



# TFL Designer – Streamlining Analysis Results Generation

Bhavin Busa

German CDISC User Group Meeting

March 14, 2024

# **Current State: Analysis Results Deliverables**

Manual process in designing TFL shells/layout and ADaM specifications

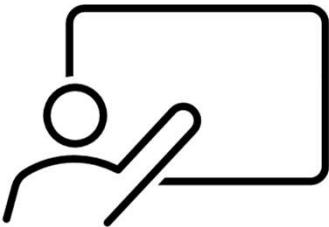
Programmer writes the SAS code to generate analysis deliverables (sometime with macros or re-using the code)

Too much variability across studies, disease areas, and organization

Static results with No or limited linking facility (e.g., to Protocol, SAP, ADaM data)

There is no industry standards for analysis results

## Poll Summary\*



\* Results from the live poll (*n=253 responders*) conducted during the TFL Designer Virtual Design Thinking Workshop (Part I), 13<sup>th</sup> Sep 2022, Bhavin Busa

### Who attended the workshop:

- Clinical / Statistical Programmer (63%)
- Biostatisticians (14%)
- Data Standards Expert (13%)
- Other (12%)

**74%** organization have TFL standards or templates

**40%** annotate their TFL mock-up shells to provide results metadata information

### Top 5 pain points:

1. Too much variability across studies / disease areas / organizations
2. No industry-wide standards exist
3. TFL metadata and shells are not machine-readable
4. Multiple manual steps in the process
5. Limited or no automation exist

Programming is more of a Science (50%) than it is an Art (39%)!



Who generates TFL shells (mock-ups)?  
**57%** - Biostatistician  
**31%** - Biostats & Programmers

**87%** responders confirmed their TFL shells are NOT machine-readable

**65%** responders uses MS Word / RTF for TFL shells generation

**76%** do not generate analysis results metadata prospectively to use in their TFL program

**82%** confirmed not having machine-readable TFL analysis results metadata

Out of the responders who use machine-readable ARM: MS Excel (14%) and SAS (10%) are top 2 format choices

**CDISC Analysis Result Standards – Releasing April 2024!**

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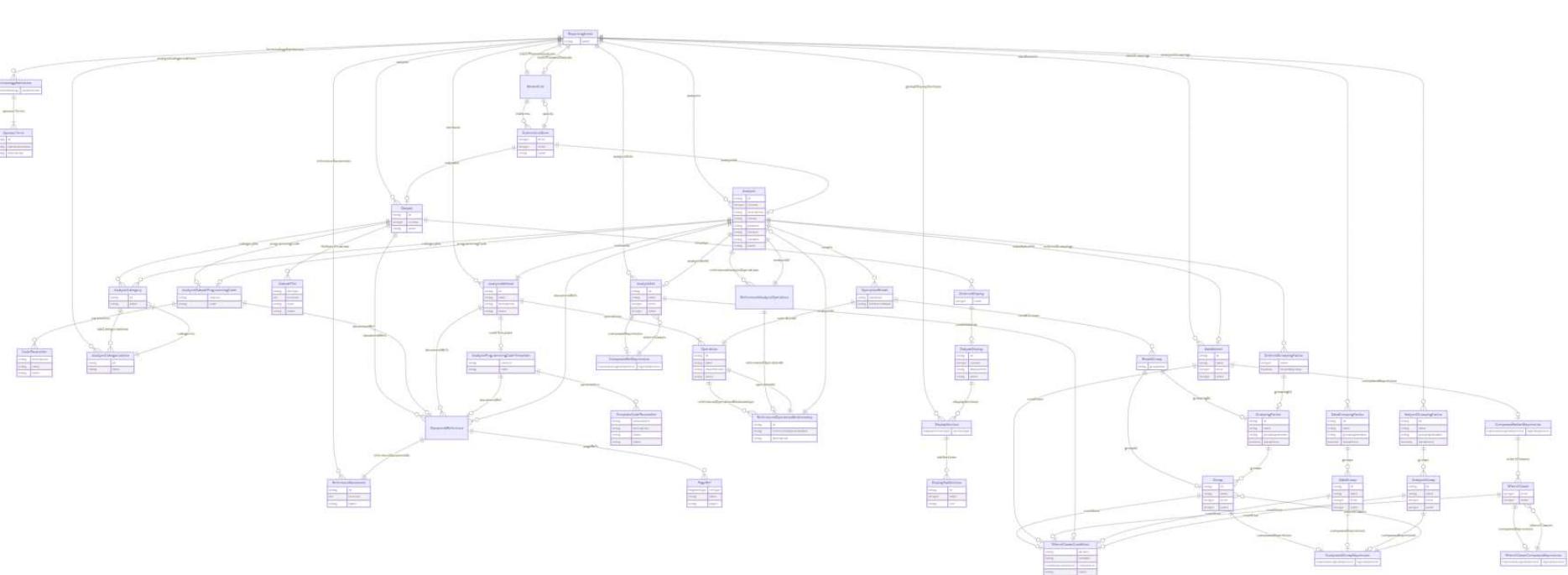


# DS04: Getting Started with the New CDISC Analysis Results Standard

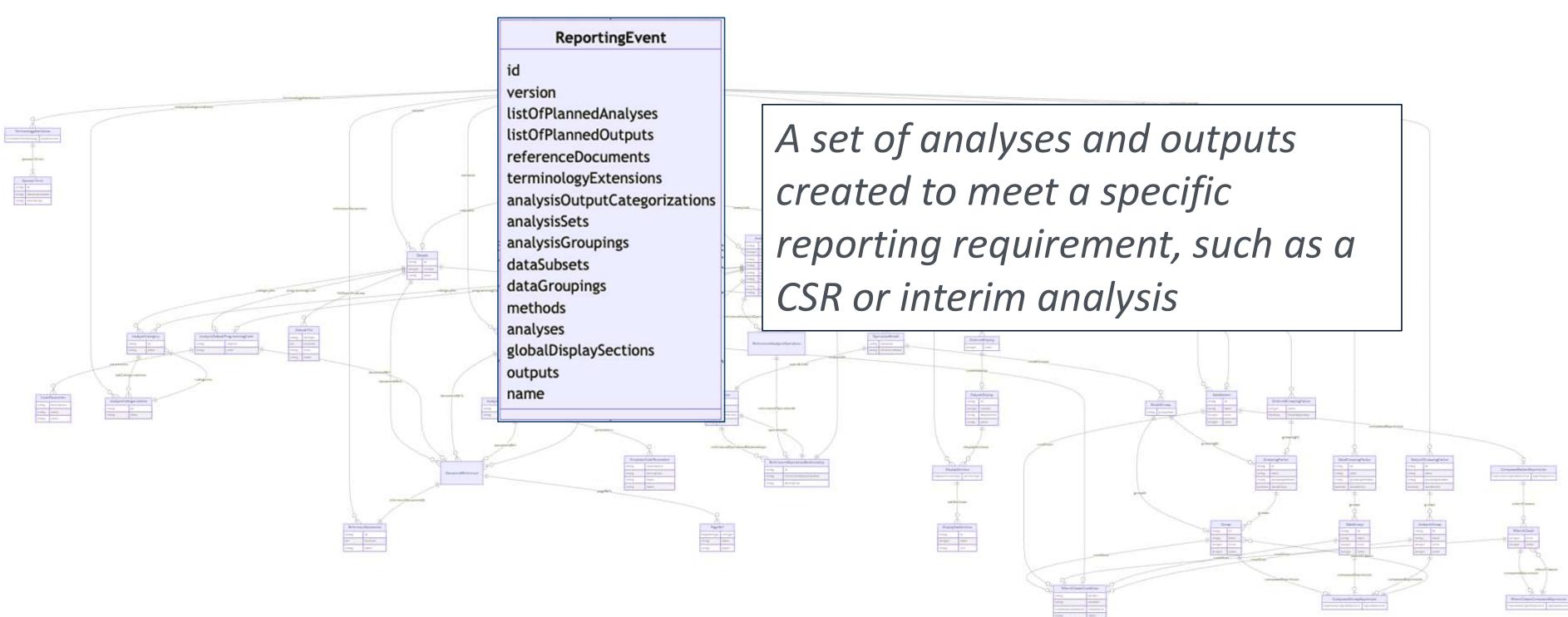
**Bess LeRoy, Head of Standards Innovation, CDISC**

