SAS[®] GLOBAL FORUM 2021

Paper 1102-2021 Generating .xpt files with SAS, R and Python Todd Case and YuTing Tian

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ABSTRACT

The primary purpose of this paper is to first lay out a process of generating a simplified Transport (.xpt) file with RStudio and Python to meet study electronic data submission requirements of the Food & Drug Administration (FDA). The second purpose of this paper is to compare the .xpt files created from three different languages: R, Python and SAS. The paper is the expansion of the original FDA guideline document "CREATING SIMPLIFIED TS.XPT FILES", published in November, 2019.

Transport files can be created by SAS, *as well as open source software*, including R and Python. According to the FDA guideline document mentioned above, .xpt files can be created by R and Python. This may allow Pharmaceutical companies to expand use of R and Python beyond data visualization and statistical analysis currently being generated by these two languages. Hopefully, readers can use the process shown in the paper as a template to create .xpt files.

INTRODUCTION

Transport files are in use in the pharmaceutical industry as a result of FDA e-data submission requirements. The "Creating Simplified TS.XPT Files" is the specific guide to help sponsors create TS (ts.xpt) files with R and Python to meet study data submission requirements.

The paper is divided into four sections which correspond to the steps outlined in our suggested processes: the first three sections introduce how to produce .xpt files with SAS,R and Python, respectively, and the fourth section compares the results of simplified .xpt files generated from the different software packages using SAS Universal Viewer (the application to view SAS Transport .xpt files).

- 1) Using SAS to generate final datasets in .xpt format
 - 1.1) Code to generate raw TS domain in SAS
 - 1.2) Export TS.SAS with .xpt format
 - 1.3) Using macro to create .xpt files with SAS
- 2) Using RStudio to generate final datasets in xpt format
 - 2.1) Code to generate raw TS domain in RStudio
 - 2.2) Export TS.R file with .xpt format
 - 2.3) A fast way to create All .xpt files with RStudio
- 3) Using Python to generate final datasets in .xpt format
 - 3.1) Code to generate raw TS domain in Python
 - 3.2) Export TS.py file with .xpt format
- 4) Review and Compare generated ts.xpt file in SAS Universal Viewer

1). USING SAS TO GENERATE FINAL DATASETS IN .XPT FORMAT

1.1) CODE TO GENERATE RAW TS DOMAIN IN SAS

The Trial Summary domain is used to record basic information about the study such as protocol title, trial phase, etc. The purpose of this paper is to introduce the process of generating xpt files with different languages in terms of the original paper "Creating Simplified TS.XPT Files. We assume the readers have a fundamental knowledge of using SAS, RStudio and Python already, therefore we start our paper with raw data already cleaned with SAS, then the same raw Trial Summary dataset is applied into SAS, RStudio and Python to generate a ts.xpt file separately. The fabricated TS raw dataset is produced as Figure1 shown below:

