



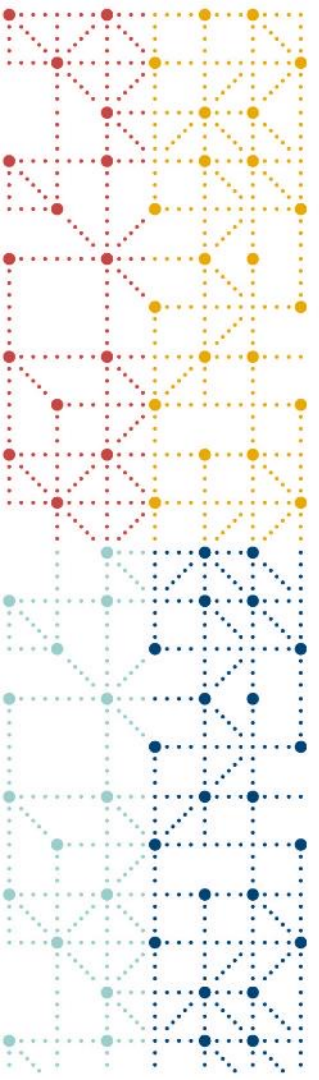
# Analysis Results Public Review Webinar

Bess LeRoy, Head of Standards Innovation, CDISC

Richard Marshall, Principal Data Modeler, CDISC

Bhavin Busa, Principal & Co-Founder, Clymb Clinical





# Agenda

- Project Background
- Overview of ARS Model and User Guide
- Next Steps
- Q&A

# CDISC Foundational Standards

Data Collection  
**CDASH**

Data Aggregation  
**SDTM**

Analysis  
**ADaM**

Results  
**???**

Table 4.2.2: HbA1c Longitudinal Repeated Measures Analysis Results Metadata	
Metadata Field	Metadata
DISPLAY IDENTIFIER	Table 4.2.1/Figure 4.2.1
DISPLAY NAME	Mean Change from Baseline in HbA1c (Percent) Longitudinal Repeated Measures Analysis, 24-Week Short-term Double-blind Treatment Period, Intention-to-treat Population
RESULT IDENTIFIER	Treatment difference results (LSMean, confidence interval, p-value)
PARAM	HbA1c (%)
PARAMCD	HBA1C
ANALYSIS VARIABLE	CHG (Change from baseline)
ANALYSIS REASON	SPECIFIED IN SAP
ANALYSIS PURPOSE	PRIMARY OUTCOME MEASURE
ANALYSIS DATASET	ADHBA1C

**ARM for Define.XML**

# Analysis Results Key Objectives

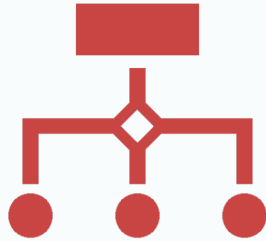


Leverage analysis results metadata to drive the automation of results



Support storage, access, processing, traceability and reproducibility of results

# Analysis Results Standards Key Results



Logical model that describes analysis results and associated metadata



User Guide to illustrate and exercise model with common safety displays

# Focus on Concepts, Not Layout

- Focus on concepts presented in data displays not on subjective layout and formatting of displays
- Representative displays therefore condense concepts
- For example, side-by-side Visit and Change-from-baseline summaries consolidates more concepts into an easy-to-read summary table

Parameter (Units) Visit	Treatment X (N=XX)	Treatment Y (N=XX)	Total (N=XX)
<Parameter 1> (<unit>)			
Baseline			
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
< Visit n >			
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
< Visit n Change from Baseline >			
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

Parameter (Units) Visit	Treatment X (N=XX)		Treatment Y (N=XX)		Total (N=XX)	
	Observed	CFB	Observed	CFB	Observed	CFB
<Parameter 1> (<unit>)						
Baseline						
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	
...						
<Visit n>						
n	XX	XX	XX	XX	XX	XX
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)	XX.X (XX.XX)
Median	XX.X	XX.X	XX.X	XX.X	XX.X	XX.X
Q1, Q3	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X	XX.X, XX.X
Min, Max	XX, XX	XX, XX	XX, XX	XX, XX	XX, XX	XX, XX

# FDA Standard Safety Tables and Figures: Integrated Guide



## STANDARD SAFETY TABLES AND FIGURES: *INTEGRATED GUIDE*

Center for Drug Evaluation and Research (CDER)  
Biomedical Informatics and Regulatory Review Science  
(BIRRS) Team

Please email [ONDbiomedicalinformatics@fda.hhs.gov](mailto:ONDbiomedicalinformatics@fda.hhs.gov) with any questions.

Version Date: August 2022

Table 2. Baseline Demographic and Clinical Characteristics, Safety Population, Pooled Analyses (or Trial X)

Characteristic	Drug Name	Drug Name	Placebo	Active Control	Total
	Dosage X	Dosage Y			
	N = XXX	N = XXX	N = XXX	N = XXX	Population
	n (%)	n (%)	n (%)	n (%)	N = XXX
	n (%)	n (%)	n (%)	n (%)	n (%)
<b>Sex, n (%)</b>	n (%)	n (%)	n (%)	n (%)	n (%)
Male	n (%)	n (%)	n (%)	n (%)	n (%)
Female	n (%)	n (%)	n (%)	n (%)	n (%)
<b>Age, years</b>	<b>X.X (Y.Y)</b>	<b>X.X (Y.Y)</b>	<b>X.X (Y.Y)</b>	<b>X.X (Y.Y)</b>	<b>X.X (Y.Y)</b>
Mean (SD)	X.X (Y.Y)	X.X (Y.Y)	X.X (Y.Y)	X.X (Y.Y)	X.X (Y.Y)
Median (min, max)	X.X (Y.Y, Z.Z)	X.X (Y.Y, Z.Z)	X.X (Y.Y, Z.Z)	X.X (Y.Y, Z.Z)	X.X (Y.Y, Z.Z)
<b>Age groups (years), n (%)</b>	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 (%)
<b>Race, n (%)</b>	n (%)	n (%)	n (%)	n (%)	n (%)
American Indian or Alaska Native	n (%)	n (%)	n (%)	n (%)	n (%)
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].

<sup>1</sup> 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

# Analysis Results Standard Model and User Guide

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

Analysis Results Standard (ARS) Search

**Analysis Results Standard (ARS)**

Schema Diagram

Classes

Slots

Enumerations

Types

Subsets

## 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\_idm

### 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
<a href="#">NamedObject</a>	An object with a name
<a href="#">ReportingEvent</a>	A set of analyses and outputs created to meet a specific reporting requiremen...
<a href="#">NestedList</a>	A list of items (analyses or outputs) that may be organized within sub-lists

## 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



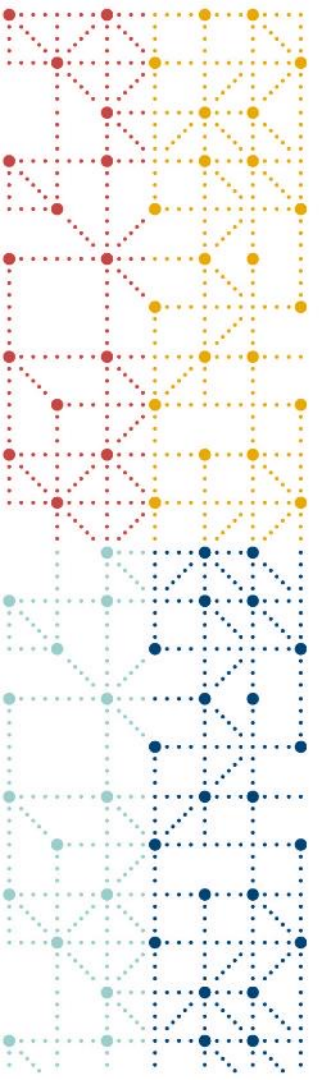


---

## Wait, What About ARM for Define.XML?

- At this point, no changes to ARM for Define.XML
- Retrospective documentation to aid in traceability
- Fills a specific regulatory need

