

# Auto Subgroup TLG Generation With R Shiny

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1. Background
2. Case Study
3. Discussion

# Background

# What Is A Subgroup

- As per the guideline of EMA/CHMP,

The term 'subgroup' will be used to refer to a subset of the clinical trial population defined by one or more intrinsic and extrinsic factors (see ICH-E5) of the patients under investigation, usually measured at baseline.

- As per guideline of China NMPA,

亚组人群（简称亚群）指总体人群中具有某些特征的一个子集，亚组是全人群中的一个子集，亚组通常由患者的一个或多个内在和/或外在因素（见 ICH E5）来定义，而且应具有一定的临床意义。

EMA/CHMP/539146/2013: Guideline on the investigation of subgroups in confirmatory clinical trials.

NMPA:药物临床试验亚组分析指导原则（试行）

# Typical List of Subgroup



**Region**



**Age**



**Sex**



**Race**



**Baseline severity  
measure**



**Clinical events in  
previous years**



**Baseline medication**



**Baseline blood  
biomarker**

# What Are Subgroup Analyses

- **Content**

- Splitting all the participant data into subgroups, to make comparisons between them
- Usually done for subsets of participants (e.g. males/females or different geographical locations)

- **Purpose**

- Investigating heterogeneous results, or to answer specific questions about particular patient groups, types of intervention or types of study

“Most trials report subgroup analysis (median = 4 subgroups)”

- Assmann SF, Lancet 2000; 355:1064-1069;

# Case Study

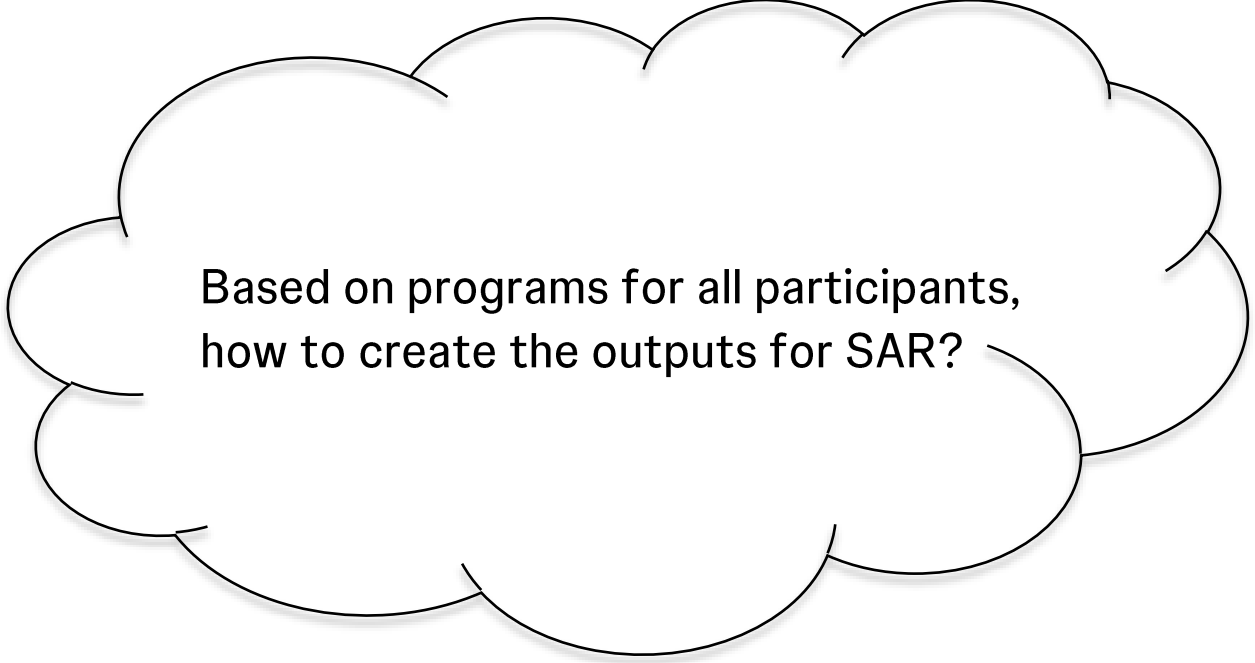
**A Subgroup Analysis in Support of  
NMPA Submission**