| Data TS; | | | | | | | | |
|---------------------|----------------------------------|----------------------------|-------------|-------------------------|------------------|---------------------|-----------------|-----------------|
| infile datalines t | runcover firstok | s=2; | | | | | | |
| length STUDYID \$3 | 20 DOMAIN \$2 | TSSEQ 8. | TSGR | PID \$40. | TSPARMCI |) \$ <mark>8</mark> | TSPARM | \$ 40 |
| TSVAL | TSVALNF | TSVALCD \$2 | 00 TSVC | DREF \$ <mark>20</mark> | TSVCDVER | R \$ 10; | | |
| attrib | | | | | | | | |
| STUDYI |) label = | 'Study Iden | tifier' | | | | | |
| DOMAI | N label = | 'Domain Abb | reviation | • | | | | |
| TSSEQ | label = | 'Sequence N | umber' | | | | | |
| TSGRP | ID label = | 'Group ID' | | | | | | |
| TSPAR | MCD label = | • 'Trial Summ | ary Param | eter Short | Name' | | | |
| TSPAR | 4 label = | 'Trial Summ | ary Param | eter' | | | | |
| TSVAL | label = | 'Parameter | Value' | | | | | |
| TSVAL | NF label = | 'Parameter | Null Flav | or' | | | | |
| TSVAL | CD label = | 'Parameter | Value Cod | e' | | | | |
| TSVCD: | REF label = | Name of th | e Referen | ce Terminol | Logy' | | | |
| TSVCD | /ER label=' | Version of t | he Refere | nce Termino | ology'; | | | |
| input @1 STUDYID \$ | 1-3 @5 DOMAIN | I\$ <mark>5-6</mark> @ 8] | 'SSEQ 1. | 010 TSGRPII | \$ 10-22 | | | |
| 023 TSPARMC |) \$ 23-31 @ 32 TS | PARM \$ 32-57 | @ 59 | TSVAL \$ 59- 7 | 72 @74 TS | SVALNF | \$ 74-75 | |
| 076 TSVALCD | \$ 76-82 @83 | TSVCDREF \$ 8 | 3-91 @93 | TSVCDVER \$ | 93-103 @ | | | |
| ; | | | | | | | | |
| datalines; | | | | | | | | |
| 123456789012345678 | 9012345678901234 | 567890123456 | 789012345 | 67890123456 | 5789012345 | 5678901 | 235678901 | 234567890123456 |
| 001 TS 1 | ACTSUB Act | ual Number o | of Subject | s 10 | | | | |
| 001 TS 1 | ADAPT Ada | ptive Design | 1 | N | | C49487 | CDISC | 2019-12-20 |
| 001 TS 1 | AGEMAX Max | imum Age of | Subjects | P65Y | | | ISO 8601 | |
| 001 TS 1 | AGEMIN Mir | imum Age of | Subjects | P18Y | | | ISO 8601 | |
| 001 TS 1 | DCUTDESC Dat | a Cutoff Des | cription | DATABASE | E LOCK | | | |
| 001 TS 1 | DCUTDTC Dat | a Cutoff Dat | e | 2020-11- | <mark>-26</mark> | | | |
| 001 TS 1 group1,dr | ugl DOSE Dos | e per Admini | stration | 400 | | | | |
| 001 TS 2 group2,dr | 1g2 DOSE Dos | e per Admini | stration | 15 | | | | |
| 001 TS 1 drug1 | DOSFRM Dos | e Form | | TABLET | | C42998 | ISO 8601 | |
| 001 TS 1 drug2 | DOSFRM Dos | e Form | | CAPSULES | 5 | C42998 | ISO 8601 | |
| ; | | | | | | | | |
| run: | | | | | | | | |



Figure1 is a simulated dataset - we have created the eleven standard CDISC (Clinical Data Interchange Standards Consortium) variables that exist in the TS domain. We assume the readers are already familiar with the Trial Summary dataset, so we do not explain each variable and value in detail.

This fabricated data shows some parameters that are required or expected in the Trial Summary dataset, the STUDYID is "001"; the DOMAIN is "TS"; there are two drugs: -drug1 and drug2; TSPARMCD is the short name of the parameter, such as "ACTSUB", "AGEMAX" here; TSPARM is the term for the parameter in corresponding with TSPARMCD, such as "Actual Number of Subjects", "Maximum Age of Subjects"; TSVAL is the value of TSPARM, such as the number of subjects in this study is "10", the maximum age of subjects is P65Y, meaning 65 years old, etc.

| | | STUDYID | DOMAIN | TSSEQ | TSGRPID | TSPARMCD | TSPARM | TSVAL | TSVALNF | TSVALCD | TSVCDREF | TSVCDVER |
|----|-----|---------|--------|-------|-----------------|----------|---------------------------|---------------|-----------|---------|----------|------------|
| 1 | 001 | | TS | 1 | | ACTSUB | Actual Number of Subjects | 10 | 0.000.000 | | | |
| 2 | 001 | | TS | 1 | | ADAPT | Adaptive Design | N | | C49487 | CDISC | 2019-12-20 |
| 3 | 001 | | TS | 1 | | AGEMAX | Maximum Age of Subjects | P65Y | | | ISO 8601 | |
| 4 | 001 | | TS | 1 | | AGEMIN | Minimum Age of Subjects | P18Y | | | ISO 8601 | |
| 5 | 001 | | TS | 1 | | DCUTDESC | Data Cutoff Description | DATABASE LOCK | | | | |
| 6 | 001 | | TS | 1 | | DCUTDTC | Data Cutoff Date | 2020-11-26 | | | | |
| 7 | 001 | | TS | 1 | group 1, drug 1 | DOSE | Dose per Administration | 400 | | | | |
| 8 | 001 | | TS | 2 | group2,drug2 | DOSE | Dose per Administration | 15 | | | | |
| 9 | 001 | | TS | 1 | drug1 | DOSFRM | Dose Form | TABLET | | C42998 | ISO 8601 | |
| 10 | 001 | | TS | 1 | drug2 | DOSFRM | Dose Form | CAPSULES | | C42998 | ISO 8601 | |

Figure2

Figure2 shows the result of the code from Figure1 in SAS.