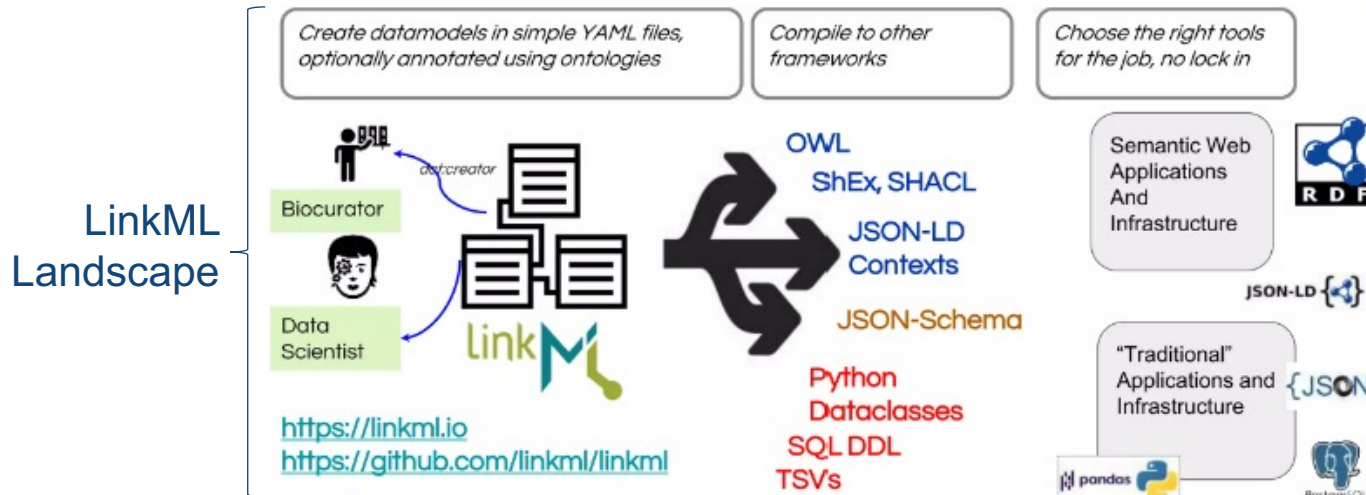




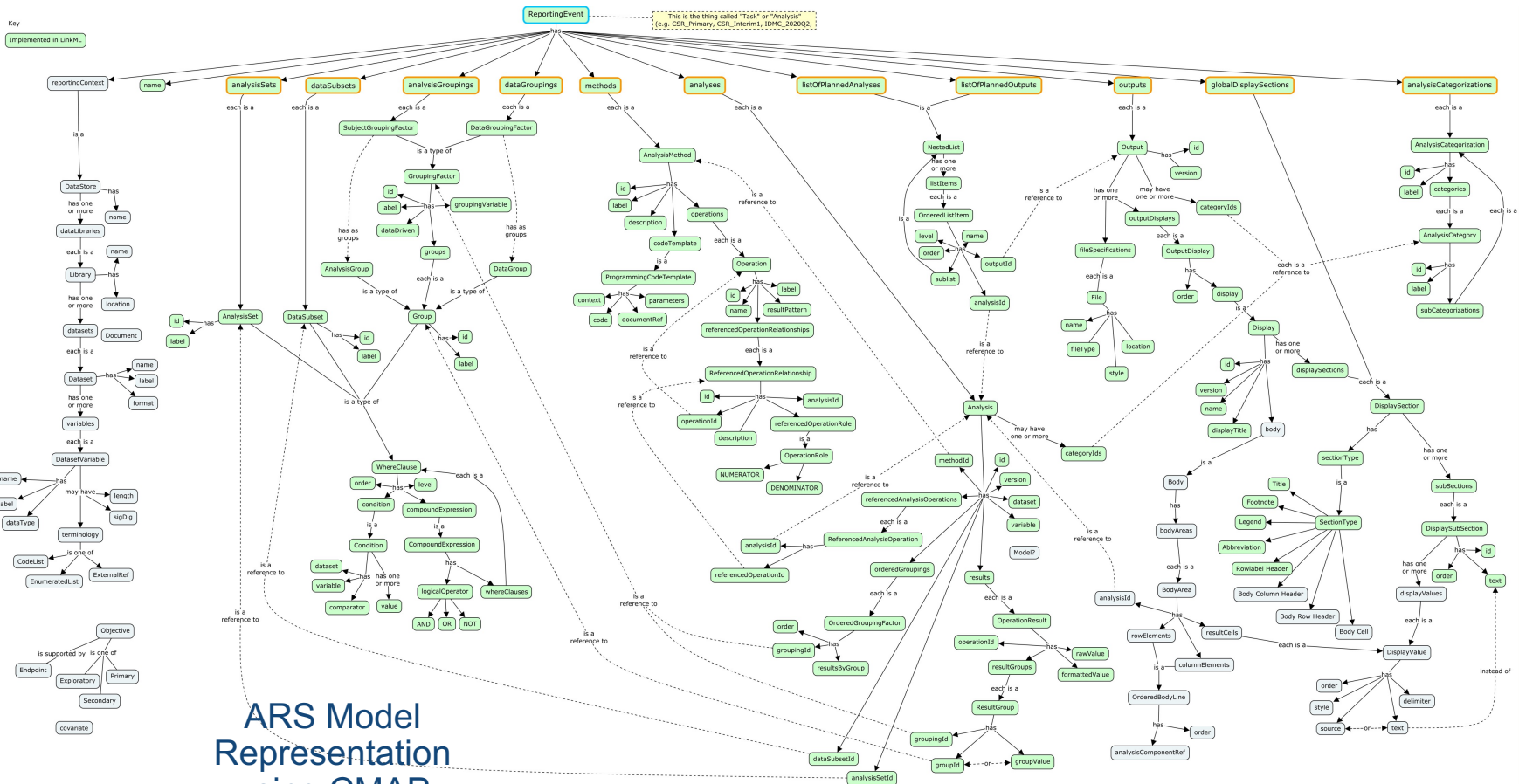
# Overview of ARS Model and User Guide

# Using LinkML to Create Analysis Results Model

- LinkML is a general-purpose modeling language that can be used with linked data, JSON, and other formalisms



