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*****;
* Project          : ZRHM-REXA-07-JP
*
* Program name     : T150204240201_ZRHM-REXA-07_V1.sas
*
* Author          : L. Yan
*
* Date created     : 05/20/2015
*
* Purpose         : Table T150204240201
*
* Revision History :
*
* Date          Author      Ref      Revision (Date in YYYYMMDD format)
*
*****;

%let prgname=T150204240201_ZRHM_REXA_07_JP_V1;
options mprint;

options sasautos=("W:\pmp07\macros" sasautos) notes;
%init(delivery=9);

%titlecsv(prgname=&prgname., version=5);

%put &title1;
%put &title2;
%put &APPENDIX;
%put &endpoint;
%put &outname.;

options missing="";

%macro cal_sumary_pvalue(wher=, outnum=, var=, in=, pflg=, used=, paramcd=, avisit=);
title2 h=10pt j=1 "&used";

proc sort data=&in. out=anadt_&outnum.;
by usubjid;
where &wher. ;
run;

proc sort data=anadt_&outnum.;
by trtcd;
run;

proc means data = anadt_&outnum. noprint;
by trtcd;
var &var.;
output out=xlab_&outnum. n=n mean=mean median=med std=sd min=min max=max q1=q1 q3=q3 lclm=lclm uclm=uclm;
run;

data xlab_&outnum.;
set xlab_&outnum.;
n1 = trim(left(compress(put(n, 8.))));
if sd > . then mean1 = (trim(left(compress(put(mean, 8.1))))||' ( '||trim(left(compress(put(ceil(sd*100)/100, 8.2))))||')');
else mean1 = (trim(left(compress(put(mean, 8.1))))||' (NA)');
ci1=trim(left(compress(put(floor(lclm*100)/100, 8.2))))||' , '||trim(left(compress(put(ceil(uclm*100)/100, 8.2)))));
median1 = trim(left(compress(put(med, 8.1))));
q1q3 = trim(left(compress(put(q1, 8.2))))||' , '||trim(left(compress(put(q3, 8.2))));
min1 = trim(left(compress(put(min, 8.))))||' , '||trim(left(compress(put(max, 8.0))));
run;

/*
proc mixed data=anadt_&outnum.;

Class trtcd sex UCPDGR1;

Model logaval = logbase sex UCPDGR1 trtcd / outp=pred;

lsmeans trtcd / pdiff =control('mCC') alpha=0.05 cl;

ods output lsmeans=lsmeans_&outnum. (keep=trtcd lower upper estimate); *each arm;

ods output diffs=LSMeanDiffCL&outnum. (keep=trtcd lower upper probt estimate); * lsmean and C.I. for ratios;

ods output covparms=ROOTMSE&outnum. (rename=(estimate=mse)); *MSE;

run;
*/

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title3 h=10pt j=1 "Paramcd: &paramcd, &avisit. Model: MIXED, Method: Log";

proc mixed data=anadt &outnum.;
class trtp sex UCPDGR1;
model logaval = logbase sex UCPDGR1 trtp/ outp=pred;
lsmeans trtp / pdiff =control('mCC') alpha=0.05 cl;
lsmeans trtp / pdiff =control('SA') alpha=0.05 cl;
ods output lsmeans=lsmeans_&outnum. (keep=trtp lower upper estimate); *each arm;
ods output diffs=LSMeanDiffCL&outnum. (keep=_trtp trtp lower upper probt estimate where=(TRTP="THSm2.2")); * lsmean and
C.I. for ratios;
ods output covparms=estimate&outnum.(rename=(estimate=rootmse)); *MSE;
run;
ods output close;

data pval&outnum.;
set LSMeanDiffCL&outnum.;
ProbtDiff=probt;
keep trtp ProbtDiff;
run;

data lsmeans_&outnum.;
set lsmeans_&outnum.;
lowercl=lower;
uppercl=upper;
lsmean=estimate;
keep trtp lowercl uppercl lsmean;
run;

data LSMeanDiffCL&outnum.;
set LSMeanDiffCL&outnum.;
lowercl=lower;
uppercl=upper;
difference=estimate;
keep trtp _trtp lowercl uppercl difference;
run;

data lsmeans_&outnum.;
set lsmeans_&outnum.;
if TRTP="THSm2.2" then trtcd=1;
else if TRTP="mCC" then trtcd=2;
else if TRTP="SA" then trtcd=3;