1.2) EXPORT IT WITH XPT FORMAT

| <pre>libname sasfile "E:\users\tiany"; libname xptfile XPORT "E:\users\tiany\ts.xpt";</pre> | 1 |
|---|---|
| <pre>data xptfile.ts; set sasfile.ts2; run;</pre> | 2 |
| /*or another way to xport*/ | |
| <pre>proc copy in=sasfile out=xptfile memtype=data; select ts2; run;</pre> | 3 |

Figure3

Figure3 is the code to generate the ts.xpt file.

1: Define two libnames: "sasfile" for saving sas dataset and "xptile" for saving file with .xpt format.

Notice: when we create a path for storing xpt file, we should be specific the name of the .xpt file we are creating into path using XPORT engine, such as: "E:\users\tiany\ts.xpt" here.

There are two different ways of creating xpt file as below:

2: create xpt file with "data step"

3: to create xpt file with "proc copy" statement.

The process is straightforward as shown in figure2.

| This PC 🔸 | (E:) ► users ► tiany | | | | | | | |
|-----------|----------------------|--------------------|-------------------|-------|--|--|--|--|
| | Name | Date modified | Туре | Size | | | | |
| | 🗮 ts | 11/26/2020 3:10 PM | SAS Xport Transpo | 10 KB | | | | |

Figure4

Finally, the ts.xpt file is created in SAS. In Windows Explorer we can see the file create, shown in Figure 4.

1.3) A USEFUL MACRO TO CREATE XPT FILES WITH SAS

%*createxpt*(inlib=sdtm, xptdir=.\xpt);

In the pharmaceutical company, a macro we call %createxpt can create several .xpt files effectively and efficiently. This macro includes two required parameters, inlib=XX, the name of the SAS library containing the input dataset, such as SDTM; xptdir is the folder where the .xpt files will be created. When invoking the macro %createxpt, then xpt files are created, Figure5 is shown as below.

| This PC | | | ▶ xpt ▶ | |
|---------|------|-------------------|-------------------|-------|
| | Name | Date modified | Туре | Size |
| | 📬 ae | 4/23/2020 3:15 PM | SAS Xport Transpo | 25 KB |
| | 📬 cm | 4/23/2020 3:15 PM | SAS Xport Transpo | 14 KB |
| | 📬 da | 4/23/2020 3:15 PM | SAS Xport Transpo | 8 KB |
| 1 | 📑 dm | 4/23/2020 3:15 PM | SAS Xport Transpo | 8 KB |

Figure5

2)USING RSTUDIO TO GENERATE FINAL DATASETS IN XPT FORMAT

2.1) CODE TO GENERATE RAW TS DOMAIN IN RSTUDIO

R is a programming language developed cooperatively and noncommercially; RStudio is a commercial product - it is an integrated development environment as a tool for statistical computing and graphics. In this section, as an extension of the original document "Creating Simplified TS.XPT Files", we use RStudio as one of the programming languages to create raw TS dataset, SDTM.TS dataset and export it with xpt format.

| <pre>##option 1 package## Install.packages('SASxport')</pre> | 1 |
|--|---|
| library(SASxport) | |
| library(Hmisc) | 2 |
| Library(sas7bdat) | 3 |
| ##option 2 package## | |
| Install.packages('haven') | 4 |
| library(Hmisc) | |
| library(haven) | |
| | |
| | |

Figure6

After installing R and RStudio, then you can use either option1 "SASXport" package or option2 "Haven" package. For each package, there are three steps separately as shown below:

1: Install the package "SASXport". This package is to provide functions to read, list contents and write SAS export files.

2: Invoke the library function to load it into the current R session: library(SASxport), library(Hmisc).

The Hmisc library contains many functions such as useful data analysis. We need to use data frame function and label function under Hmisc library.

3: then we can use library(sas7bdat) to read SAS datasets into R.