# Background

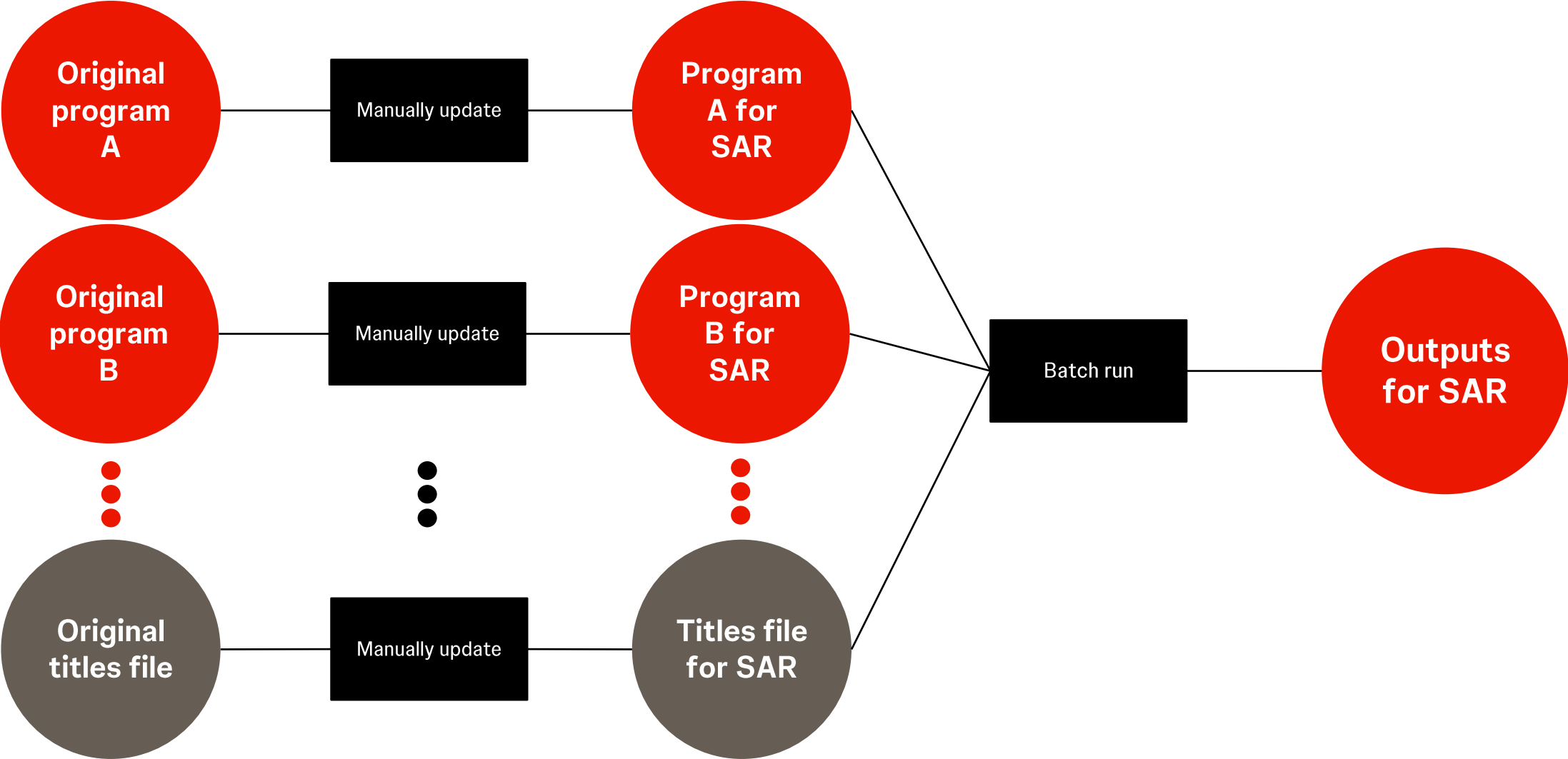
- A subgroup analysis report (SAR) is required to evaluate the consistency of results among all participants. This report includes safety and efficacy analyses conducted on specific sub-populations within a multi-regional clinical trial (MRCT).
- Global clinical study report (CSR) has been finalized for the MRCT, with the R programs to create outputs for all participants available.
- Sub-populations:
  - ✓ East Asia
  - ✓ China\*

