```
TA
```

```
libname output '/home/u58485303/output';
proc import datafile= '/home/u58485303/raw/Trial design specification.xlsx'
out= ta dbms= xlsx replace;
sheet= 'TA';
run;
data output.ta (label= 'Trial Arms');
attrib STUDYID label= 'Study Identifier' length= $200
              DOMAIN label= 'Domain Abbreviation' length= $200
              ARMCD label= 'Planned Arm Code' length= $200
              ARM label= 'Description of Planned Arm' length= $200
              TAETORD label= 'Planned Order of Element within Arm' length= 8
              ETCD label= 'Element Code' length= $200
              ELEMENT label= 'Description of Element' length= $200
              TABRANCH label= 'Branch' length= $200
              TATRANS label= 'Transition Rule' length= $200
              EPOCH label= 'Epoch' length= $200;
set ta;
keep STUDYID DOMAIN
                             ARMCD ARM TAETORD ETCD ELEMENT TABRANCH TATRANS
       EPOCH;
run;
TE
libname output '/home/u58485303/output';
proc import datafile= '/home/u58485303/raw/Trial design specification.xlsx'
out= te dbms= xlsx replace;
```

```
sheet= 'TE';
run;
data output.te (label= 'Trial Elements');
attrib STUDYID label= 'Study Identifier' length= $200
               DOMAIN label= 'Domain Abbreviation' length= $200
               ETCD label= 'Element Code' length= $200
               ELEMENT label= 'Description of Element' length= $200
               TESTRL label= 'Rule for Start of Element' length= $200
               TEENRL label= 'Rule for End of Element' length= $200
               TEDUR label= 'Planned Duration of Element' length= $200;
set te;
keep STUDYID DOMAIN
                              ETCD ELEMENT TESTRL
                                                            TEENRL TEDUR;
run;
<u>TI</u>
libname output '/home/u58485303/output';
proc import datafile= '/home/u58485303/raw/Trial design specification.xlsx'
out= ti dbms= xlsx replace;
sheet= 'TI';
run;
data output.ti (label= 'Trial Inclusion/Exclusion Criteria');
attrib STUDYID label= 'Study Identifier' length= $200
               DOMAIN label= 'Domain Abbreviation' length= $200
               IETESTCD label= 'Incl/Excl Criterion Short Name' length= $200
               IETEST label= 'Inclusion/Exclusion Criterion' length= $200
               IECAT label= 'Inclusion/Exclusion Category' length= $200;
```

```
set ti;
keep STUDYID DOMAIN
                              IETESTCD IETEST
                                                     IECAT;
run;
<u>TV</u>
libname output '/home/u58485303/output';
proc import datafile= '/home/u58485303/raw/Trial design specification.xlsx'
out= tv dbms= xlsx replace;
sheet= 'TV';
run;
data output.tv (label= 'Trial Visits');
attrib STUDYID label= 'Study Identifier' length= $200
               DOMAIN label= 'Domain Abbreviation' length= $200
               VISITNUM label= 'Visit Number' length= 8
               VISIT label= 'Visit Name' length= $200
               VISITDY label= 'Planned Study Day of Visit' length= 8
               ARMCD label= 'Planned Arm Code' length= $200
               ARM label= 'Description of Planned Arm' length= $200
               TVSTRL label= 'Visit Start Rule' length= $200
               TVENRL label= 'Visit End Rule' length= $200;
set tv;
keep STUDYID DOMAIN
                              VISITNUM VISIT VISITDY ARMCD ARM TVSTRL TVENRL;
run;
<u>TS</u>
```

libname output '/home/u58485303/output';

```
proc import datafile= '/home/u58485303/raw/Trial design specification.xlsx'

out= ts dbms= xlsx replace;
sheet= 'TS';
run;

data output.ts (label= 'Trial Summary');
attrib STUDYID label= 'Study Identifier' length= $200

DOMAIN label= 'Domain Abbreviation' length= $200

TSSEQ label= 'Sequence Number' length= 8

TSPARMCD label= 'Trial Summary Parameter Short Name' length= $200

TSPARM label= 'Trial Summary Parameter' length= $200

TSVAL label= 'Parameter Value' length= $200;
set ts;
keep STUDYID DOMAIN

TSSEQ TSPARMCD TSPARM TSVAL;
run;
```