    Estimate1 = exp(lsmean); /* Ratio of geometric mean */
    LowerCL   = exp(lowercl); /* 95% CI lower bound */
    UpperCL   = exp(uppercl); /* 95% CI upper bound */
run;

data ROOTMSE&outnum.;
set estimate&outnum.;
*CVperc=100*sqrt(exp(rootmse**2)-1);
cvperc=100*sqrt(exp(rootmse)-1);
run;

proc sort data=lsmeans_&outnum. nodupkey;
by trtcd;
run;

data lsmeans_&outnum.;
length geomean geoci $100;
set lsmeans_&outnum.;
geomean=strip(put(ESTIMATE1, 8.2));
geoci=strip(put(floor(LowerCL*100)/100, 8.2)||", "||strip(put(ceil(UpperCL*100)/100, 8.2)));
keep trtcd geomean geoci;
run;

proc sort data=LSMeanDiffCL&outnum. nodup;
by TRTP _TRTP;
run;

data LSMeanDiffCL&outnum.;
set LSMeanDiffCL&outnum.;
myord=1;
run;

data ROOTMSE&outnum.;
set ROOTMSE&outnum.;
myord=1;
run;

data LSMeanDiffCL&outnum.;
merge LSMeanDiffCL&outnum. ROOTMSE&outnum.;
by myord;
run;

data LSMeanDiffCL&outnum.;
length geomean geoci $100;
set LSMeanDiffCL&outnum.;

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if _TRTP eq "mCC" then trtcd=4;
if _TRTP eq "SA" then trtcd=5;

    difference = 100*exp(difference); /* Ratio of geometric mean */
    lowercl = 100*exp(lowercl); /* 95% CI lower bound */
    uppercl = 100*exp(uppercl); /* 95% CI upper bound */

geomean=strip(put(difference, 8.2))||" ("||strip(put(ceil(CVperc*100)/100, 8.2)) ||")";
geoci=strip(put(floor(lowercl*100)/100, 8.2))||", "||strip(put(ceil(uppercl*100)/100, 8.2));
keep trtcd geomean geoci;
run;

data pval&outnum.;
set pval&outnum.;
if _n_=1;
trtcd=4;
keep trtcd ProbtDiff;
format ProbtDiff PVALUE6.3;
run;

proc sort data=LSMeanDiffCL&outnum.;
by trtcd;
run;

data LSMeanDiffCL&outnum.;
merge LSMeanDiffCL&outnum. pval&outnum.;
by trtcd;
run;

data mrep_&outnum.;
set lsmeans_&outnum. LSMeanDiffCL&outnum.;
run;

proc sort data=mrep_&outnum.;
by trtcd;
run;

proc sort data=xlab_&outnum.;
by trtcd;
run;

data xlab_&outnum.;
merge xlab_&outnum.(in=a) mrep_&outnum.;
by trtcd;
run;

data xlab_&outnum.;
set xlab_&outnum.;
if geoci ne "";
run;

proc transpose data = xlab_&outnum. out=xlab_1_&outnum.;
id trtcd;
var n1 geomean geoci ProbtDiff;
run;

data rep_&outnum.;
length _name_ _1 _2 _3 ord1 $100;
set xlab_1_&outnum.;
ord1="&outnum";
ordnum=input(ord1, best.);
if upcase(_name_)="N1" then do; _name_="n"; sord=0; end;
if upcase(_name_)="GEOMEAN" then do; _name_="Geometric LS Mean (CV%)"; sord=1; end;
if upcase(_name_)="GEOCI" then do; _name_="95% CI"; sord=2; end;