**PHUSE US Connect 24**

# ARS Logical Model Schema Diagram



# ARS Logical Model Schema Diagram: Reporting Event



# Model Components

## Reporting Event

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

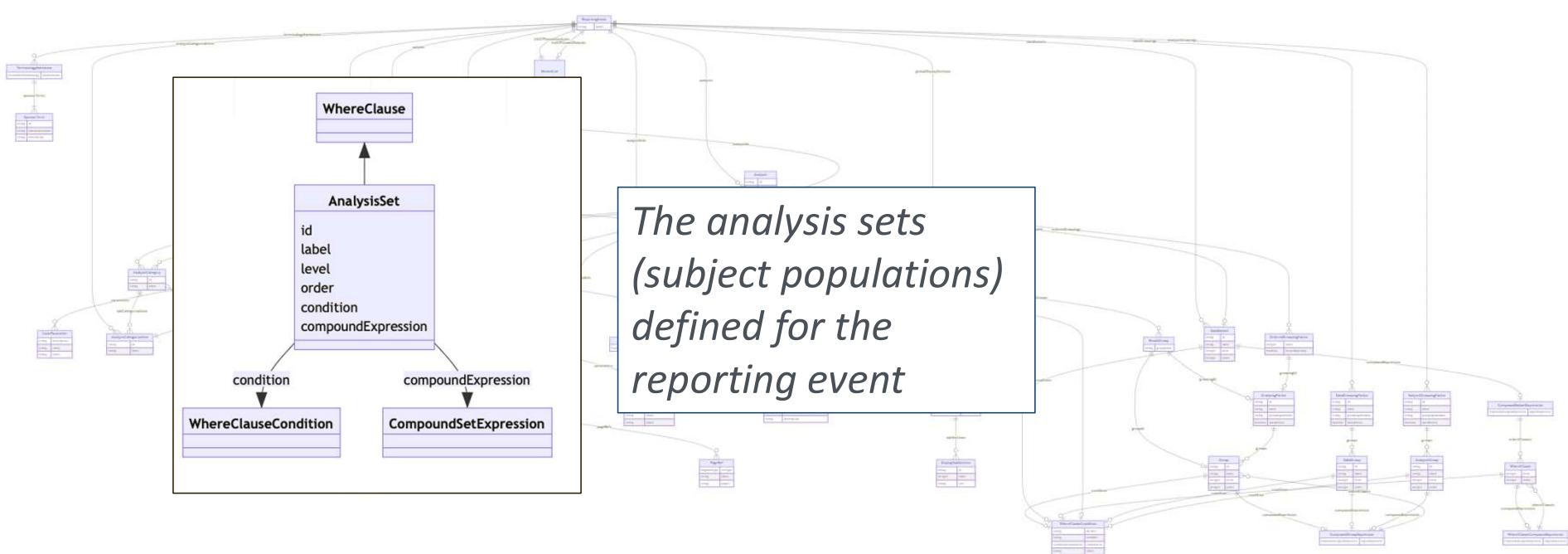
### Summary of TEAE by SOC and PT

Study - CDISC 360				Page x of y
System Organ Class Preferred Term [a], n (%)	Table 14.3.1.1 Summary of TEAE by System Organ Class and Preferred Term Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Notes: TEAE=Treatment-Emergent Adverse Events.  
Subjects are counted once within each system organ class and preferred term.  
[a] All investigators adverse events were coded using MedDRA version xx.x.

Source dataset: adae, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

# ARS Logical Model Schema Diagram: Analysis Set



# Model Components

## Analysis Set

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

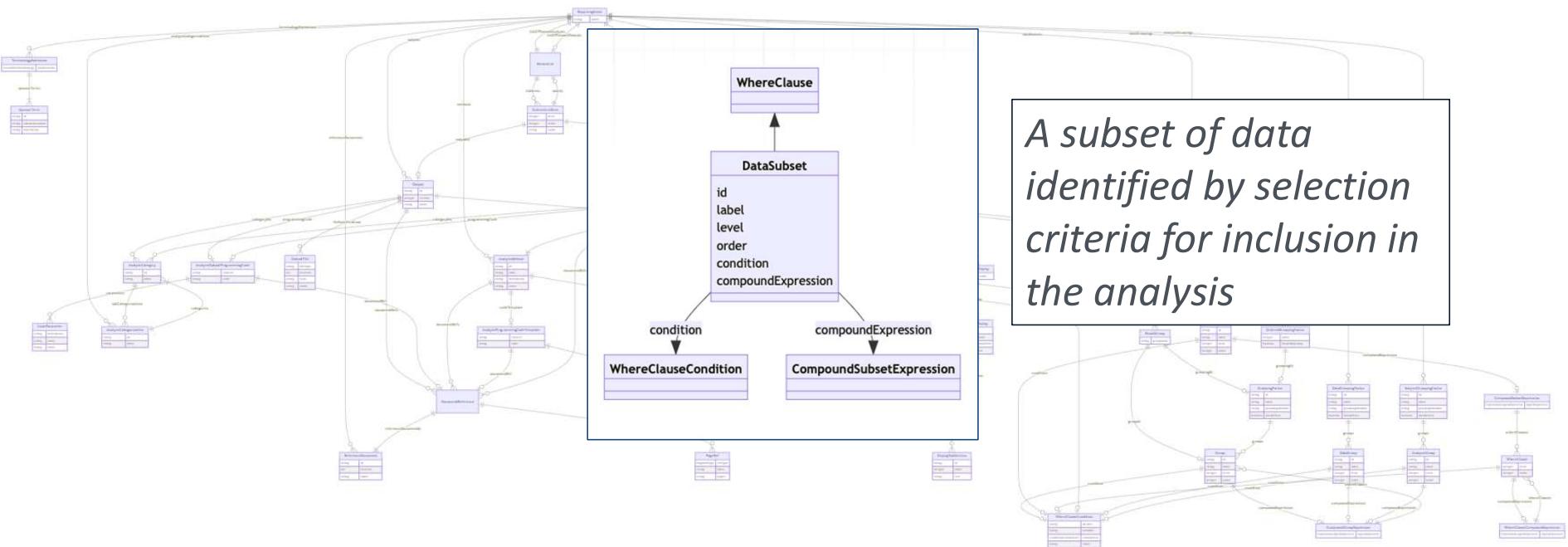
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Study - CDISC 360				Page x of y
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	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Notes: TEAE=Treatment-Emergent Adverse Events.  
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# ARS Logical Model Schema Diagram: Data Subset