\* China mainland



Based on programs for all participants,  
how to create the outputs for SAR?

# Regular Approach



# A Closer Look at The 'Manually Update'

Original program tsflab04

- tsflab01dc1.r
- tsflab01di.r
- tsflab01ec1.r
- tsflab01fc1.r
- tsflab01fccp.r
- tsflab04.r
- tsflab05.r
- tsflabrb05.r
- tsflabrb05cp.r
- tsflabrb06.r

### Code update:

- Update program header
- Filtering on the input dataset
- Change output id

```
# Program Name : tsflab04ea.r  
  
adsl <- read_sas(read_path(a_in,"adsl.sas7bdat")) %>%  
  filter(REGION == "East Asia")  
adlb <- read_sas(read_path(a_in,"adlb.sas7bdat")) %>%  
  filter(REGION == "East Asia")  
  
##define the ID of TLGs  
tblid<-"TSFALB04ea"
```



Next original program tsflab05

A sip of water

Program A for SAR

- tsflab01dc1.r
- tsflab01di.r
- tsflab01ec1.r
- tsflab01fc1.r
- tsflab01fccp.r
- tsflab04.r
- tsflab05.r
- tsflabrb05.r
- tsflabrb05cp.r
- tsflabrb06.r



Repeat it for China sub-population X2

# A Closer Look at The 'Manually Update'

Original titles file



A sip of water

89	TSFAE02	TITLE	Number of Subjects With Treatment-emergent Adverse Events by System Organ Class and Preferred Term; Safety Analysis Set (Study XXXXXXX)
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89	TSFAE02ea	TITLE	Number of Subjects With Treatment-emergent Adverse Events by System Organ Class and Preferred Term for East Asia Population; Safety Analysis Set (Study XXXXXXX)
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Titles file for SAR