if upcase(_name_)="MEAN1" then do; _name_="Mean (SD)"; sord=3; end;
if upcase(_name_)="CI1" then do; _name_="95% CI"; sord=4; end;
if upcase(_name_)="MEDIAN1" then do; _name_="Median"; sord=5; end;
if upcase(_name_)="Q1Q3" then do; _name_="Q25, Q75"; sord=6; end;
if upcase(_name_)="MIN1" then do; _name_="Min, Max"; sord=7; end;
if upcase(_name_)="PROBTDIFF" then do; _name_="p-value (one-sided)"; sord=9; end;
run;

data rep;
set rep rep_&outnum.;
run;

%mend;

%macro mainloop(flgl, outn=, where=);

proc sort data=adam.adsl out=trt;
by usubjid;
where &flgl="Y";
run;

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data trt;
set trt;
if TRT01A="THSm2.2" then trtcd=1;
else if TRT01A="mCC" then trtcd=2;
else if TRT01A="SA" then trtcd=3;
run;

data anldata1;
set adam.adbx;
used="The where clause used on the dataset adam.adbx: anl02f1=Y and anl03f1=Y and &flg =Y";
if trta in ("mCC" "THSm2.2" "SA") and aval>. and &flg.="Y" and paramcd="CYP1A2" and &where. and anl02f1="Y" and anl03f1
="Y"/*and dtype ne "LOCF"*/;
run;

data anldata1;
set anldata1;

if avisitn=105 and not (anl01f1="Y" and anl02f1="Y") then delete;

if aval>. then logaval=log(aval);
if base>. then logbase=log(base);
run;

proc sort data=anldata1 out=check(keep=paramn avisitn avisit used paramcd) nodupkey;
by paramn avisitn avisit;
run;

data trt_1;
set trt;
run;

data anldata1;
set anldata1;
if TRTP="THSm2.2" then trtcd=1;
else if TRTP="mCC" then trtcd=2;
else if TRTP="SA" then trtcd=3;
run;

data check;
set check;
ord=_n_;
run;

%*cal_summary_pvalue(where=1, outnum=1, var=aval, in=anldata1, pflg=1);

data rep;
run;

data _null_;
set check;
call execute ('%cal_summary_pvalue(where=%str(avisitn=||avisitn|| and paramn=||paramn|| ), outnum=||ord||, var
=logaval, in=anldata1, used=||used||, paramcd=||paramcd||, avisit=||avisit||)');
run;

data frep;
set rep;
ord=ORDNUM;
run;

data frep;
merge frep(in=a) check;
by ord;
if a;
if avisitn>.;
run;

proc sort data=trt_1 nodupkey;
by trtcd usubjid;
run;

proc freq data = trt_1 noprint;
tables trtcd/ out= denom;
run;

data _null_;
set denom end=eof;

retain total 0;

total = total+count;

if trtcd= 1 then do;

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        call symput('trt1', trim(left(put(count,8))));
    end;
    if trtcd= 2 then do;
        call symput('trt2', trim(left(put(count,8))));
    end;
    if trtcd= 3 then do;
        call symput('trt3', trim(left(put(count,8))));
    end;

run;

%put trt1=&trt1 trt2=&trt2 trt3=&trt3;

proc format;
value grp
    49      ="Nicotine-Glucuronide (ng/mL)"
    3       ="Cotinine (ng/mL)"
;
run;

%macro cal_part_main();

data frep;
set frep;

    %do i = 1 %to 100;
        if (&i-1)*4<ordnum<=&i*4 then pagen=&i;
    %end;

run;

%mend;

%cal_part_main();
data frep&outn.;
set frep;
space=" ";
run;

%mend;

%trtrtfg(pgmname=&loutname., pgmid=1, new=0, style=, bookmark=%lowcase(&outname.));

title1 bold j=1 "&title1 &title2";

%mainloop(flag=fasfl, outn=1, where=%str(avisitn in (105, 190)));