# Model Components

## Data Subset

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

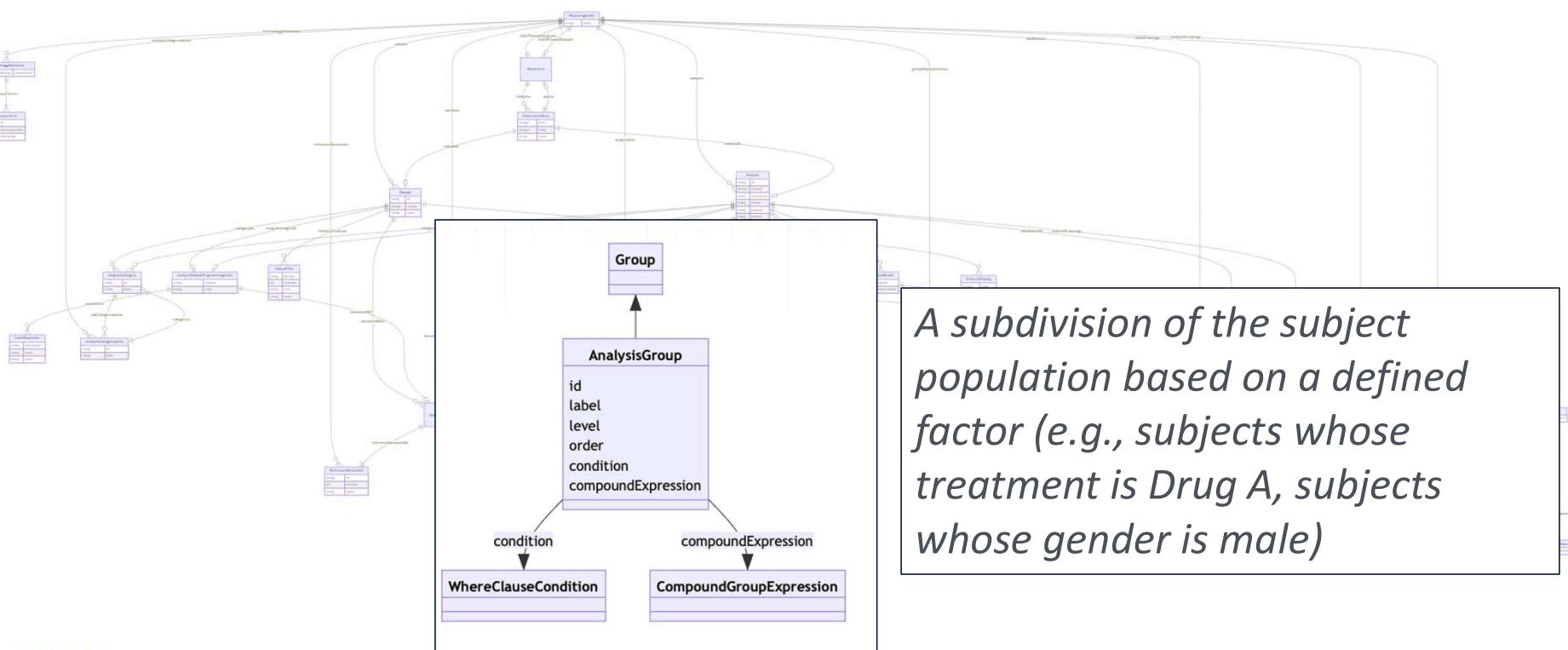
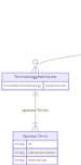
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System Organ Class Preferred Term [a], n (%)	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Notes: TEAE=Treatment-Emergent Adverse Events.  
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# ARS Logical Model Schema Diagram: Analysis Grouping



# Model Components

## Analysis Grouping

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

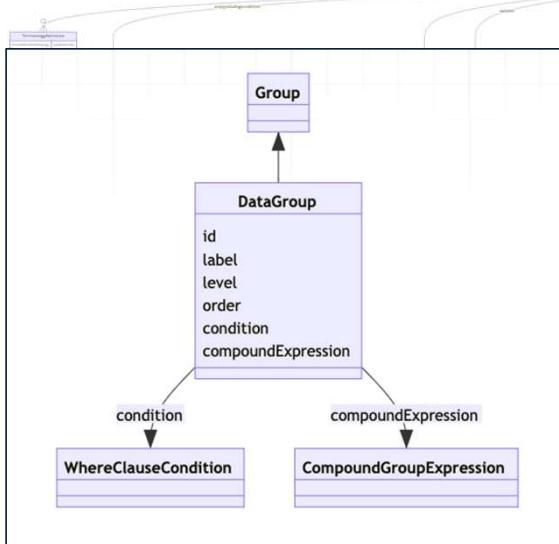
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Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

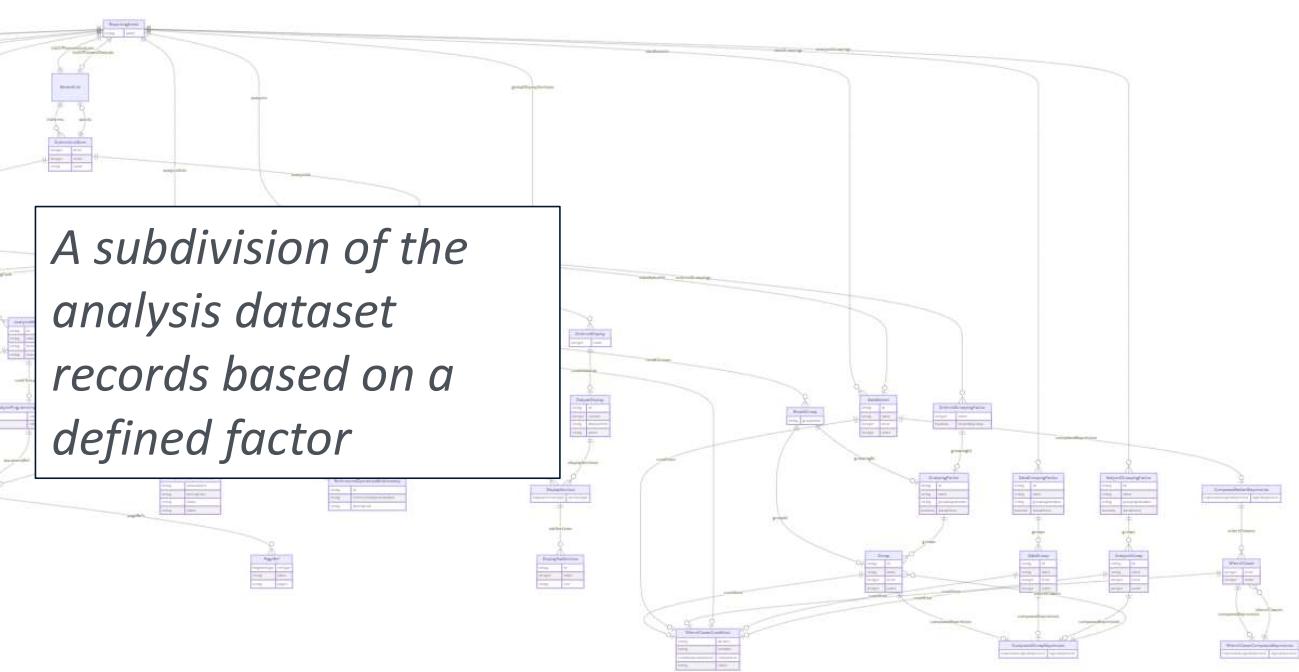
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Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