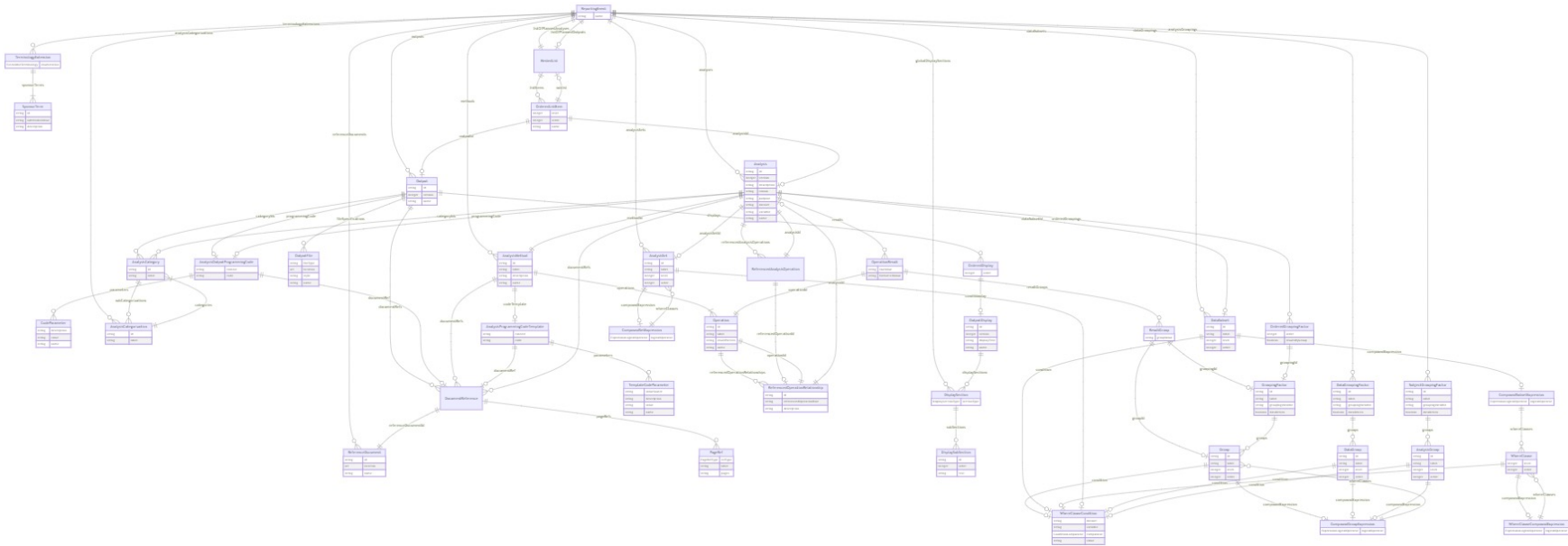
Reference: <https://www.slideshare.net/cmungall/linkml-intro-july-2022pptx>



# ARS Model Representation using CMAP (DRAFT)



# ARS Model Representation using Mermaid Markdown (DRAFT)



# Review Examples

Analysis Set

Data Subset

Analysis Grouping

Data Grouping

Method

Analysis

Results

## Summary of Demographics

Study - CDISC 360 Page x of y

Table 14.1.1  
Summary of Demographics  
Safety Population

Characteristics	Placebo (N=XX)	Xanomeline Low Dose (N=XX)	Xanomeline High Dose (N=XX)
<b>Age (years)</b>			
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
<b>Age Group, n (%)</b>			
< 65 years	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
≥ 65 years	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<b>Gender, n (%)</b>			
Male	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
Female	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<b>Ethnicity, n (%)</b>			
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>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<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>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<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

# Review Examples

## Summary of Demographics

Study - CDISC 360 Page x of y

Table 14.1.1  
Summary of Demographics  
Safety Population

Characteristics	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

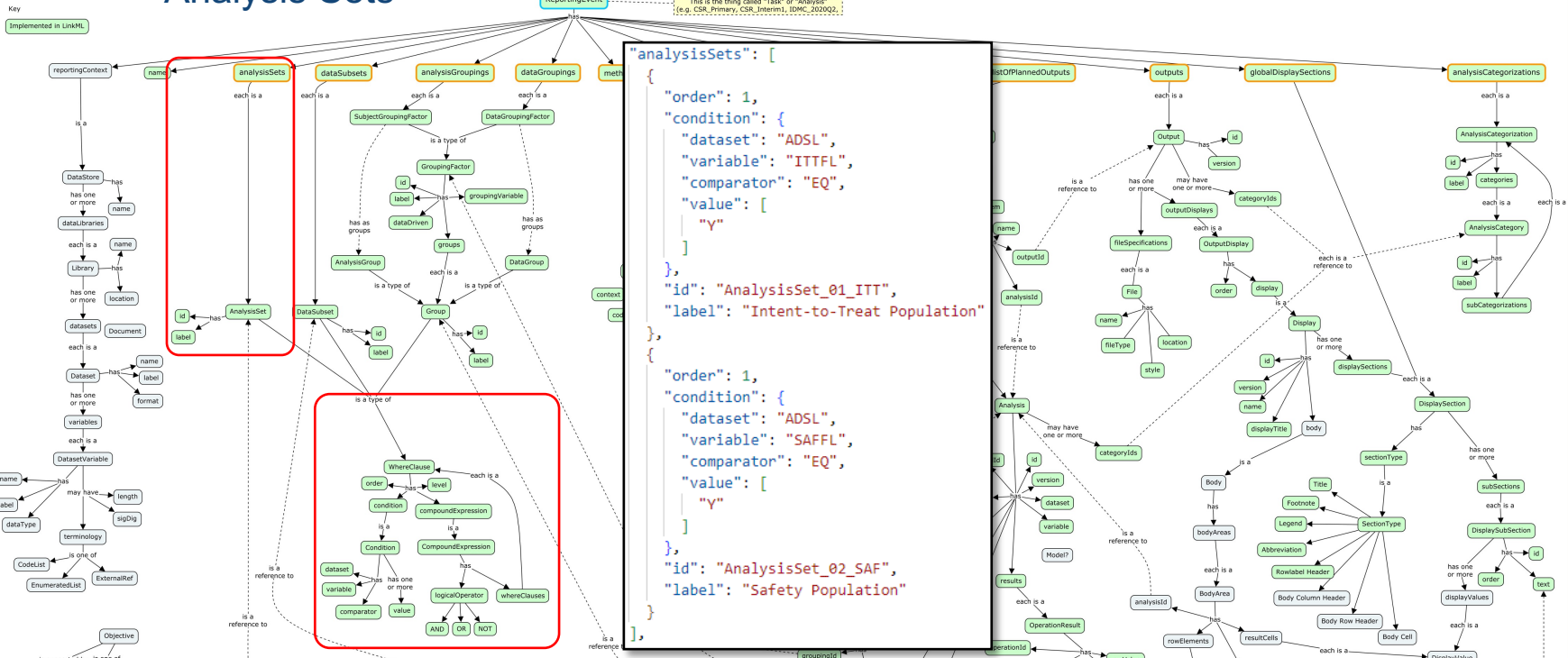
Table 14.3.1.1  
Summary of TEAE by System Organ Class and Preferred Term  
Safety Population

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>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<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>	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
<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