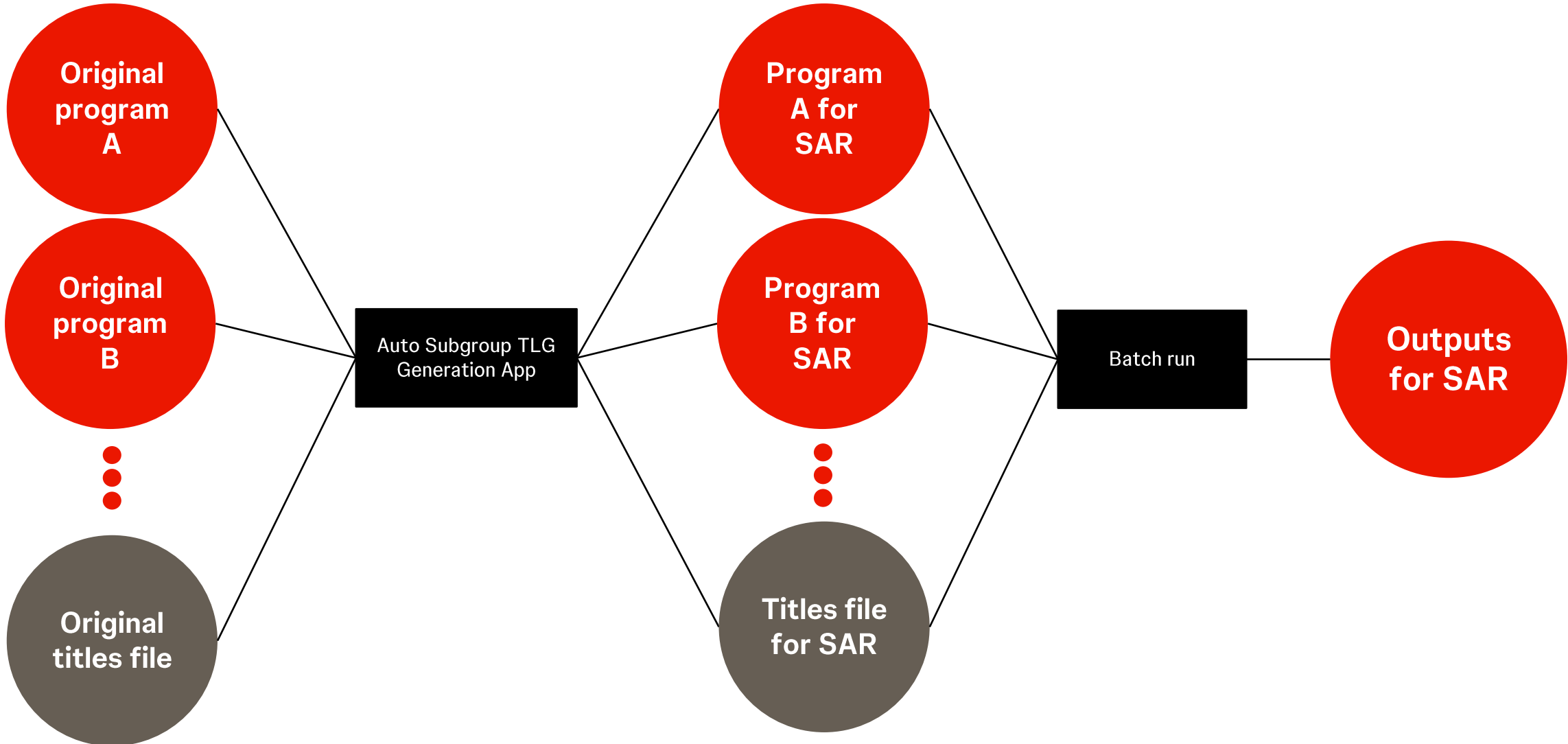
It's never too bad to absorb water and walk around, but is there a way to improve the process?

Customized user inputs

R programs processing in batch

**R Shiny!**

# New Workflow



# Auto Subgroup TLG Generation App

## Auto Subgroup TLG Generation

### 1. Choose programs you want to update

#### Select Level of Original Programs

Other Path

#### Programs

Alphabetic order

### 2. Update titles.xlsx based on your selections in step.1

#### Upload titles.xlsx

#### Add to title ID

Suggest: Please add only lowercase letters as suffix. Uppercase letters will be converted to lowercase and '-' will be removed when generating the rtf output

#### Add to title

Note: A new titles.xlsx will be created in your pdev input folder. Please avoid unconscious update

### 3. Update the programs

#### Add output ID from step 2

Other output ID

#### Group variable

#### Subgroup population



# 1. Select Programs for SAR

- shinyWidgets::pickerInput
  - Multiple choices
  - Search box

1. Choose programs you want to update

Select Level of Original Programs

PREPROD

Other Path

Programs

tsflab04.r, tsflab05.r, tsflabrb09.r, tsflabrb10.r

Select All

Deselect All

tsflab01cc1.r  
tsflab01ci.r  
tsflab01dc1.r  
tsflab01di.r  
tsflab01ec1.r  
tsflab01fc1.r  
tsflab01fccp.r  
tsflab04.r  
tsflab05.r  
tsflabrb05.r  
tsflabrb05cp.r  
tsflabrb06.r  
tsflabrb06cp.r  
tsflabrb08.r  
tsflabrb08cp.r  
tsflabrb09.r  
tsflabrb10.R  
tsflabrb10cp.r  
tsflabrb12i.r  
tsfvit01.r  
tsfvit04.r  
tsidem01saf.r