4: the second way is to install the package "Haven". Then we invoke library(Hmisc) and library(haven) to read SAS datasets into R.

As Figure6 is shown above.

| ts<-data.frame | (STUDYID | = c("001","001","001","001","001","001","001","001","001","001"), |
|----------------|-----------|--|
| | DOMAIN | = "TS", |
| | TSSEQ | = c(1,1,1,1,1,1,1,2,1,1), |
| | TSGRPID | <pre>= c("","","","","",",",",",",",",",",",","</pre> |
| | TSPARMCD | = c("ACTSUB", "ADAPT", "AGEMAX", "AGEMIN", |
| | | "DCUTDESC", "DCUTDTC", "DOSE", "DOSE", "DOSFRM", "DOSFRM"), |
| | TSPARM | = c("Actual Number of Subjects", |
| | | "Adaptive Design", |
| | | "Maximum Age of Subjects", |
| | | "Minimum Age of Subjects", |
| | | "Data Cutoff Description", |
| | | "Data Cutoff Date", |
| | | "Dose per Administration", |
| | | "Dose per Administration", |
| | | "Dose Form", |
| | | "Dose Form"), |
| | TSVAL | = c("10", "N", "P65Y", "P18Y", |
| | | "DATABASE LOCK", |
| | | strftime(as.Date("2020-11-26",format="%Y-%m-%d")), |
| | | "400", "15", |
| | | "TABLET", "CAPSULES"), |
| | TSVALNF | = c(""), |
| | TSVALCD | = c("","C49487","","","","","","C42998","C42998"), |
| | TSVCDREF | = c("","CDISC","ISO 8601","ISO 8601","","","","","ISO 8601","ISO 8601"), |
| | TSVCDVER | = c("",strftime(as.Date("2019-12-20",format="%Y-%m-%d")),"","","", |
| | | "","","","",""),stringsAsFactors = FALSE) |
| | | |
| View(ts) | | |

Figure7

Figure7 shows how to create Trial Summary dataset using data.frame function in RStudio. We use data.frame function to collect variables which share many of the properties of lists, then we use a series of c vectors to store values. As we mentioned above, we assume readers have a rudimental knowledge of using RStudio, therefore we won't explain the code in detail.

| STUDYID [‡] | DOMAIN [‡] | TSSEQ [‡] | TSGRPID [‡] | TSPARMCD [‡] | † | †SVAL [‡] | TSVALNF [‡] | TSVALCD [‡] | TSVCDREF [‡] | TSVCDVER [‡] |
|----------------------|---------------------|--------------------|----------------------|-----------------------|---------------------------|--------------------|----------------------|----------------------|-----------------------|-----------------------|
| 001 | TS | 1 | | ACTSUB | Actual Number of Subjects | 10 | | | | |
| 001 | TS | 1 | | ADAPT | Adaptive Design | N | | C49487 | CDISC | 2019-12-20 |
| 001 | TS | 1 | | AGEMAX | Maximum Age of Subjects | P65Y | | | ISO 8601 | |
| 001 | TS | 1 | | AGEMIN | Minimum Age of Subjects | P18Y | | | ISO 8601 | |
| 001 | TS | 1 | | DCUTDESC | Data Cutoff Description | DATABASE LOCK | | | | |
| 001 | тs | 1 | | DCUTDTC | Data Cutoff Date | 2020-11-26 | | | | |
| 001 | TS | 1 | group1,drug1 | DOSE | Dose per Administration | 400 | | | | |
| 001 | TS | 2 | group2,drug2 | DOSE | Dose per Administration | 15 | | | | |
| 001 | TS | 1 | drug1 | DOSFRM | Dose Form | TABLET | | C42998 | ISO 8601 | |
| 001 | TS | 1 | drug2 | DOSFRM | Dose Form | CAPSULES | | C42998 | ISO 8601 | |

Figure8

Figure8 is the result code from figure7.

| Library(sas7bdat) ts<-read.sas7bdat("E:\\users\\tiany\\ts2.sas7bdat", NULL) | 1 |
|--|---|
| ##another way## | |
| Library(haven) | 2 |
| <pre>ts<-read_sas("E:\\users\\tiany\\ts2.sas7bdat", NULL)</pre> | |
| | |
| View(ts) | |

Figure9

Besides creating the Trial Summary dataset using data.frame funcition in RStudio, there are two other methods of importing SAS datasets into R directly.