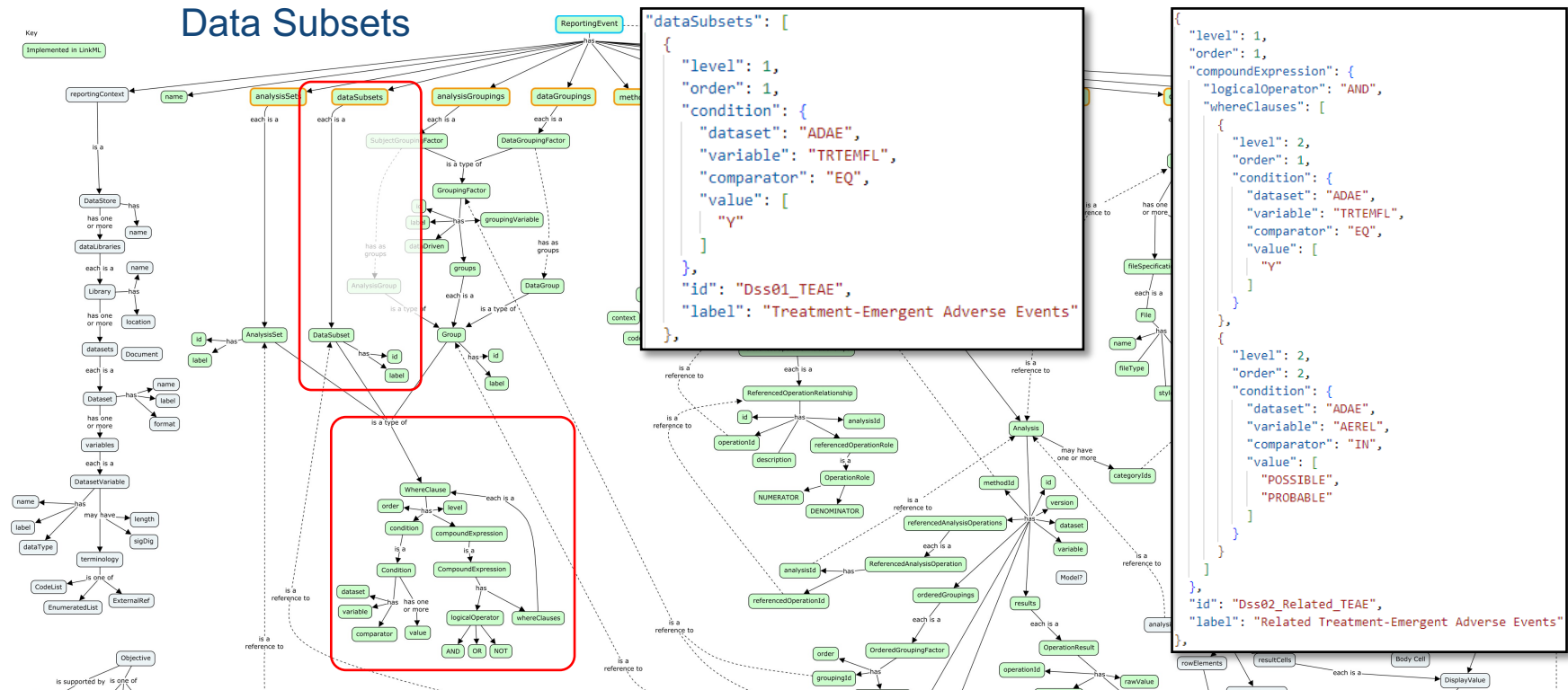
# Analysis Sets



id	label	order	condition_dataset	condition_variable	condition_comparator	condition_value
AnalysisSet_01_ITT	Intent-to-Treat Population	1	ADSL	ITTFL	EQ	Y
AnalysisSet_02_SAF	Safety Population	1	ADSL	SAFFL	EQ	Y

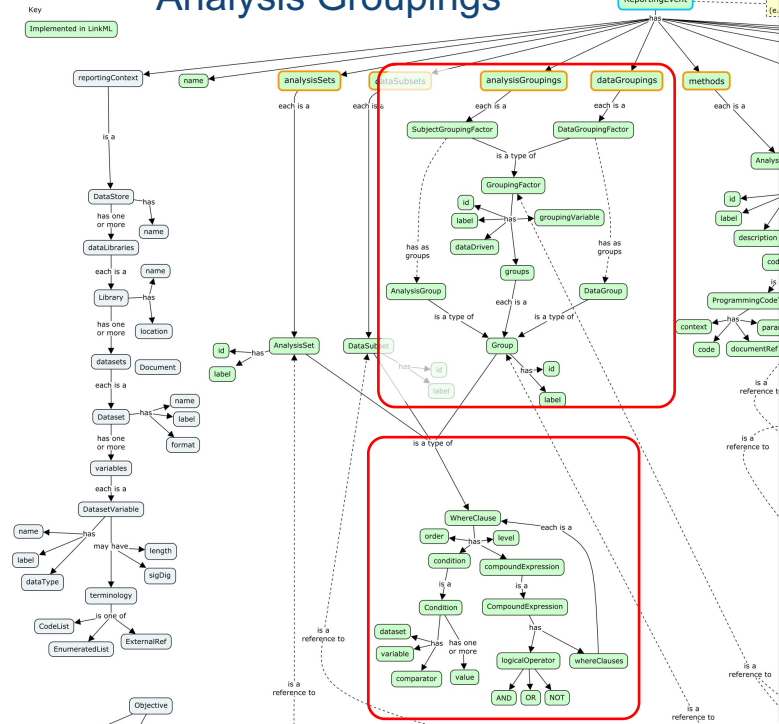


# Data Subsets



id	label	level	order	compoundExpression_logicalOperator	condition_dataset	condition_variable	condition_comparator	condition_value
Dss01_TEAE	Treatment-Emergent Adverse Events	1	1		ADAE	TRTEMFL	EQ	Y
Dss02_Related_TEAE	Related Treatment-Emergent Adverse Events	1	1	AND				
Dss02_Relateds_TEAE	Related Treatment-Emergent Adverse Events	2	1		ADAE	TRTEMFL	EQ	Y
Dss02_Relateds_TEAE	Related Treatment-Emergent Adverse Events	2	2		ADAE	AEREL	IN	POSSIBLE   PROBABLE

# Analysis Groupings

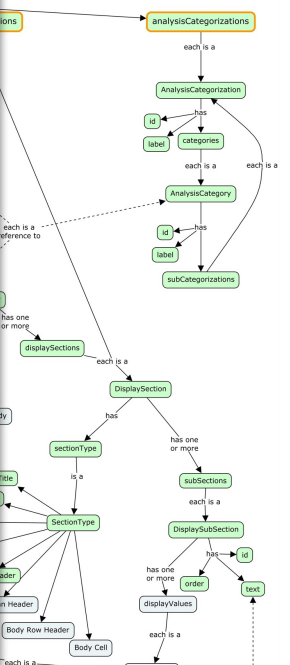


```

"analysisGroupings": [
  {
    "id": "AnlsGrouping_01_Sex",
    "dataDriven": false,
    "label": "Gender",
    "groupingVariable": "SEX",
    "groups": [
      {
        "order": 1,
        "condition": {
          "dataset": "ADSL",
          "variable": "SEX",
          "comparator": "EQ",
          "value": [
            "M"
          ]
        }
      },
      {
        "order": 2,
        "condition": {
          "dataset": "ADSL",
          "variable": "SEX",
          "comparator": "EQ",
          "value": [
            "F"
          ]
        }
      }
    ]
  },
  {
    "id": "AnlsGrouping_01_Sex_1",
    "label": "Male"
  },
  {
    "id": "AnlsGrouping_01_Sex_2",
    "label": "Female"
  }
]
  
```

```

{
  "id": "AnlsGrouping_02_Trt",
  "dataDriven": false,
  "label": "Treatment",
  "groupingVariable": "TRT01A",
  "groups": [
    {
      "order": 1,
      "condition": {
        "dataset": "ADSL",
        "variable": "TRT01A",
        "comparator": "EQ",
        "value": [
          "Placebo"
        ]
      }
    },
    {
      "order": 2,
      "condition": {
        "dataset": "ADSL",
        "variable": "TRT01A",
        "comparator": "EQ",
        "value": [
          "Xanomeline Low Dose"
        ]
      }
    },
    {
      "order": 3,
      "condition": {
        "dataset": "ADSL"
      }
    }
  ]
}
  