# 2. Update titles.xlsx File

## 2. Update titles.xlsx based on your selections in step.1

### Upload titles.xlsx

Browse... No file selected

### Add to title ID

ea|chn

Suggest: Please add only lowercase letters as suffix. Uppercase letters will be converted to lowercase and '-' will be removed when generating the rtf output

### Add to title

for East Asia Population|for China Populatio

Note: A new titles.xlsx will be created in your pdev input folder. Please avoid unconscious update

Create titles.xlsx

- Multiple subgroups at the same time

TSF AE03	TITLE	Number of Subjects With Treatment-emergent Serious Adverse Events by System Organ Class and Preferred Term; Safety Analysis Set (Study XXXXXXXX)
TSF AE03	FOOTNOTE1	Key: SAE = serious adverse event.
TSF AE03ae	TITLE	Number of Subjects With Treatment-emergent Serious Adverse Events by System Organ Class and Preferred Term for East Asia Population; Safety Analysis Set (Study XXXXXXXX)
TSF AE03ea	FOOTNOTE1	Key: SAE = serious adverse event.
TSF AE03chn	TITLE	Number of Subjects With Treatment-emergent Serious Adverse Events by System Organ Class and Preferred Term for China Population; Safety Analysis Set (Study XXXXXXXX)
TSF AE03chn	FOOTNOTE1	Key: SAE = serious adverse event.

# 3. Update Programs

- Previous inputs will be carried over
- Programs will be updated with the selected filtering criteria by simply clicking (no need to open up files/type codes/save files)
- Programs can be updated for multiple subgroups at the same time

```
## apply population filter  
subpop_subject_list <- read_sas(read_path(a_in, 'adsl.sas7bdat')) %>%  
  filter(COUNTRY %in% c('CHN', 'JPN', 'KOR')) %>%  
  pull(USUBJID)
```

```
filter(USUBJID %in% subpop_subject_list)
```

3. Update the programs

Add output ID from step 2  
ea

Other output ID

Group variable  
COUNTRY

Subgroup population  
CHN, JPN, KOR

Select All	Deselect All
COUNTRY	
	CHN ✓
	JPN ✓
	FRA
	RUS
	CAN
	TUR
	USA
	GBR
	KOR ✓
	DEU

3. Update the programs

Add output ID from step 2  
ea

Other output ID

Group variable  
COUNTRY

Subgroup population  
CHN, JPN, KOR

Confirm Undo

ID	Var	Selection
ea	COUNTRY	CHN, JPN, KOR
chn	COUNTRY	CHN

Create programs

# Outputs for SAR

TSIDEM01: Demographics and Baseline Characteristics; Safety Analysis Set (Study XXXXXX)				
	Dummy Treatment A	Dummy Treatment B	Dummy Treatment C	Total
Analysis set: Safety Analysis				
Set	27	8	19	54
Sex				
N	27	8	19	54
Female	15 (55.6%)	3 (37.5%)	8 (42.1%)	26 (48.1%)
Male	12 (44.4%)	5 (62.5%)	11 (57.9%)	28 (51.9%)
Age, years				
N	27	8	19	54
Mean (SD)	35.6 (9.95)	27.3 (9.95)	33.6 (11.15)	33.6 (10.58)
SE	1.92	3.52	2.56	1.44
Median	37.0	24.0	30.0	32.0
Range	(18; 53)	(18; 49)	(18; 55)	(18; 55)
18 - <=30 years	9 (33.3%)	6 (75.0%)	10 (52.6%)	25 (46.3%)
>30 - <=45 years	14 (51.9%)	1 (12.5%)	6 (31.6%)	21 (38.9%)
>45 - <=55 years	4 (14.8%)	1 (12.5%)	3 (15.8%)	8 (14.8%)