%*mainloop(flag=PPROT4FL, outn=4, where=%str(avisitn in (190)));

ods listing;
ods rtf close;

data odata.&prgname.;
set frep1 (in=a) /*frep4 (in=d)*/;
if a then group="fasfl";
*if d then group="PPROT4FL";
run;

proc sort data=anldata1 out=fmt(keep=paramn param) nodupkey;
by paramn param;
run;

data fmt;
set fmt;
fmtname="grp";
start=paramn;
label="Parameter: "||strip(param);
run;

proc format cntlin=fmt;
run;

data final;
set frep1 /*frep4*/;
run;

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proc sort data=final;
by pagen;
run;

%global totalpage;

data _null_;
  set final end=eof;

  if eof then do;
    call symput('totalpage', trim(left(put(pagen,8)))));
  end;

run;

%put totalpage=&totalpage;

%*title(prgname1=&prgname.);

%trtrtfpg(pgmname=&outname., pgmid=1, new=0, style=, bookmark=%lowercase(&outname.));

proc report data=final headskip headline spacing=4 nowd split='~' style=[outputwidth=100%] style(header column)=[protec
tspecialchars=off];
  column pagen paramn avisitn avisit sord _name_ _1 space _2 space _3 space _4 space _5;
  define pagen /order order=internal noprint;
  define paramn /order order=internal noprint;
  define avisitn /order order=internal noprint;
  define avisit /order "Time Point" flow style(column)=[cellwidth=7% just=l];
  define sord /order order=internal noprint;

  define _name_ /display "Statistic" flow style(column)=[cellwidth=18% just=l];
  define _1 /display "THSm2.2" flow style(column)=[cellwidth=10% just=c];
  define space /display " " flow style(column)=[cellwidth=0.5% just=c];

  define _2 /display "mCC" flow style(column)=[cellwidth=10% just=c];
  define space /display " " flow style(column)=[cellwidth=0.5% just=c];

  define _3 /display "SA" flow style(column)=[cellwidth=10% just=c];
  define space /display " " flow style(column)=[cellwidth=0.5% just=c];

  define _4 /display "THSm2.2 : mCC Ratio (%)" flow style(column)=[cellwidth=10% just=c];
  define space /display " " flow style(column)=[cellwidth=0.5% just=c];

  define _5 /display "THSm2.2 : SA Ratio (%)" flow style(column)=[cellwidth=10% just=c];

/*
  COMPUTE before paramn ;
  LINE @1 paramn grp.;
  ENDCOMP;
*/
  COMPUTE after avisitn ;
  LINE @1 "";
  ENDCOMP;

  compute before pagen;
  line @1 "";
  endcomp;

  compute before _page_ /style=[fontweight=bold fontsize=3.75];
  line @1 "&title1 &title2";
  line @1 " ";
  line @1 paramn grp.;
  line @1 " ^R/RTF'\brdrb\brdrs\brdrw30\brsp20\b ' ";
  endcomp;

  compute after _page_ /style=[fontsize=1.75];
  line @1 "Note: mCC = Menthol conventional cigarettes; SA = Smoking abstinence; THSm2.2 = Tobacco Heating System 2.2 Ment
hol.";
  line @1 "Note: Adjusted geometric least squares (LS) means and confidence intervals (CIs) from an mixed model conducted
on log-transformed";
  line @1 "values with log-transformed baseline value, study arm, sex and mCC consumption reported at screening as fixed e
ffect factors. ";
  line @1 "Geometrical CV% of the ratio is estimated from the residual mean squares.";
  line @1 "Note: p-value for one-sided test is for comparison between THSm2.2 and mCC";
  line @1 " ";
  line @1 "&APPENDIX.";
  line @1 "Study ID:ZRHM-REXA-07-JP Program: &fprgname..sas Status: &repversion./&fdate. Page: 1 o
f &totalpage";
  endcomp;

break after pagen/page;

run;

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ods listing;  
ods rtf close;
```