1: if we invoke "SASxport" package, then we should use read.sas7bdat function to import SAS datasets into R.

2: If we invoke "Haven" package, then we should use read_sas function to read SAS datasets into R;

Notice, the "haven" package can read both uncompressed and compressed SAS datasets, however, "sas7bdat" package can only read uncompressed SAS datasets.

Finally, we use View function to look at the whole Trial Summary dataset in RStudio.

| STUDYID [‡] | DOMAIN [÷] | TSSEQ [‡] | TSGRPID [‡] | TSPARMCD [‡] | † | †SVAL [‡] | TSVALNF [‡] | TSVALCD [‡] | TSVCDREF [‡] | TSVCDVER [‡] |
|----------------------|---------------------|--------------------|----------------------|-----------------------|---------------------------|--------------------|----------------------|----------------------|-----------------------|-----------------------|
| 001 | TS | 1 | | ACTSUB | Actual Number of Subjects | 10 | | | | |
| 001 | TS | 1 | | ADAPT | Adaptive Design | N | | C49487 | CDISC | 2019-12-20 |
| 001 | TS | 1 | | AGEMAX | Maximum Age of Subjects | P65Y | | | ISO 8601 | |
| 001 | TS | 1 | | AGEMIN | Minimum Age of Subjects | P18Y | | | ISO 8601 | |
| 001 | TS | 1 | | DCUTDESC | Data Cutoff Description | DATABASE LOCK | | | | |
| 001 | TS | 1 | | DCUTDTC | Data Cutoff Date | 2020-11-26 | | | | |
| 001 | TS | 1 | group1,drug1 | DOSE | Dose per Administration | 400 | | | | |
| 001 | TS | 2 | group2,drug2 | DOSE | Dose per Administration | 15 | | | | |
| 001 | TS | 1 | drug1 | DOSFRM | Dose Form | TABLET | | C42998 | ISO 8601 | |
| 001 | TS | 1 | drug2 | DOSFRM | Dose Form | CAPSULES | | C42998 | ISO 8601 | |

Figure10

Figure10 is the result code from Figure9, in addition:

Figure 8 and figure10 are outputs using data.frame function and read.sas7bdat/ read_sas function in RStudio are identical.

label(ts) <- 'Trial Summary'
label(ts\$STUDYID)<-'Study Identifier'
label(ts\$TSDMAIN)<-'Domain Abbreviation'
label(ts\$TSSEQ)<-'Sequence Number'
label(ts\$TSGRPID)<-'Group ID'
label(ts\$TSPARMCD)<-'Trial Summary Parameter Short Name'
label(ts\$TSPARM)<-'Trial Summary Parameter'
label(ts\$TSVAL)<-'Parameter Value'
label(ts\$TSVALNF)<-'Parameter Null Flavor'
label(ts\$TSVALCD)<-'Parameter Value Code'
label(ts\$TSVCDREF)<-'Name of the Reference Terminology'
label(ts\$TSVCDVER)<-'Version of the Reference Terminology</pre>

Figure11

Figure11 shows of using label function to assign labels for each variable.

2.2) EXPORT TS.R FILE WITH .XPT FORMAT

write.xport(ts,file="E:/users/tiany/ts_R.xpt")

Figure12

Finally, we use write.xport function to export TS file with xpt format. Figure12 is shown as above.

| This PC 🕨 | (E:) ► users ► tiany | | | | |
|-----------|----------------------|--------------------|-------------------|------|------|
| | Name | Date modified | Туре | Size | |
| | 💼 ts_R | 11/29/2020 2:20 PM | SAS Xport Transpo | • | 4 KB |

Figure 13

Figure13 is the ts.xpt file created by RStudio.