TSIDEM01ea: Demographics and Baseline Characteristics for East Asia Population; Safety Analysis Set (Study XXXXXX)				
	Dummy Treatment A	Dummy Treatment B	Dummy Treatment C	Total
Analysis set: Safety Analysis				
Set	10	2	11	23
Sex				
N	10	2	11	23
Female	8 (80.0%)	1 (50.0%)	5 (45.5%)	14 (60.9%)
Male	2 (20.0%)	1 (50.0%)	6 (54.5%)	9 (39.1%)
Age, years				
N	10	2	11	23
Mean (SD)	34.1 (11.00)	24.0 (1.41)	34.7 (12.19)	33.5 (11.24)
SE	3.48	1.00	3.68	2.34
Median	34.0	24.0	32.0	30.0
Range	(18; 50)	(23; 25)	(18; 55)	(18; 55)
18 - <=30 years	5 (50.0%)	2 (100.0%)	5 (45.5%)	12 (52.2%)
>30 - <=45 years	3 (30.0%)	0	4 (36.4%)	7 (30.4%)
>45 - <=55 years	2 (20.0%)	0	2 (18.2%)	4 (17.4%)

TSIDEM01chn: Demographics and Baseline Characteristics for China Population; Safety Analysis Set (Study XXXXXX)				
	Dummy Treatment A	Dummy Treatment B	Dummy Treatment C	Total
Analysis set: Safety Analysis				
Set	7	2	2	11
Sex				
N	7	2	2	11
Female	5 (71.4%)	1 (50.0%)	0	6 (54.5%)
Male	2 (28.6%)	1 (50.0%)	2 (100.0%)	5 (45.5%)
Age, years				
N	7	2	2	11
Mean (SD)	36.4 (11.25)	24.0 (1.41)	21.5 (4.95)	31.5 (11.26)
SE	4.25	1.00	3.50	3.40
Median	38.0	24.0	21.5	25.0
Range	(23; 50)	(23; 25)	(18; 25)	(18; 50)
18 - <=30 years	3 (42.9%)	2 (100.0%)	2 (100.0%)	7 (63.6%)
>30 - <=45 years	2 (28.6%)	0	0	2 (18.2%)
>45 - <=55 years	2 (28.6%)	0	0	2 (18.2%)

# Discussion

# Benefit



## Efficiency gained

With the introduction of automation, manual work is significantly minimized, reducing hours of labor to minutes.



## Less error-prone

It reduces human errors that often occur due to the tedious and repetitive nature of tasks.



## Accessibility

Once published online, a Shiny App is readily accessible as long as an internet connection is available.



## User-friendly interface

It allows for easy navigation and user input. This approach can be applied not only for regional subgroup analysis in China or Japan submissions but also for any other subgroup analysis (such as sex, age groups)

# Limitations and Looking Forward



It currently only works for subgroup analysis with the same layout. Side by side table is not applicable yet.



It cannot automatically handle the case where statistical analysis fails due to reduced sample size after data subsetting. Verification is still required.

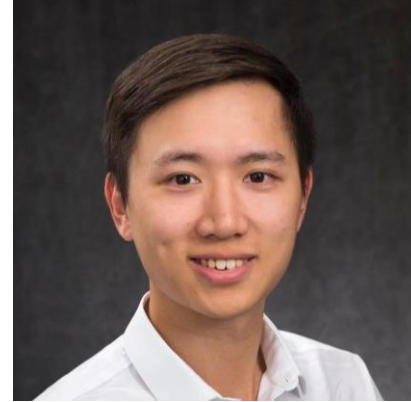


Moving forward, the use of natural language for updating programs offers a more generalized approach, showing great promise.

# Team Members



**Jiaqi Song**  
**SAS Macro**



**Yufan Chen**  
**R Shiny App**



# Acknowledge

Thanks for the support and valuable feedbacks from He Liu and Paul Jenkins!

# Thank you

If you have more questions, please contact:  
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