```

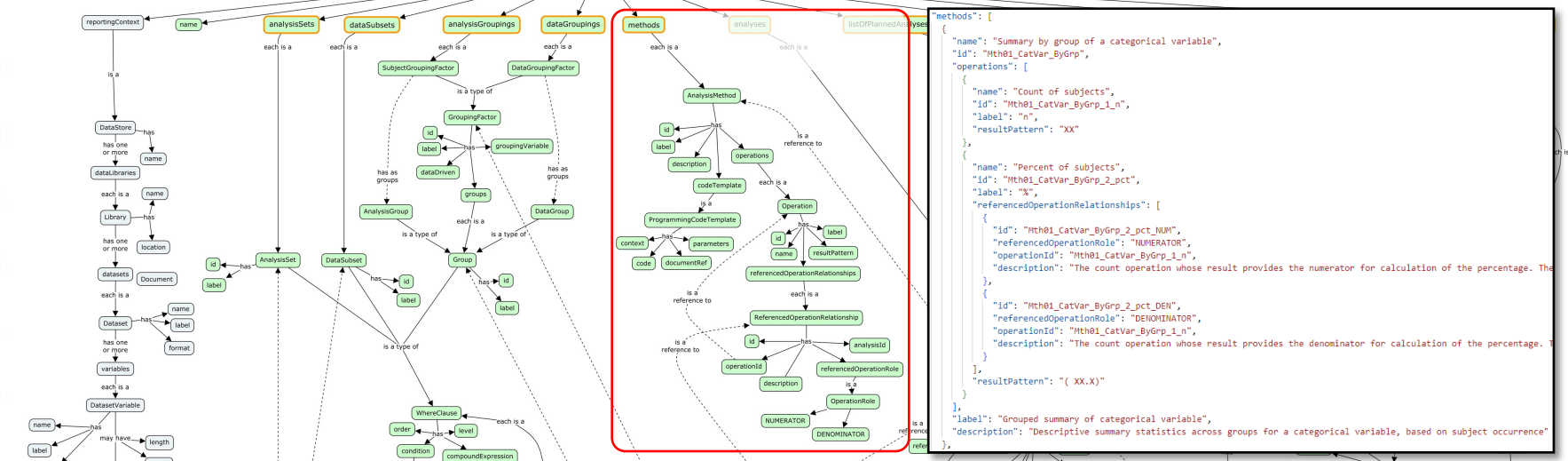


id	label	groupingVariable	dataDriven	group_id	group_label	group_order	group_condition_dataset	group_condition_variable	group_condition_comparator	group_condition_value
AnlsGrouping_01_Sex	Gender	SEX	FALSE	AnlsGrouping_01_Sex_1	Male	1	ADSL	SEX	EQ	M
AnlsGrouping_01_Sex	Gender	SEX	FALSE	AnlsGrouping_01_Sex_2	Female	2	ADSL	SEX	EQ	F
AnlsGrouping_02_Trt	Treatment	TRT01A	FALSE	AnlsGrouping_02_Trt_1	Placebo	1	ADSL	TRT01A	EQ	Placebo
AnlsGrouping_02_Trt	Treatment	TRT01A	FALSE	AnlsGrouping_02_Trt_2	Xanomeline Low Dose	2	ADSL	TRT01A	EQ	Xanomeline Low Dose
AnlsGrouping_02_Trt	Treatment	TRT01A	FALSE	AnlsGrouping_02_Trt_3	Xanomeline High Dose	3	ADSL	TRT01A	EQ	Xanomeline High Dose

# Methods

Key  
Implemented in LinkML

This is the thing called "Task" or "Analysis"  
(e.g. CSR\_Primary, CSR\_Interim, IDMC\_2020Q2)



```

"methods": [
  {
    "name": "Summary by group of a categorical variable",
    "id": "Mth01_CatVar_ByGrp",
    "operations": [
      {
        "name": "Count of subjects",
        "id": "Mth01_CatVar_ByGrp_1_n",
        "label": "n",
        "resultPattern": "XX"
      },
      {
        "name": "Percent of subjects",
        "id": "Mth01_CatVar_ByGrp_2_pct",
        "label": "%",
        "referencedOperationRelationships": [
          {
            "id": "Mth01_CatVar_ByGrp_2_pct_NUM",
            "referencedOperationRole": "NUMERATOR",
            "operationId": "Mth01_CatVar_ByGrp_1_n",
            "description": "The count operation whose result provides the numerator for calculation of the percentage. The"
          },
          {
            "id": "Mth01_CatVar_ByGrp_2_pct_DEN",
            "referencedOperationRole": "DENOMINATOR",
            "operationId": "Mth01_CatVar_ByGrp_1_n",
            "description": "The count operation whose result provides the denominator for calculation of the percentage. T"
          }
        ]
      }
    ],
    "label": "Grouped summary of categorical variable",
    "description": "Descriptive summary statistics across groups for a categorical variable, based on subject occurrence"
  }
]
    
```

name	label	description	operation_id	operation_name	operation_order	operation_label	operation_resultPattern
Summary by group of a categorical variable	Grouped summary of a categorical variable	Descriptive summary statistics across groups for a categorical variable, based on subject occurrence	Mth01_CatVar_ByGrp_1_n	Count of subjects	1	n	XX
Summary by group of a categorical variable	Grouped summary of a categorical variable	Descriptive summary statistics across groups for a categorical variable, based on subject occurrence	Mth01_CatVar_ByGrp_2_pct	Percent of subjects	2	%	{ XX.X }

operation_referencedResultRelation	operation_referencedResultRelation	operation_referencedResultRelationship	operation_referencedResultRelation	operation_referencedResultRelationship
Mth01_CatVar_ByGrp_2_pct_NUM	Mth01_CatVar_ByGrp_1_n	The count operation whose result provides the numerator for calculation of the percentage. The referenced analysis should be the analysis that contains this percent operation.	Mth01_CatVar_ByGrp_2_pct_DEN	Mth01_CatVar_ByGrp_1_n



# Analyses

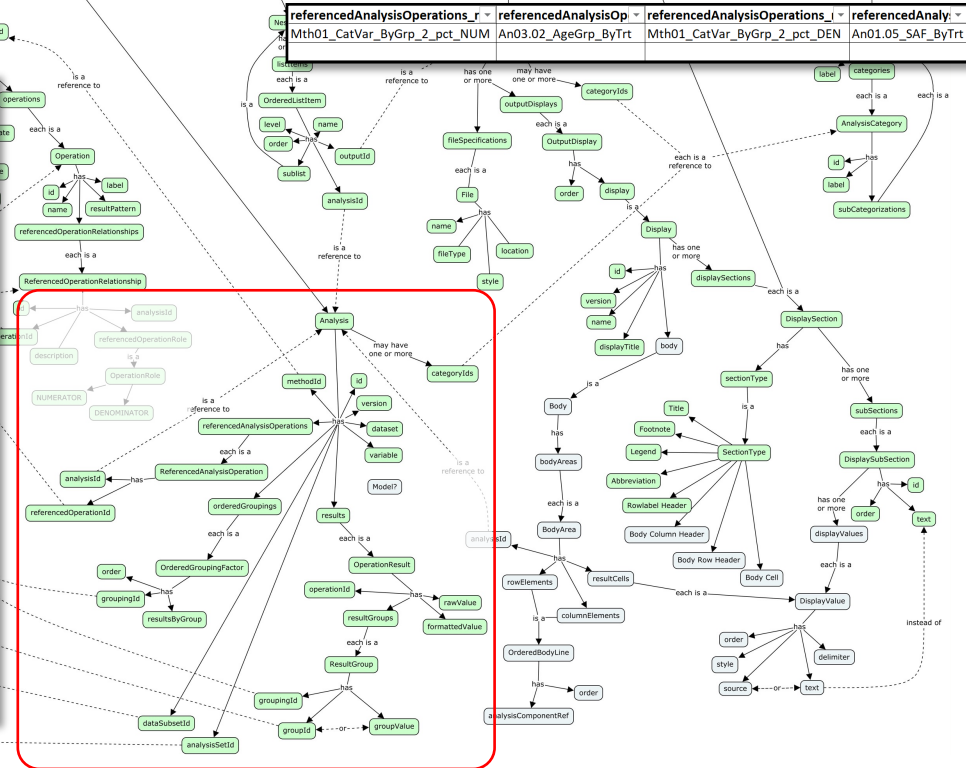
Key  
Implemented in LinkML

ReportingEvent  
This is the thing called "Task" or "Analysis"  
(e.g. CSR\_Primary, CSR\_Interim, IDMC\_2020Q2)