2.3) A FAST WAY TO CREATE ALL XPT FILES IN A DIRECTORY WITH RSTUDIO

| <pre>##import all files##</pre> | |
|--|---|
| Library(haven) | |
| <pre>setwd("E:\\Users\\Tiany")</pre> | 1 |
| <pre>ldf <- list()</pre> | 2 |
| listsas <- dir(pattern = "*.sas7bdat") | |
| <pre>for (k in 1:length(listsas)) {</pre> | |
| <pre>ldf[[k]] <- read_sas(listsas[k])</pre> | 3 |
| <pre>m<-substr(listsas[[k]],1,2)</pre> | |
| <pre>path<-paste("E:\\Users\\Tiany\\sas\\",m,"_new.xpt",sep="")</pre> | 4 |
| <pre>write_xpt(ldf[[k]],path)</pre> | 5 |
| } | |
| | |
| | |
| | |
| | |

Figure14

- 1: create a list
- 2: create the list of all the sas7bdat files in the directory
- 3: using for loop function to import all sas7bdat files from the SAS library to RStudio.

[1] "ae.sas7bdat" "cm.sas7bdat" "dm.sas7bdat" "ds.sas7bdat" "ts.sas7bdat"

Figure15

Figure15 shows the result of #3 in figure14.

4: concatenate the string and create the path where store all .xpt files

[1] "E:/users/tiany/sas/ae.xpt" "E:/users/tiany/sas/cm.xpt" "E:/users/tiany/sas/dm.xpt" "E:/users/tiany/sas/ds.xpt" [5] "E:/users/tiany/sas/ts.xpt"

Figure16 Figure16 shows the result of #4 in figure14

5: export all xpt files with write_xpt function.

This PC ► (E:) ► users ► tiany ► sas

| Name | Date modified | Туре | Size |
|--------|--------------------|-------------------|-------|
| 💼 ts_R | 12/15/2020 5:09 PM | SAS Xport Transpo | 22 KB |
| 💼 ds_R | 12/15/2020 5:09 PM | SAS Xport Transpo | 32 KB |
| 💼 dm_R | 12/15/2020 5:09 PM | SAS Xport Transpo | 17 KB |
| 💼 cm_R | 12/15/2020 5:09 PM | SAS Xport Transpo | 6 KB |
| 💼 ae_R | 12/15/2020 5:09 PM | SAS Xport Transpo | 20 KB |

Figure17

Figure17 is the result of code from #5 in figure15.

3) USING PYTHON TO GENERATE FINAL DATASETS IN XPT FORMAT

3.1) CODE TO GENERATE RAW TS DOMAIN IN PYTHON

Python is an interpreted, object-oriented, high-level programming language. In the pharmaceutical industry, there is a growing acceptance towards using Python to deal with data and to deliver programming outputs efficiently and effectively.

After installing python, open the CMD.exe command line interface, set the directory where python is installed and type in pip install xport, as shown on below Figure 18.



Figure18

Open the IDLE application, which is an integrated development environment for Python, then start to create a new file.





Figure19, we import two packages, "XPORT" and "PANDAS".

1: "XPORT" is a module for providing load function for reading data from a SAS file;

2: "PANDAS" is a standard for data analysis and management, it offers an easy way to import data, modulate variables, etc.

The following process shows how to import statement "pandas" with "as" to define a short name "pd".

| >>> ts frame= |
|---|
| pd.DataFrame(|
| {"STUDYID": |
| ["001","001","001","001","001","001","001","001","001","001"], |
| "DOMAIN": ["TS","TS","TS","TS","TS","TS","TS","TS" |
| "TSSEQ" : [1,1,1,1,1,1,2,1,1], |
| "TSGRPID": |
| ["","","","","","","","group1,drug1","group2,drug2","drug1","drug2"], |
| "TSPARMCD" :["ACTSUB","ADAPT","AGEMAX","AGEMIN","DCUTDESC", |
| "DCUTDTC", "DOSE","DOSFRM", "DOSFRM"], |
| "TSPARM" : ["Actual Number of Subjects", |
| "Adaptive Design", |
| "Maximum Age of Subjects", "Minimum Age of Subject, |
| "Data Cutoff Description","Data Cutoff Date", |
| "Dose per Administration", |
| "Dose per Administration", |
| "Dose Form", |
| "Dose Form"], |
| "TSVAL" : ["10","N","P65Y","P18Y","DATABASE LOCK", |
| "2020-11-26","400" ,"15","TABLET","CAPSULES"], |
| "TSVALNF" :["","","","","","","","","], |
| "TSVALCD" :["","C49487","","","","","","C42998","C42998"], |
| "TSVCDREF" :["","CDISC","ISO 8601","ISO 8601","","","","","ISO 8601","ISO |
| 8601"], |
| "TSVCDVER" :["","2019-12-20","","","", |
| "", "", "", "", ""], |
| } |
| |
| >>> pd.set_option("display.max_columns", None) |
| >>> pd.set_option("display.max_rows", None) |
| >>> ts_frame.head() |

Figure20

1: Showing how to use the "dataframe" function of "PANDAS" to create data frame columns and rows.