# ARS Logical Model Schema Diagram: Data Grouping



*A subdivision of the analysis dataset records based on a defined factor*



# Model Components

## Data Grouping

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
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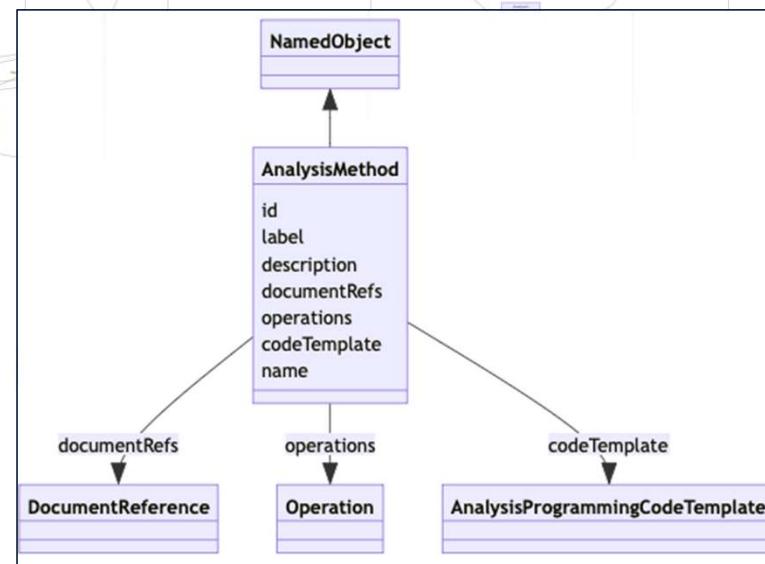
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Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Notes: TEAE=Treatment-Emergent Adverse Events.  
Subjects are counted once within each system organ class and preferred term.  
[a] All investigators adverse events were coded using MedDRA version xx.x.

Source dataset: adae, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

# ARS Logical Model Schema Diagram: Analysis Method



*A set of one or more statistical operations*

# Model Components

## Analysis Method

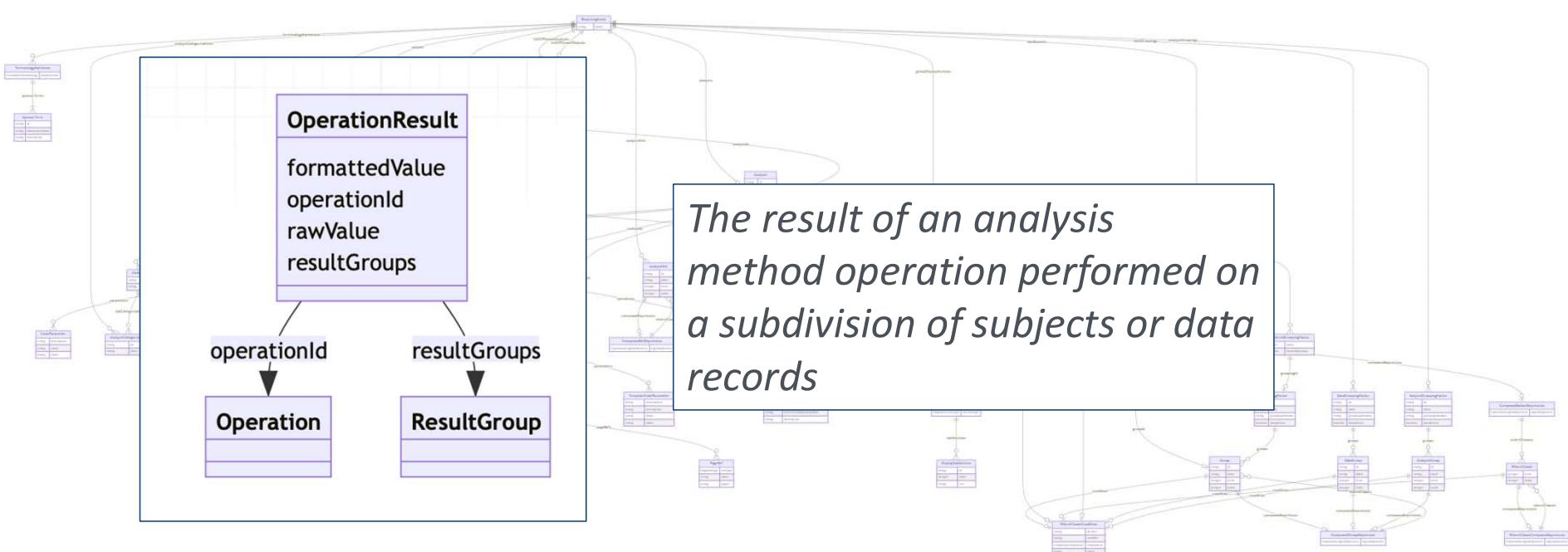
### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, [n (%)]				
< 65 years	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
≥ 65 years	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Gender, [n (%)]				
Male	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Female	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Ethnicity, [n (%)]				
Hispanic or Latino	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Not Hispanic or Latino	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Source dataset: adsl, Generated on: DDMONYYYY:HH:MM Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM				

### Summary of TEAE by SOC and PT

Study - CDISC 360				Page x of y
Table 14.3.1.1 Summary of TEAE by System Organ Class and Preferred Term Safety Population				
System Organ Class Preferred Term [a], [n (%)]	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Number of subjects with at least one event	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
<SOC 1>				
<Preferred Term 1>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
...	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
<Preferred Term n>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
<SOC 2>				
<Preferred Term 1>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
...	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
<Preferred Term n>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)	
Notes: TEAE=Treatment-Emergent Adverse Events. Subjects are counted once within each system organ class and preferred term. [a] All investigators adverse events were coded using MedDRA version xx.x.				
Source dataset: adae, Generated on: DDMONYYYY:HH:MM Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM				

# ARS Logical Model Schema Diagram: Results



# Model Components

## Results

### Summary of Demographics

Study - CDISC 360				Page x of y
Characteristics	Table 14.1.1 Summary of Demographics Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Age (years)				
n	XX	XX	XX	
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	
Median	XX.X	XX.X	XX.X	
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	
Min, Max	XX, XX	XX, XX	XX, XX	
Age Group, n (%)				
< 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
≥ 65 years	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Gender, n (%)				
Male	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Female	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Ethnicity, n (%)				
Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	
Not Hispanic or Latino	XX (XX.X)	XX (XX.X)	XX (XX.X)	

Source dataset: adsl, Generated on: DDMONYYYY:HH:MM  
Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

### Summary of TEAE by SOC and PT

Study - CDISC 360				Page x of y
System Organ Class Preferred Term [a], n (%)	Table 14.3.1.1 Summary of TEAE by System Organ Class and Preferred Term Safety Population			
	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)	
Number of subjects with at least one event	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 1>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<SOC 2>				
<Preferred Term 1>	XX (XX.X)	XX (XX.X)	XX (XX.X)	
...	XX (XX.X)	XX (XX.X)	XX (XX.X)	
<Preferred Term n>	XX (XX.X)	XX (XX.X)	XX (XX.X)	

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Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