id	versi	name	categoryIds	analysisSetId	groupingId1	groupingId2	groupingId3	dataSubsetId	data	variable	method_id
An03.02_AgeGrp_ByTrt	1	Summary of Subjects by Treatment and Age Group		AnalysisSet_02_SAF	AnlsGrouping_02_Trt	AnlsGrouping_03_AgeGrp			ADSL	USUBJID	Mth01_CatVar_ByGrp
An08.02_ChgBl_ByTrt	1	Summary of Change from Baseline by Treatment, Parameter and Visit		AnalysisSet_02_SAF	AnlsGrouping_02_Trt	AnlsGrouping_08_Param	AnlsGrouping_09_Visit	Dss10_VS_NonBl_AnRec	ADVS	CHG	Mth02_ContVar_ByGrp

```

"analyses": [
  {
    "name": "Summary of Change from Baseline by Treatment, Parameter and Visit",
    "id": "An08.02_ChgBl_ByTrt",
    "methodId": "Mth02_ContVar_ByGrp",
    "version": 1,
    "analysisSetId": "AnalysisSet_02_SAF",
    "orderedGroupings": [
      {
        "order": 1,
        "groupingId": "AnlsGrouping_02_Trt"
      },
      {
        "order": 2,
        "groupingId": "AnlsGrouping_08_Param"
      },
      {
        "order": 3,
        "groupingId": "AnlsGrouping_09_Visit"
      }
    ],
    "dataSubsetId": "Dss10_VS_NonBl_AnRec",
    "dataset": "ADVS",
    "variable": "CHG",
    "results": [
    ]
  }
]
  
```



# Analysis Results

Key  
Implemented in LinkML

ReportingEvent  
This is the thing called "Task" or "Analysis"  
(e.g. CSR\_Primary\_CSR\_Interim)\_IDMC\_2020Q2

id	operation_id	resultGroup1_groupingId	resultGroup1_groupId	resultGroup2_groupingId	resultGroup2_groupId	resultGroup3_groupingId	resultGroup3_groupId	rawValue	formattedVal
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_1_n	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	249	249
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_2_Mean	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	-3.3012	-3.3
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_3_SD	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	14.60121	(14.60)
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_4_Media	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	-2	-2.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_5_Q1	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	-12	-12.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_6_Q3	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	4	4.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_7_Min	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	-38	-38
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_8_Max	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_02	40	40
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_1_n	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	243	243
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_2_Mean	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	-3.02469	-3.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_3_SD	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	15.66829	(15.67)
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_4_Media	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	-2	-2.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_5_Q1	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	-12	-12.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_6_Q3	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	6	6.0
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_7_Min	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	-48	-48
An08.02_ChgBl_ByTrt	Mth02_ContVar_ByGrp_8_Max	AnlsGrouping_02_Trt	AnlsGrouping_02_Trt_1	AnlsGrouping_08_Param	AnlsGrouping_08_Param_1	AnlsGrouping_09_Visit	AnlsGrouping_09_Visit_03	50	50

```

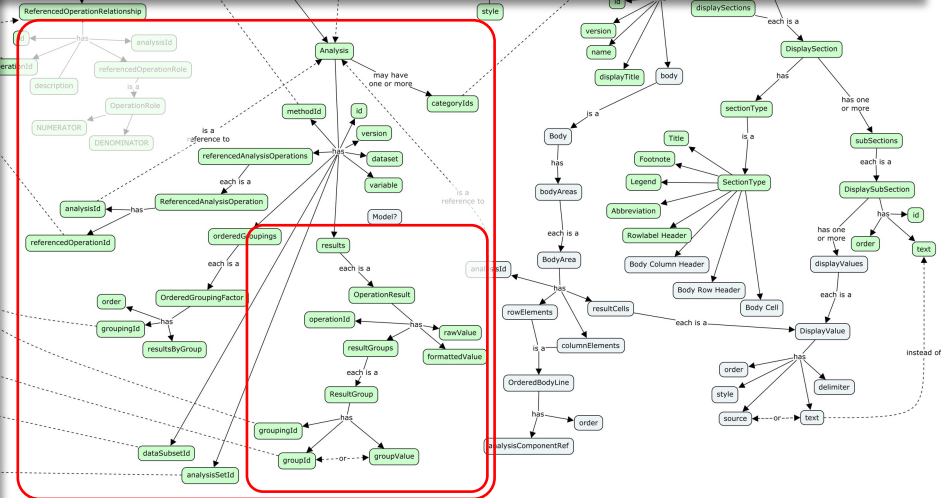
"results": {
  "operationId": "Mth02_ContVar_ByGrp_1_n",
  "resultGroups": [
    {
      "groupingId": "AnlsGrouping_02_Trt",
      "groupId": "AnlsGrouping_02_Trt_1"
    },
    {
      "groupingId": "AnlsGrouping_08_Param",
      "groupId": "AnlsGrouping_08_Param_1"
    },
    {
      "groupingId": "AnlsGrouping_09_Visit",
      "groupId": "AnlsGrouping_09_Visit_02"
    }
  ],
  "rawValue": "249",
  "formattedValue": "249"
},
"operationId": "Mth02_ContVar_ByGrp_2_Mean",
"resultGroups": [
  {
    "groupingId": "AnlsGrouping_02_Trt",
    "groupId": "AnlsGrouping_02_Trt_1"
  },
  {
    "groupingId": "AnlsGrouping_08_Param",
    "groupId": "AnlsGrouping_08_Param_1"
  },
  {
    "groupingId": "AnlsGrouping_09_Visit",
    "groupId": "AnlsGrouping_09_Visit_02"
  }
],
"rawValue": "-3.301204819",
"formattedValue": "- 3.3"
},
"operationId": "Mth02_ContVar_ByGrp_3_SD",
"resultGroups": [

```

```

"rawValue": "14",
"formattedValue": "14"
},
"operationId": "Mth01_CatVar_ByGrp_1",
"resultGroups": [
  {
    "groupingId": "AnlsGrouping_02_Trt",
    "groupId": "AnlsGrouping_02_Trt_1"
  },
  {
    "groupingId": "AnlsGrouping_03_Age",
    "groupId": "AnlsGrouping_03_AgeGrp"
  }
],
"rawValue": "72",
"formattedValue": "72"
},
"operationId": "Mth01_CatVar_ByGrp_1",
"resultGroups": [
  {
    "groupingId": "AnlsGrouping_02_Trt",
    "groupId": "AnlsGrouping_02_Trt_1"
  }
],

```



# Concepts, Not Layout

Analysis ID:	An03.2_AgeGrp_ByTrt						
Display Value:	formattedValue						
			AnlsGrouping_02_Trtr	Treatment	Placebo	Xanomeline Low Dose	Xanomeline High Dose
		AnlsGrouping_03_AgeGp	Mth01_CatVar_ByGrp				
		Age Group	Operation				
		< 65 years	n		14	8	11
		< 65 years	%		( 16.3)	( 9.5)	( 13.1)
		≥ 65 years	n		72	76	73
		≥ 65 years	%		( 83.7)	( 90.5)	( 86.9)

Analysis ID:	An03.2_AgeGrp_ByTrt								
Display Value:	formattedValue								
		AnlsGrouping_02_Trtr	Treatment		Placebo	Placebo	Xanomeline Low Dose	Xanomeline Low Dose	Xanomeline High Dose
		Mth01_CatVar_ByGrp	Operation		n	%	n	%	n
		AnlsGrouping_03_AgeGp							
		Age Group							
		< 65 years			14	( 16.3)	8	( 9.5)	11
		≥ 65 years			72	( 83.7)	76	( 90.5)	73
									( 13.1)
									( 86.9)

Analysis ID:	An03.2_AgeGrp_ByTrt						
Display Value:	formattedValue						
			Mth01_CatVar_ByGrp	Operation	n	%	
		AnlsGrouping_02_Trtr	AnlsGrouping_03_AgeGp				
		Treatment	Age Group				
		Placebo	< 65 years		14	( 16.3)	
		Placebo	≥ 65 years		72	( 83.7)	
		Xanomeline Low Dose	< 65 years		8	( 9.5)	
		Xanomeline Low Dose	≥ 65 years		76	( 90.5)	
		Xanomeline High Dose	< 65 years		11	( 13.1)	
		Xanomeline High Dose	≥ 65 years		73	( 86.9)	