2: In order to show all columns and rows of ts_frame data, we use set_option function to display all columns and rows;

3: finally, use head function to show the TS data in Python.

| TSVCDVER | TSVCDREF | TSVALCD | TSVALNE | TSVAL | TSPARM | TSPARMCD | TSGRPID | TSSEQ | DOMAIN | STUDYID |
|------------|----------|---------|---------|---------------|---------------------------|----------|---------------|-------|--------|---------|
| | | | | 10 | Actual Number of Subjects | ACTSUB | | 1 | TS | 001 |
| 2019-12-20 | CDISC | C49487 | | N | Adaptive Design | ADAPT | | 1 | TS | 001 |
| | ISO 8601 | | | P65Y | Maximum Age of Subjects | AGEMAX | | 1 | TS | 001 |
| | ISO 8601 | | | P18Y | Minimum Age of Subjects | AGEMIN | | 1 | TS | 001 |
| | | | | DATABASE LOCK | Data Cutoff Description | DCUTDESC | | 1 | TS | 001 |
| | | | | 2020-11-26 | Data Cutoff Date | DCUTDTC | | 1 | TS | 001 |
| | | | | 400 | Dose per Administration | DOSE | group1, drug1 | 1 | TS | 001 |
| | | | | 15 | Dose per Administration | DOSE | group2, drug2 | 2 | TS | 001 |
| | ISO 8601 | C42998 | | TABLET | Dose Form | DOSFRM | drug1 | 1 | TS | 001 |
| | ISO 8601 | C42998 | | CAPSULES | Dose Form | DOSFRM | drug2 | 1 | TS | 001 |

Figure21

Figure21 is the output of code from Figure20 in Python. It is the same result with Figure2, the one created by SAS , and Figure10, the one created by RStudio.

3.2) EXPORT TS.PY FILE WITH .XPT FORMAT

>>> with open("E:/users/tiany/python file/ts_python.xpt","wb") as
f:xport.from_columns(ts_frame,f)

Figure22

In figure22, We use (with open) function to export ts_frame.py with ts_python.xpt format and store it to the location "E:/users/tiany/python file.

| This PC | (E:) ► users ► tiany ► python file | | | | | | | | | | |
|---------|------------------------------------|--------------------|-------------------|-------|--|--|--|--|--|--|--|
| | Name | Date modified | Туре | Size | | | | | | | |
| | ats_python | 11/30/2020 9:24 PM | SAS Xport Transpo | 18 KB | | | | | | | |

Figure 23

The result shows on figure23.

4)REVIEW AND COMPARE GENERATED SIMPLIFIED TS.XPT FILE IN SAS UNIVERSAL VIEWER

The SAS Universal Viewer is a replacement for the SAS System Viewer, we use SAS Universal Viewer to open the ts.xpt file that we created.

| | | ts.xpt | | | | | | | | | | | | | | |
|-----|------------|--------|------|----|----------|-----------|--------|--------|----------|--------------------------------------|--|--|--|--|--|--|
| Lib | rary Prope | rties | 55 | | | | | | | | | | | | | |
| N | ame | Obs | Vars | # | Variable | Туре | Length | Format | Informat | Label | | | | | | |
| 1 | S | 10 | 11 | 1 | STUDYID | Character | 20 | | | Study Identifier | | | | | | |
| | | | | 2 | DOMAIN | Character | 2 | | | Domain Abbreviation | | | | | | |
| | | | | 3 | TSSEQ | Numeric | 8 | | | Sequence Number | | | | | | |
| | | | | 4 | TSGRPID | Character | 40 | | | Group ID | | | | | | |
| | | | | 5 | TSPARMCD | Character | 8 | | | Trial Summary Parameter Short Name | | | | | | |
| | | | | 6 | TSPARM | Character | 40 | | | Trial Summary Parameter | | | | | | |
| | | | | 7 | TSVAL | Character | 200 | | | Parameter Value | | | | | | |
| | | | | 8 | TSVALNF | Character | 200 | | | Parameter Null Flavor | | | | | | |
| - | | | | 9 | TSVALCD | Character | 200 | | | Parameter Value Code | | | | | | |
| | | | | 10 | TSVCDREF | Character | 20 | | | Name of the Reference Terminology | | | | | | |
| | | | | 11 | TSVCDVER | Character | 10 | | | Version of the Reference Terminology | | | | | | |