# Creating Analysis Results Metadata: JSON

Table 2. Baseline Demographic and Clinical Characteristics, Safety Population, Pooled Analyses (or Trial X)					
Characteristic	Drug Name Dosage X N = XXX n (%)	Drug Name Dosage Y N = XXX n (%)	Placebo N = XXX n (%)	Active Control N = XXX n (%)	Total N = XXX n (%)
Sex, n (%)	n (%)	n (%)	n (%)	n (%)	n (%)
Male	n (%)	n (%)	n (%)	n (%)	n (%)
Female	n (%)	n (%)	n (%)	n (%)	n (%)
Age, years	XX (Y.Y) X.X (Y.Y) Median (min, max)	XX (Y.Y) X.X (Y.Y) XX (Y.Y, Z.Z)	XX (Y.Y) X.X (Y.Y) XX (Y.Y, Z.Z)	XX (Y.Y) X.X (Y.Y) XX (Y.Y, Z.Z)	XX (Y.Y) X.X (Y.Y) XX (Y.Y, Z.Z)
Age groups (years), n (%)	n (%)	n (%)	n (%)	n (%)	n (%)
≥17 to <65	n (%)	n (%)	n (%)	n (%)	n (%)
≥65	n (%)	n (%)	n (%)	n (%)	n (%)
≥65 to <75	n (%)	n (%)	n (%)	n (%)	n (%)
≥75	n (%)	n (%)	n (%)	n (%)	n (%)
Race, n (%)	n (%)	n (%)	n (%)	n (%)	n (%)
American Indian or Alaska Native Asian	n (%)	n (%)	n (%)	n (%)	n (%)
Black or African American	n (%)	n (%)	n (%)	n (%)	n (%)
Native Hawaiian or Other Pacific Islander	n (%)	n (%)	n (%)	n (%)	n (%)
White	n (%)	n (%)	n (%)	n (%)	n (%)
Other	n (%)	n (%)	n (%)	n (%)	n (%)

Source: [Include Applicant source, datasets and/or software tools used].  
 1 Difference is shown between [treatment arms] (e.g., difference is shown between Drug Name dosage X vs. placebo).  
 Abbreviations: N, number of patients in treatment arm; n, number of patients with given characteristic; SD, standard deviation.

