB
", "MONO", "NEUTAB", "NEUT");
%let
str2=%nrstr(BASOAB, BASO, EOSINAB, EOSIN, LYMPHAB, LYMPH, MONOAB, MONO, NEUTAB
, NEUT);
%let
str3=%nrstr(BASOAB||BASO||EOSINAB||EOSIN||LYMPHAB||LYMPH||MONOAB||MONO
||NEUTAB||NEUT);
/*-----Generic re-usable code starts here-----
/* Create subset of lab standard values dataset for the required tests
*/
proc sql;
create table lbsub as select usubjid, lbdtc, lbtpt, lbcat, lbtestcd,
lbstresn, lbnrind,
case when substr(lbnrind, 1, 1) ^='' then
trim(put(lbstresn,best12.)||' '||trim(substr(lbnrind,1,1))||', N')
else ' ' end as lbstresn2 from r.lb
where lbtestcd in (%unquote(&str));
quit;
/* Transpose the data subset to get standard values with corresponding
lab test column headings */
proc transpose data=lbsub out=lbsub;
by usubjid lbdtc lbtpt;
id lbtestcd;
var lbstresn2;
run;
/* Merge the lab subset with SC and DM datasets to get randomization
numbers and demographic data */
data lbsub;
length sex $10;
merge lbsub(in=a)
        r.sc(in=b keep=usubjid scorres where=(scorres like ("RN%")))
        r.dm(in=c keep=usubjid age sex race);
by usubjid;
if a;
if sex='F' then sex="Female";
if sex='M' then sex="Male";
run;
/* Create new variables required for reporting by concatenating data
data lbsub;
length rsub $25;
length asr $45;
length lbdtc2 8;
```

```
set lbsub;
rsub=trim(trim(scorres)||'/'||substr(usubjid, 15, 5));
asr=propcase(trim(trim(age)||'/'||trim(sex)||'/'||trim(race)));
lbdtc2=input(substr(lbdtc,1,10),yymmdd10.);
format lbdtc2 date9.;
run;
/* Create final dataset with required variables for reporting */
proc sql;
create table final as select rsub, asr, propcase(lbtpt) as lbtpt,
lbdtc2, substr(lbdtc, 12, 5) as time, %unquote(&str2)
from lbsub
where index((%unquote(&str3)),',') ^=0;
/*----End Generic code-----
____*/
/* Make changes here for additional lab reports */
/*Create macro variable for sysdate system macro variable*/
data null;
today = put(date(), date9.);
call symput('dtnull', today);
run;
%let dtnull2=%substr(&dtnull,1,2)-%substr(&dtnull,3,3)-
%substr(&dtnull, 6, 4);
/* Proc Report Step */
dm "odsresults; clear;";
ods escapechar='^';
options nonumber nodate orientation=landscape;
ods listing close;
ods rtf
file="\\nacl03\users ab\valabojr\Documents\OTR1021\reports\OTR1021 14-
3-5-2d.rtf" style=Styles.custom;
proc report
     data = final
     nowd
     ls=122
     ps=35
     headline
     headskip
     split='*'
     style(report) = {font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.5in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
```

```
where (basoab^='' or baso^='' or eosinab^='' or eosin^='' or
lymphab^='' or lymph^='');
     columns rsub asr lbtpt lbdtc2 time BASOAB BASO EOSINAB EOSIN
LYMPHAB LYMPH;
     define rsub / display group order order=internal center
"Randomization*No/Subject No" style={cellwidth=12% just=1};
     define asr / display group center "Age/Gender/*Race" flow
                                 style={cellwidth=12.125% just=c};
     define lbtpt / display center "Time-point"
                                      style={cellwidth=10% just=c};
     define lbdtc2 / display "Date"
                                            style={cellwidth=9%
just=c};
     define time / display "Actual*Time" center
                                      style={cellwidth=8% just=c};
     define BASOAB / display center "BASO Abs*(10^6/L)"
                                      style={cellwidth=8% just=c};
     define BASO / display center "BASO %*(fraction)"
                                 style={cellwidth=8% just=c};
     define EOSINAB / display center "EOS Abs*(10^6/L)"
                                      style={cellwidth=8% just=c};
     define EOSIN / display center "EOS %*(fraction)"
                                 style={cellwidth=8% just=c};
     define LYMPHAB / display center "LYMPH Abs*(10^6/L)"
                                 style={cellwidth=8% just=c};
     define LYMPH / display center "LYMPH %*(fraction)"
                                      style={cellwidth=8% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-2d
DATE: &dtnull2 TIME: &systime";
     title2 j=c "Subjects with Abnormal Hematology Laboratory Data
Based on the Reference Ranges: Differential Counts (Part 1 of 2)";
     *title3 j=c "(Page 1 of 1)";
     title3 j=c "Page ^{thispage} of ^{lastpage}";
     title4;
     title5 j=1 "Population: Randomized Safety";
     footnote1 j=1 "NOTE: H = laboratory value above the reference
laboratory range";
     footnote2 j=1 '09'x "L = laboratory value below the reference
laboratory range";
     footnote3 j=1 '09'x "A = laboratory value abnormal";
     footnote4 j=1 '09'x "C = laboratory value deemed clinically
significant by the investigator";
     footnote5 j=1 '09'x "N = laboratory value deemed not clinically
significant by the investigator.";
```

```
footnote6 j=l '09'x "Values out of the normal range are
flagged.";
     footnote7 j=1 '09'x "Lab units are presented as SI units.";
     footnote8 j=1 '09'x "Source: Listing 16.2.8-3b";
     footnote9 j=1 '09'x "Reference Range Source: Listing 16.2.8-1";
 run; quit;
proc report
     data = final
     nowd
     1s=122
     ps=35
     headline
     headskip
     split='*'
     style(report)={font face=arial font size=2.5 bordercolor=black}
     style(column)={just=center font face=arial font size=2.5
font size=2.5 cellheight=.5in bordercolor=black}
     style(header)={font face=arial font size=2.5 foreground=black
bordercolor=black flow}
where (monoab^='' or mono^='' or neutab^='' or neut^='');
     columns rsub asr lbtpt lbdtc2 time MONOAB MONO NEUTAB NEUT;
     define rsub / display group order order=internal center
"Randomization*No/Subject No" style={cellwidth=14% just=1};
     define asr / display group center "Age/Gender/*Race" flow
                                 style={cellwidth=12.125% just=c};
     define lbtpt / display center "Time-point"
                                      style={cellwidth=13% just=c};
     define lbdtc2 / display "Date"
                                            style={cellwidth=10%
just=c};
     define time / display "Actual*Time" center
                                      style={cellwidth=10% just=c};
     define MONOAB / display center "MONO Abs*(10^6/L)"
                                      style={cellwidth=10% just=c};
     define MONO / display center "MONO %*(fraction)"
                                 style={cellwidth=10% just=c};
     define NEUTAB / display center "NEUT Abs*(10^6/L)"
                                      style={cellwidth=10% just=c};
     define NEUT / display center "NEUT %*(fraction)"
                                 style={cellwidth=10% just=c};
     compute before;
     line @1 ' ';
     endcomp;
     title1 j=l "OTR1021
                Table 14.3.5-2d
DATE:&dtnull2 TIME:&systime";
```

```
title2 j=c "Subjects with Abnormal Hematology Laboratory Data
Based on the Reference Ranges: Differential Counts (Part 2 of 2)";
    *title3 j=c "(Page 1 of 1)";
    title3 j=c "Page ^{thispage} of ^{lastpage}";
    title4;
    title5 j=l "Population: Randomized Safety";
run;quit;
ods listing;
ods rtf close;
```