Figure24

Figure24 is the view of ts.xpt file created by SAS.

| 8 | | | | | | | SAS Univ | ersal Viewer | r - [ts_r.xp | ot] | | |
|---------|------------|--------|------|------|----|----------|-----------|--------------|--------------|----------|--------------------------------------|-------|
| File | Tools | Window | Help | | | | | | | | | _ 8 × |
| Addre | ss | | | | | | | | | | | ~ |
| Library | Properties | s | | | | | | | | | | |
| Name | | | Obs | Vars | # | Variable | Туре | Length | Format | Informat | Label | |
| TS | | | 10 | 11 | 1 | STUDYID | Character | 3 | | | Study Identifier | |
| | | | | | 2 | DOMAIN | Character | 2 | | | Domain Abbreviation | |
| | | | | | 3 | TSSEQ | Numeric | 8 | | | Sequence Number | |
| | | | | | 4 | TSGRPID | Character | 12 | | | Group ID | |
| | | | | | 5 | TSPARM | Character | 8 | | | Trial Summary Parameter Short Name | |
| | | | | | 6 | TSPARM | Character | 25 | | | Trial Summary Parameter | |
| | | | | | 7 | TSVAL | Character | 13 | | | Parameter Value | |
| | | | | | 8 | TSVALNF | Character | 1 | | | Parameter Null Flavor | |
| | | | | | 9 | TSVALCD | Character | 6 | | | Parameter Value Code | |
| | | | | | 10 | TSVCDR | Character | 8 | | | Name of the Reference Terminology | |
| | | | | | 11 | TSVCDV | Character | 10 | | | Version of the Reference Terminology | |

Figure25

Figure25 is the view of ts.xpt file created by RStudio

| 8 | | | | | | SAS Univ | versal Vie | wer - [t | ts_python.x | pt] | _ 🗆 🗙 |
|---------|--------------|------|------|----|----------|-----------|------------|----------|-------------|------|-------|
| | | | | | | | | | | | - 8 × |
| File | Tools Window | Help | | | | | | | | | |
| Addres | s | | | | | | | | | | ~ |
| Library | Properties | | | | | | | | | | |
| Name | | Obs | Vars | # | Variable | Туре | Length | ormat | Informat | abel | |
| | | 10 | 11 | 1 | STUDYID | Character | 3 | | | | |
| | | | | 2 | DOMAIN | Character | 2 | | | | |
| | | | | 3 | TSSEQ | Numeric | 8 | | | | |
| | | | | 4 | TSGRPID | Character | 12 | | | | |
| | | | | 5 | TSPARMCD | Character | 8 | | | | |
| | | | | 6 | TSPARM | Character | 25 | | | | |
| | | | | 7 | TSVAL | Character | 13 | | | | |
| | | | | 8 | TSVALNF | Character | 1 | | | | |
| | | | | 9 | TSVALCD | Character | 6 | | | | |
| | | | | 10 | TSVCDREF | Character | 8 | | | | |
| | | | | 11 | TSVCDVER | Character | 10 | | | | |

Figure26

Figure26 is the view of ts.xpt file created by Python.

While SAS, R and Python create ts.xpt file, we want to compare among them. Firstly, when we look at ts.xpt file generated by SAS, the length of variables is corresponding with the standard variables' length in SDTM specification document. SAS allows us to change the variables' length. However, in both R and Python, the variables' length depends on the length of elements in a vector. Secondly, when we look at the "label" column, both SAS and R can assign variables' labels, however, it does not work in Python because the Python Xport Module does not have advanced properties capabilities such as variable labels cannot be assigned. In conclusion, SAS and R include more advanced capabilities to edit .xpt files.

REFRENCE

CREATING SIMPLIFIED TS.XPT FILES https://www.fda.gov/media/132457/download

Study Data Technical Conformance Guide_v4.5 https://www.fda.gov/media/136460/download

Technical Rejection Criteria for Study Data https://www.fda.gov/media/100743/download

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