# Outputs

Key  
Implemented in LinkML

This is the thing called "Task" or "Analysis" (e.g. CSR\_Primary, CSR\_Interim, IDMC\_2020Q2)

id	name	version	displayTitle	displaySection_sectionType	displaySection_sectionId	displaySection_subSection_id	ion_order/displaySection_subSection_text
Disp14.1.1	Demog	1	Summary of Demographics	Title	Disp14.1.1_Title_1	1	Table 14.1.1
Disp14.1.1	Demog	1	Summary of Demographics	Title	Disp14.1.1_Title_2	2	Summary of Demographics
Disp14.1.1	Demog	1	Summary of Demographics	Title	Disp14.1.1_Title_3	3	Safety Population
Disp14.1.1	Demog	1	Summary of Demographics	Footnote	Disp14.1.1_Fnote_1	1	Source dataset: adsl, Generated on: DDMONYYYY:HH:MM
Disp14.1.1	Demog	1	Summary of Demographics	Footnote	Disp14.1.1_Fnote_2	2	Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM

```

"outputDisplays": [
  {
    "order": 1,
    "display": {
      "name": "Demog",
      "id": "Disp14.1.1",
      "version": 1,
      "displayTitle": "Summary of Demographics",
      "displaySections": [
        {
          "sectionType": "Title",
          "subSections": [
            {
              "id": "Disp14.1.1_Title_1",
              "order": 1,
              "text": "Table 14.1.1"
            },
            {
              "id": "Disp14.1.1_Title_2",
              "order": 2,
              "text": "Summary of Demographics"
            },
            {
              "id": "Disp14.1.1_Title_3",
              "order": 3,
              "text": "Safety Population"
            }
          ]
        },
        {
          "sectionType": "Footnote",
          "subSections": [
            {
              "id": "Disp14.1.1_Fnote_1",
              "order": 1,
              "text": "Source dataset: adsl, Generated on: DDMONYYYY:HH:MM"
            },
            {
              "id": "Disp14.1.1_Fnote_2",
              "order": 2,
              "text": "Program: <pid>.sas, Output: <pid><oid>.rtf, Generated on: DDMONYYYY:HH:MM"
            }
          ]
        }
      ]
    }
  }
]

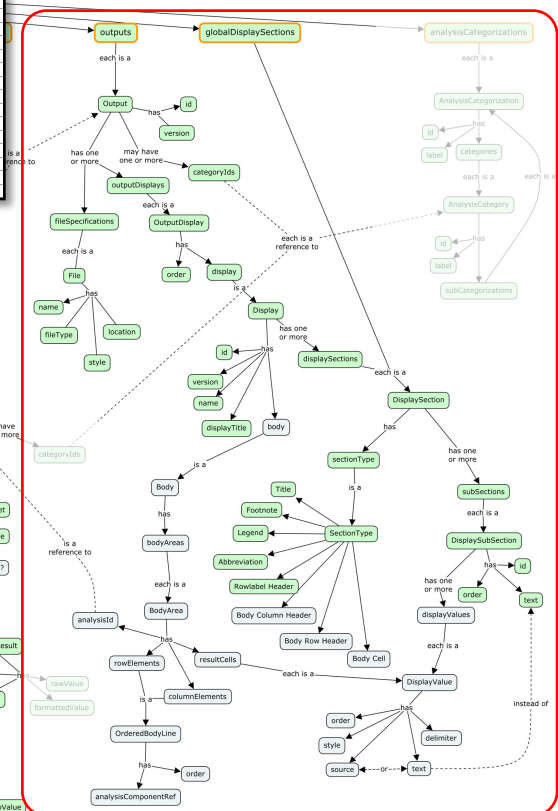
```

Table 14.1.1  
Summary of Demographics  
Safety Population

	Placebo (N=XX)	Xarelto (N=XX)
<b>Characteristics</b>		
Age (years)		
n	XX	XX
Mean (SD)	XX.X (XX.XX)	XX.X (XX.XX)
Median	XX.X	XX.X
Q1, Q3	XX.X, XX.X	XX.X, XX.X
Min, Max	XX, XX	XX, XX
Age Group, n (%)		
< 65 years	XX (XX.X)	XX (XX.X)
≥ 65 years	XX (XX.X)	XX (XX.X)
Gender, n (%)		
Male	XX (XX.X)	XX (XX.X)
Female	XX (XX.X)	XX (XX.X)
Ethnicity, n (%)		
Hispanic or Latino	XX (XX.X)	XX (XX.X)
Not Hispanic or Latino	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

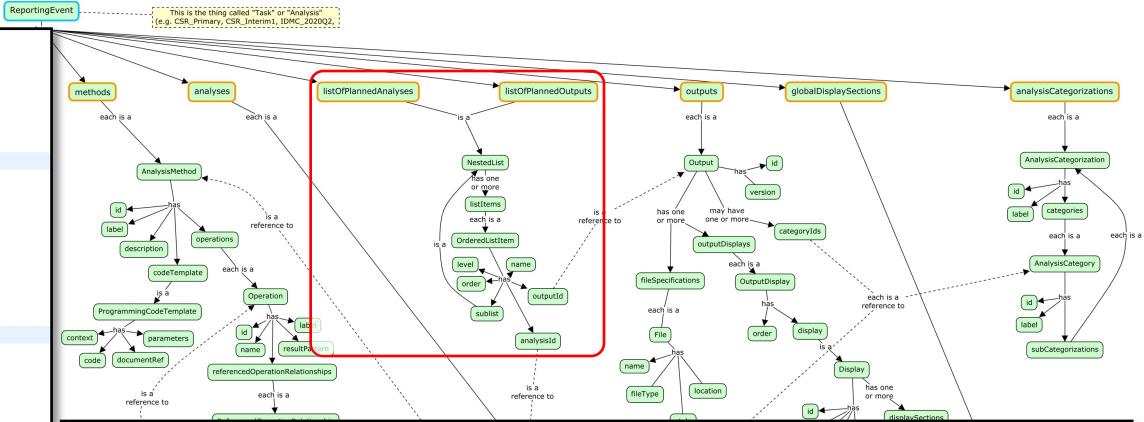
"name": "Demog",  
"id": "Disp14.1.1",  
"sectionType": "Rowlabel Header",



# List of Planned Analyses/Outputs

```

Key
"listOfPlannedAnalyses": {
  "listItems": [
    {
      "name": "Summary of Demographics",
      "level": 1,
      "order": 1,
      "sublist": {
        "listItems": [...
      ]
    },
    "outputId": "Out14.1.1"
  },
  {
      "name": "Overall Summary of Treatment-Emergent Adverse Events",
      "level": 1,
      "order": 2,
      "sublist": {
        "listItems": [...
      ]
    },
    "outputId": "Out14.3.1.1"
  },
  {
      "name": "Summary of TEAE by System Organ Class and Preferred Term",
      "level": 1,
      "order": 3,
      "sublist": {
        "listItems": [
          {
            "name": "Summary of Subjects by Treatment and System Organ Class ",
            "level": 2,
            "order": 1,
            "analysisId": "An07.09_Soc_ByTrt"
          },
          {
            "name": "Summary of Subjects by Treatment, System Organ Class and Preferred Term ",
            "level": 2,
            "order": 2,
            "analysisId": "An07.10_SocPt_ByTrt"
          }
        ]
      },
    },
    "outputId": "Out14.3.2.1"
  },
  ],
  "covariate":
}
    
```