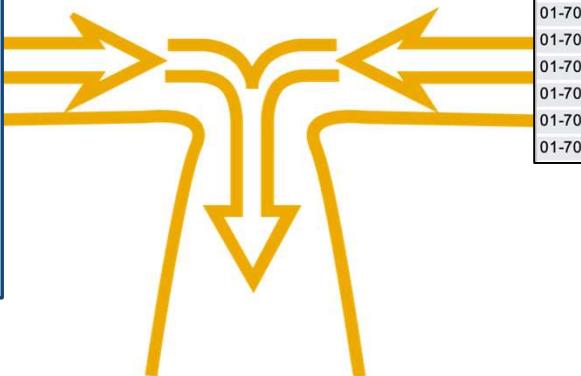


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{
  "name": "FDA Standard Safety Tables and Figures - Integrated Guide, Table 2",
  "id": "FDA_STF_T2",
  "listOfPlannedAnalyses": {
    "listItems": [
      {
        "name": "Table 2. Baseline Demographic and Clinical Characteristics, Safety Population, Trial CDISCPILOT01",
        "level": 1,
        "order": 1,
        "outputId": "O_FDA_STF_T2",
        "sublist": {
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              "level": 2,
              "order": 1,
              "analysisId": "A_SAF_CNT_USUBJID_TRT"
            },
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              "level": 2,
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              "analysisId": "A_SAF_CNT_USUBJID"
            },
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              "level": 2,
              "order": 3,
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                    "level": 3,
                    "order": 1,
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                    "level": 3,
                    "order": 2,
                    "analysisId": "A_SAF_SUM_USUBJID_SEX"
                  }
                ]
              }
            }
          ]
        }
      }
    ]
  }
}
```

# Leveraging ARS Metadata to Drive Results Automation

## ARS Metadata

{
"name": "FDA Standard Safety Tables and Figures - Integrated Guide, Table 2",
"id": "FDA_STF_T2",
"listItemType": "Table",
"listItemTypes": {
"listItems": [
{"name": "Table 2. Baseline Demographic and Clinical Characteristics, Safety Population, Trial CDISCPILOT01",
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"level": 2,
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},
{"name": "Summary of Subjects Total Population",
"level": 3,
"order": 2,
"analysisId": "A_SAF_SUM_USUBJID_SEX"
}
}
}
}



USUBJID	ARM	AGE	AGEGR1	AGEU	RACE	SEX
01-701-1015	Placebo	63	<65	YEARS	WHITE	F
01-701-1023	Placebo	64	<65	YEARS	WHITE	M
01-701-1028	Xanomeline High Dose	71	65+	YEARS	WHITE	M
01-701-1033	Xanomeline Low Dose	74	65+	YEARS	WHITE	M
01-701-1034	Xanomeline High Dose	77	65+	YEARS	WHITE	F
01-701-1047	Placebo	85	65+	YEARS	WHITE	F

id	operation_id	resultGroup1_groupingId	resultGroup1_groupid	resultGroup2_groupingId	resultGroup2_groupid	rawValue	formattedVal
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_1	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	14	14
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_1	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	72	72
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_2	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	8	8
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_2	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	76	76
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_3	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	11	11
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_1_n	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_3	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	73	73
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_1	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	16.27907	(16.3)
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_1	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	83.72093	(83.7)
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_2	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	9.52381	(9.5)
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_2	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	90.47619	(90.5)
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_3	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_1	13.09524	(13.1)
An03.02_AgeGrp_ByTrt	Mth01_CatVar_ByGrp_2_pct	AnlsGrouping_02_TrT	AnlsGrouping_02_TrT_3	AnlsGrouping_03_AgeGp	AnlsGrouping_03_AgeGp_2	86.90476	(86.9)

## Analysis Results Dataset

# Analysis Results Standard Model and User Guide

<https://cdisc-org.github.io/analysis-results-standard/>

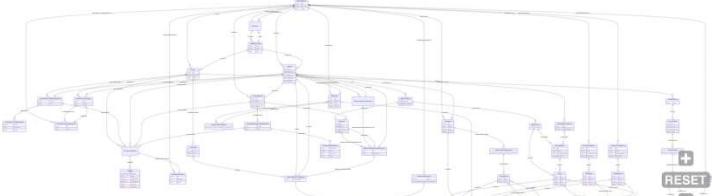
Analysis Results Standard (ARS)

Analysis Results Standard (ARS)

DRAFT Logical model to support both the prospective specification of analyses and the fully contextualized representation of the results of the analyses.

URI: <https://www.cdisc.org/ars/1-0> Name: ars\_ldm

Schema Diagram



Classes

Classes provide templates for organizing data. Data objects instantiate classes in the schema. Each class has a set of slots (aka fields, attributes) that are applicable to it. See [LinkML documentation](#) for more information.

Class	Description
NamedObject	An object with a name
ReportingEvent	A set of analyses and outputs created to meet a specific reporting requirement...
NestedList	A list of items (analyses or outputs) that may be organized within sub-lists

**cdisc**

Analysis Results Standard User Guide

Version 1.0 (Draft)

Prepared by the  
Analysis Results Standard Team

**Notes to Readers**

- This is the draft Version 1.0 of the Analysis Results Standard User Guide.
- This document is based on ADaM v2.1 and Analysis Results Metadata (ARM) v1.0 for Define-XML v2.0

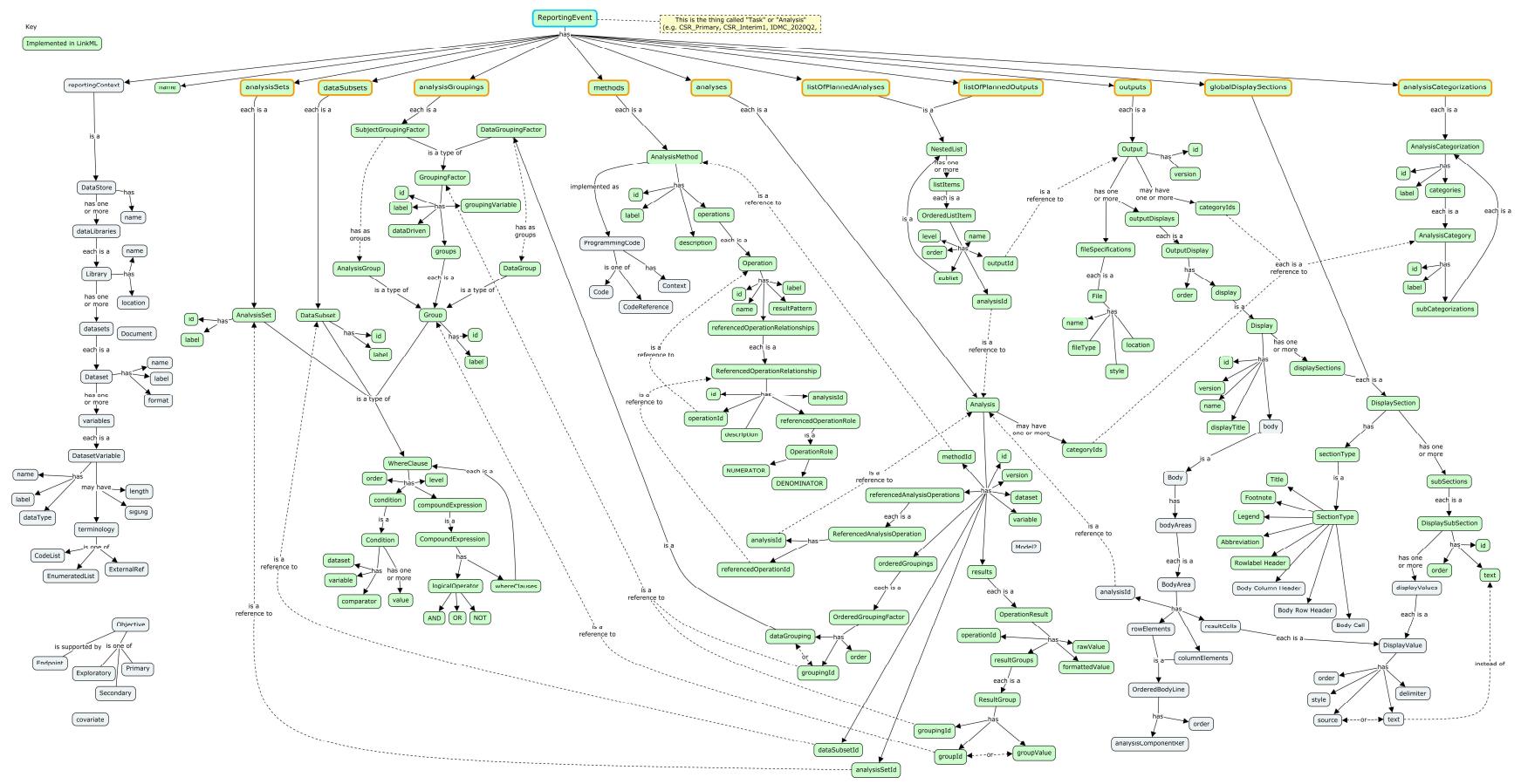
**Revision History**

Date	Version
2023-08-22	Internal Review Draft



<https://wiki.cdisc.org/display/ARSP/Analysis+Results+User+Guide>

# ARS Model Representation using CMAP

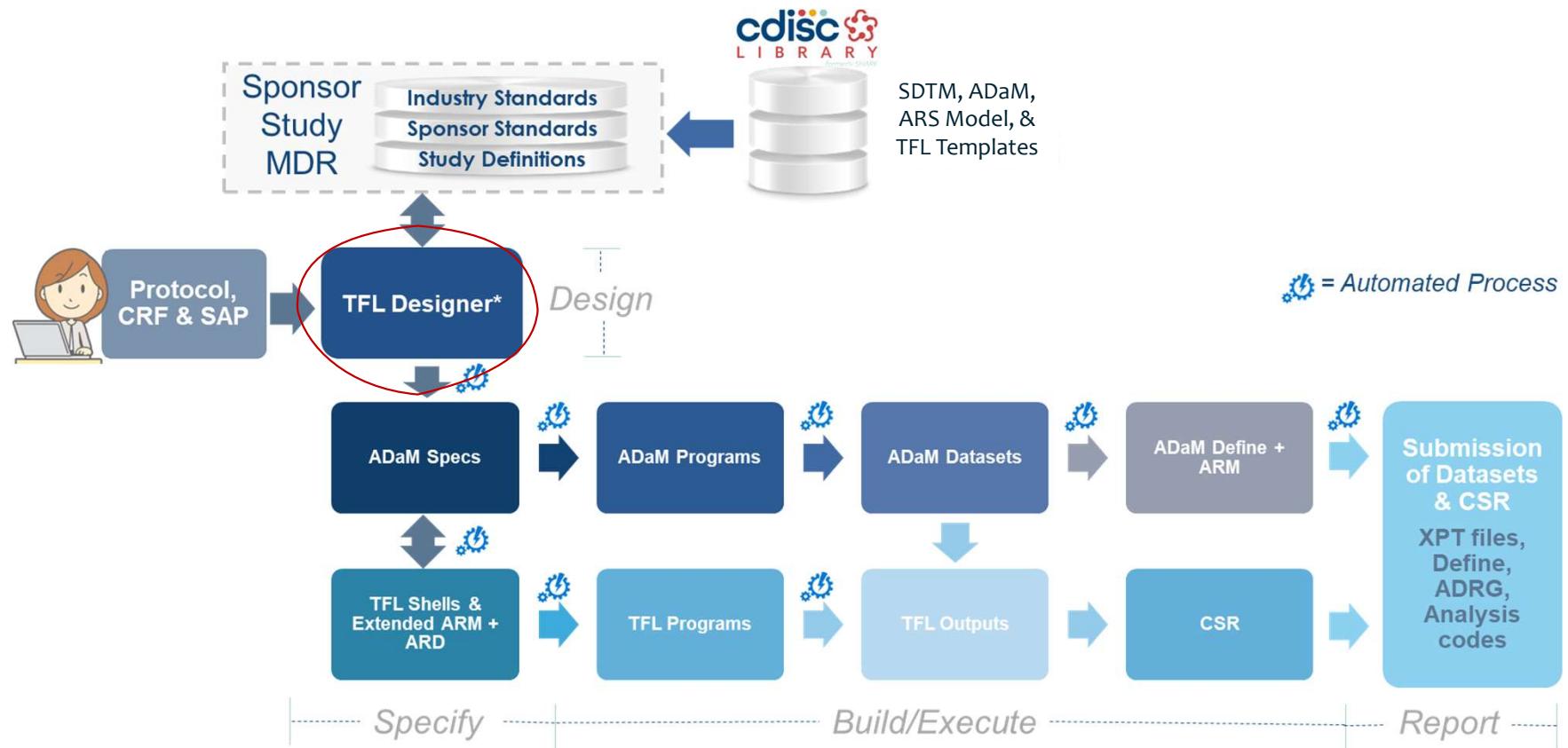


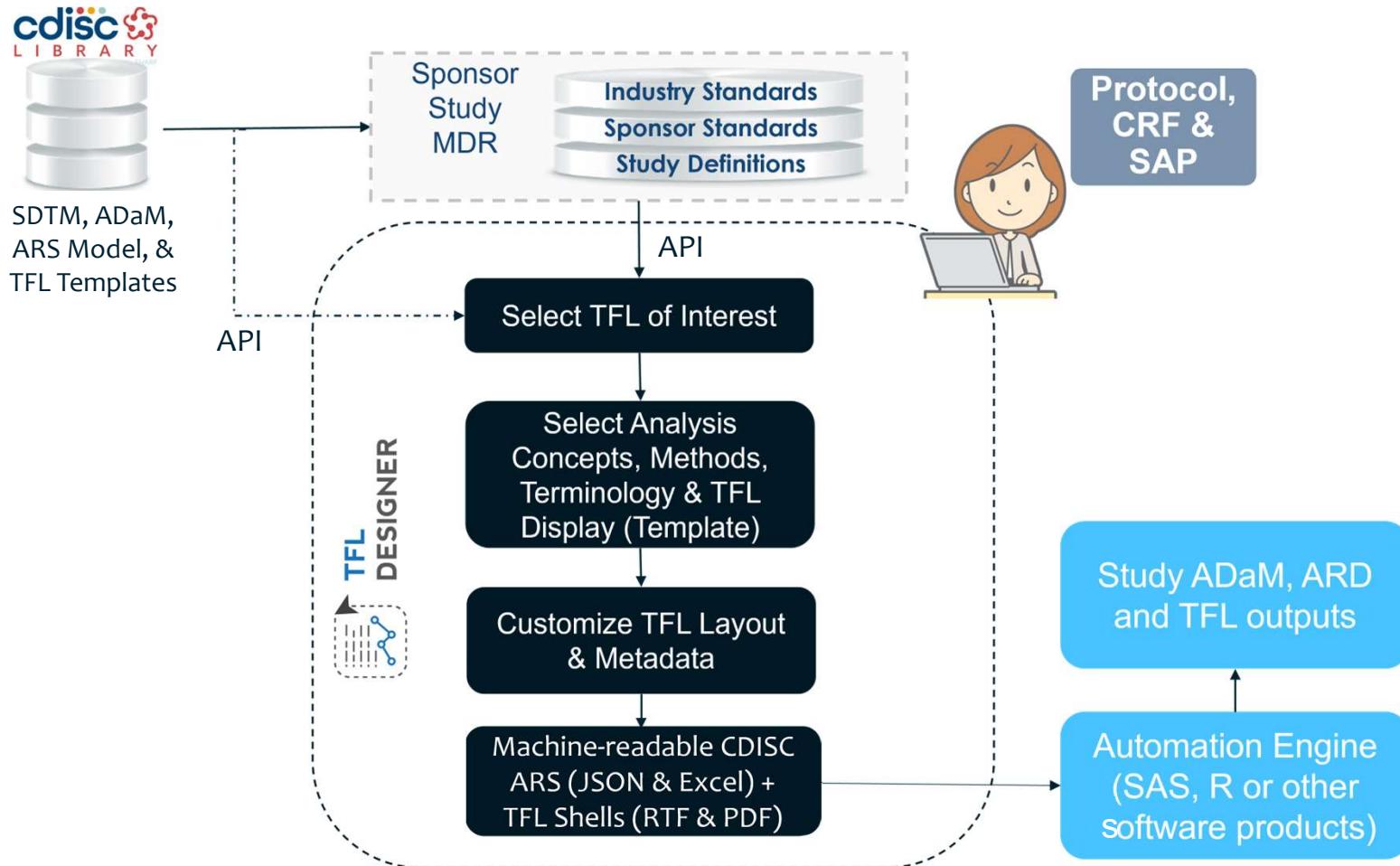
**ARS model is complex!**

**How do I operationalize it  
and generate analysis  
results metadata  
prospectively?**



# Analysis Results Workflow w/ TFL Designer







# TFL DESIGNER

## TFL Designer – Key Highlights

- Web-based solution
- Digitizes your analysis results (TFL)
- Aligned with CDISC Analysis Results Standards
- Central repository for your TFL standards, display templates, conventions and metadata
- Automates generation of TFL shells and provides machine-readable metadata
- Community & Enterprise versions



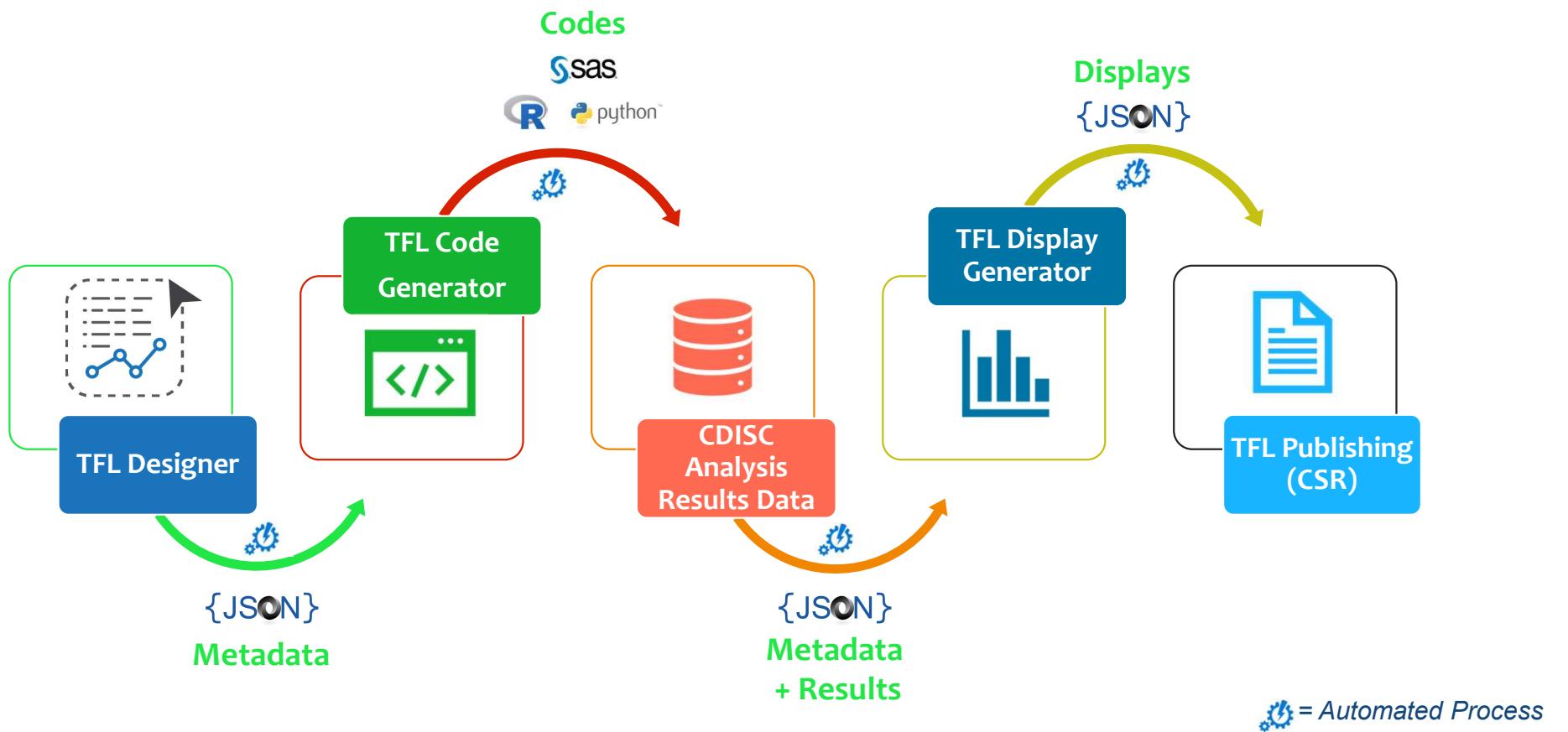
## Key Functionalities

- Central repository for your TFL standards/templates, conventions and metadata
- Access to library of TFL templates (community\* and user generated) by disease areas, TA, and indication
- Access to CDISC Standards (SDTM, ADaM, CT) via API to CDISC Library
- Develop new mock-up shells, edit/delete items
- Automatically populate items based on user inputs
- Export TFL shells in RTF & PDF formats
- Export analysis results metadata per the CDISC ARS model in JSON and Excel formats

\* including FDA STF-IG

[Will include PMDA, & PHUSE display templates in future updates]

# TFL / Analysis Results – Clymb's Development





## What are we trying to accomplish?

- Build an industry leading software solution that automates TFL design and generation process [community & enterprise versions]
- Partner with companies to improve their internal TFL standards and processes
- Quantify process improvements and continue to build future state automation (target 40-50% efficiency)
- Accelerate study timelines to allow your team to get data quicker
- CDISC 360 Vision: From PoC to Reality

# Live Demo

The screenshot shows a web browser window with the title "TFL Designer - Clymb Clinical". The URL in the address bar is "clymbclinical.com/tfl-designer/". The page itself has a dark background with green highlights. At the top left is the Clymb Clinical logo. A navigation bar at the top right includes links for Home, About Us, Solutions, Services, Careers, Resources, and a "Let's Connect!" button. Below the navigation is a large section titled "TFL Designer" with a sub-section titled "Why TFL Designer?". This section contains four items, each with an icon and text: 1. "Digitizes TFL analysis results" (calculator icon). 2. "Provides a centralized repository for TFL standards and templates" (document icon). 3. "Aligned with CDISC Analysis Results Standards and Model" (globe icon). 4. "Automates TFL shell generation and provides machine-readable metadata" (database icon). There is also a small circular icon with a stylized 'C' at the bottom left.

## TFL Designer

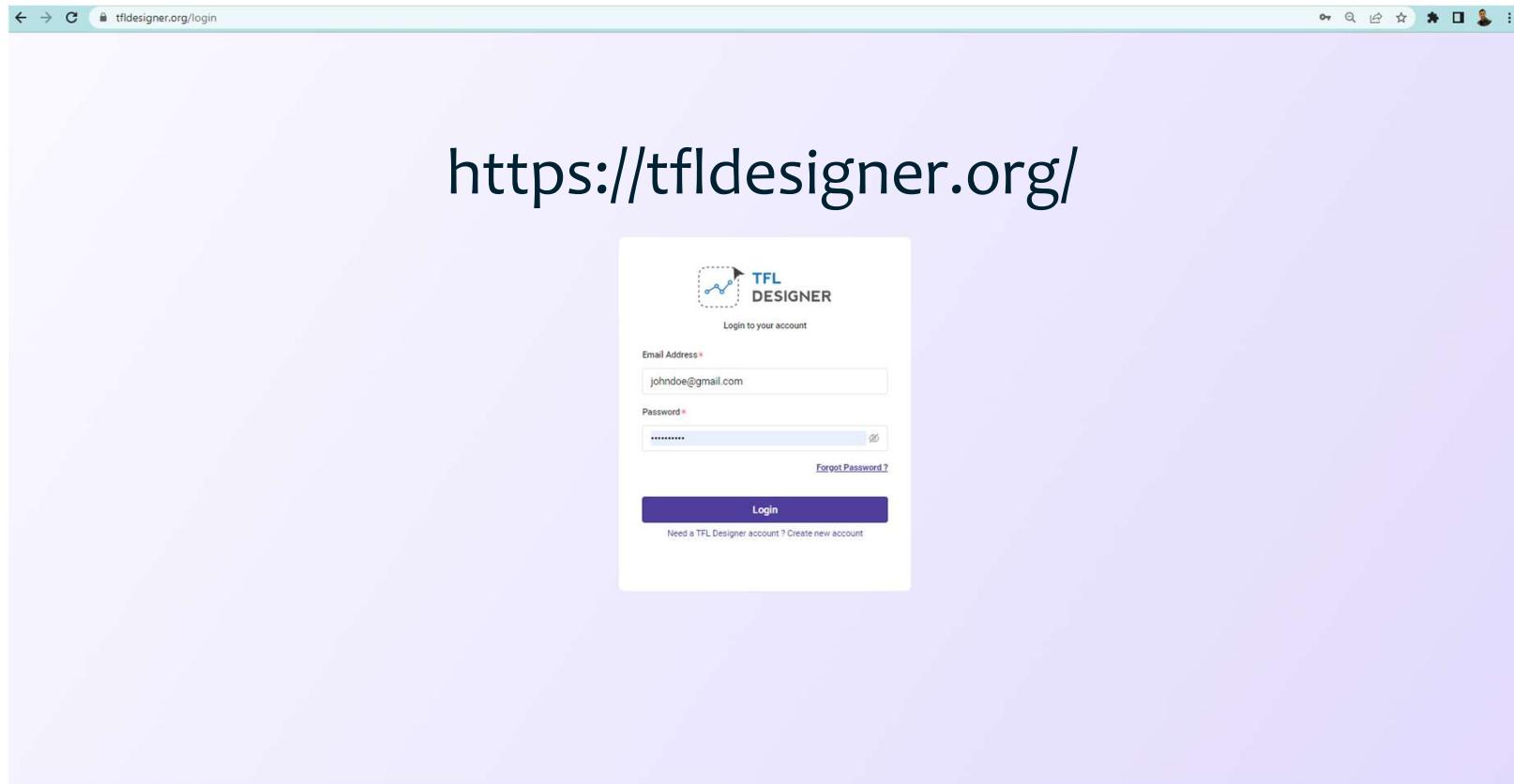
TFL Designer, available as both a Community and Enterprise version, is a leading Software as a Service (SaaS) solution that simplifies clinical trial reporting. This platform automates the creation of TFL shells and provides machine-readable metadata, which can then be seamlessly ingested for downstream automation in the programming of the TFLs. It digitizes analysis results, ensuring alignment with CDISC Analysis Results Standards (ARS), and offers a central repository for TFL standards, templates, conventions, and metadata.

Explore more

### Why TFL Designer?

- Digitizes TFL analysis results
- Provides a centralized repository for TFL standards and templates
- Aligned with CDISC Analysis Results Standards and Model
- Automates TFL shell generation and provides machine-readable metadata

# TFL Designer (Community version)



## Download files

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<http://bit.ly/3uKMAAv>

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