level	name	order	analysisId	outputId
1	Summary of Demographics	1		Out14.1.1
2	Summary of Subjects by Treatment	1	An01.05_SAF_ByTrt	
2	Summary of Age by Treatment	2	An03.01_Age_ByTrt	
2	Summary of Subjects by Treatment and Age Group	3	An03.02_AgeGrp_ByTrt	
2	Summary of Subjects by Treatment and Sex	4	An03.03_Sex_ByTrt	
2	Summary of Subjects by Treatment and Ethnicity	5	An03.04_Ethnic_ByTrt	
2	Summary of Subjects by Treatment and Race	6	An03.05_Race_ByTrt	
2	Summary of Height by Treatment	7	An03.06_Height_ByTrt	
1	Overall Summary of Treatment-Emergent Adverse Events	2		Out14.3.1.1
2	Summary of Subjects with At Least One TEAE, by Treatment	1	An07.01_TEAE_ByTrt	
2	Summary of Subjects with At Least One Related TEAE, by Treatment	2	An07.02_RelTEAE_ByTrt	
2	Summary of Subjects with At Least One Serious TEAE, by Treatment	3	An07.03_SerTEAE_ByTrt	
2	Summary of Subjects with At Least One Related Serious TEAE, by Treatment	4	An07.04_RelSerTEAE_ByTrt	
2	Summary of Subjects with At Least One TEAE Leading to Death, by Treatment	5	An07.05_TEAEld2Dth_ByTrt	
2	Summary of Subjects with At Least One Related TEAE Leading to Death, by Treatment	6	An07.06_RelTEAEld2Dth_ByTrt	
2	Summary of Subjects with At Least One TEAE Leading to Dose Modification, by Treatment	7	An07.07_TEAEld2DoseMod_ByTrt	
2	Summary of Subjects with At Least One TEAE Leading to Treatment Discontinuation, by Treatment	8	An07.08_TEAEld2TrtDsc_ByTrt	
1	Summary of TEAE by System Organ Class and Preferred Term	3		Out14.3.2.1
2	Summary of Subjects by Treatment and System Organ Class	1	An07.09_Soc_ByTrt	
2	Summary of Subjects by Treatment, System Organ Class and Preferred Term	2	An07.10_SocPt_ByTrt	
1	Summary of Observed and Change from Baseline by Scheduled Visits - Vital Signs	4		Out14.3.3.1a
2	Summary of Observed Value by Treatment, Parameter and Visit	1	An08.01_Obs_ByTrt	
2	Summary of Change from Baseline by Treatment, Parameter and Visit	2	An08.02_ChgBl_ByTrt	



# Implementations

```
> Mth02_ContVar_ByGrp_7_Min: Minimum (Min)
> Mth02_ContVar_ByGrp_8_Max: Maximum (Max)
1.3. Summary of Subjects by Treatment and Age Group
Analysis: An03.02_AgeGrp_ByTrt
Population: Safety Population [ADSL.SAFFL EQ 'Y']
Groupings:
  1. Treatment:
    1. Placebo [ADSL.TRT01A EQ 'Placebo']
    2. Xanomeline Low Dose [ADSL.TRT01A EQ 'Xanomeline Low Dose']
    3. Xanomeline High Dose [ADSL.TRT01A EQ 'Xanomeline High Dose']
  2. Age Group:
    1. < 65 years [ADSL.AGEGR1 EQ '<65']
    2. ≥ 65 years [ADSL.AGEGR1 IN ('65-80', '>80')]
Analysis Variable: ADSL.USUBJID
Method: Summary by group of a categorical variable
Operations:
  > Mth01_CatVar_ByGrp_1_n: Count of subjects (n)
  > Mth01_CatVar_ByGrp_2_pct: Percent of subjects (%)
    - Numerator: result of operation Mth01_CatVar_ByGrp_1_n for this analysis
    - Denominator: result of operation Mth01_CatVar_ByGrp_1_n for analysis An01.05_SAF_ByTrt
1.4. Summary of Subjects by Treatment and Sex
Analysis: An03.03_Sex_ByTrt
Population: Safety Population [ADSL.SAFFL EQ 'Y']
Groupings:
  1. Treatment:
    1. Placebo [ADSL.TRT01A EQ 'Placebo']
    2. Xanomeline Low Dose [ADSL.TRT01A EQ 'Xanomeline Low Dose']
    3. Xanomeline High Dose [ADSL.TRT01A EQ 'Xanomeline High Dose']
  2. Gender:
    1. Male [ADSL.SEX EQ 'M']
    2. Female [ADSL.SEX EQ 'F']
```

# Analysis Results Standard Repo on GitHub

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

The screenshot shows the GitHub repository page for 'cdisc-org/analysis-results-standard'. The repository is public and has 65 commits, 4 branches, and 3 tags. The file list includes folders like 'HowTos', 'docs', 'documents', 'images', 'model', 'project', 'utilities/python', and 'workfiles', along with files like 'CODE\_OF\_CONDUCT.md', 'CONTRIBUTING.md', 'LICENSE', 'README.md', and 'mkdocs.yml'. The 'README.md' file is selected, showing its description: 'The goals of CDISC Analysis Results Standards team is to develop: • Analysis Results Metadata Technical Specification (ARM-TS), to support automation, traceability, and creation of data disallows'. Callout boxes point to the 'model' folder, 'utilities/python' folder, 'project' folder, and the 'About' section.

**Model:** representations of the model (YAML, JSON, Mermaid ER, YUML, SVG)

**Workfiles:** CMAP, examples

**Project:** Auto-generated content (Python classes/API, documentation, model structures)

**Utilities:** Example programs

# Analysis Results Standard Model Documentation

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

The screenshot shows the main page of the Analysis Results Standard (ARS) documentation. The header includes the title "Analysis Results Standard (ARS)" and a search bar. A left sidebar lists navigation options: "Analysis Results Standard (ARS)", "Classes", "Slots", "Enumerations", "Types", and "Subsets". The main content area features the title "Analysis Results Standard (ARS)" followed by a description: "DRAFT Logical model to support both the prospective specification of analyses and the fully contextualized representation of the results of the analyses." Below this, the URI is given as "https://www.cdisc.org/ars/1-0 Name: ars\_idm". A "Classes" section lists several classes with their descriptions:

Class	Description
<a href="#">Analysis</a>	An analysis that is designed to meet a requirement of the reporting event
<a href="#">AnalysisCategorization</a>	A set of related implementer-defined categories that can be used to categoriz...
<a href="#">AnalysisCategory</a>	An implementer-defined category of analyses/outputs, which may include one or...
<a href="#">AnalysisGroup</a>	A subdivision of the subject population based on a defined factor (e
<a href="#">AnalysisMethod</a>	A set of one or more statistical operations
<a href="#">AnalysisOutputProgrammingCode</a>	Programming statements and/or a reference to the program used to perform a sp...

The screenshot shows the class documentation for "ReportingEvent". It includes a class hierarchy diagram, an inheritance list, a table of slots, and identifier/mapping information.

**Class: ReportingEvent**  
A set of analyses and outputs created to meet a specific reporting requirement, such as a CSR or clinical analysis.  
URI: ars:ReportingEvent

**Inheritance**

- ReportingEvent
- ReportingEvent

**Slots**

Name	Cardinality and Range	Description	Inheritance
analysisOutput	0..1 AnalysisOutput	A structured list of the analysis outputs for the reporting event.	class
analysisOutputList	0..1 AnalysisOutputList	An optional structured list of the outputs defined for the reporting event.	class
analysisOutputList	0..1 AnalysisOutputList	The analysis sets (subject population) defined for the reporting event.	class
analysisOutputList	0..1 AnalysisOutputList	Characteristics used to subdivide the subject population in	class
analysisOutputList	0..1 AnalysisOutputList	Characteristics used to subdivide sites (reports or the analysis domain) in	class
analysisOutputList	0..1 AnalysisOutputList	Display function	class
analysisOutputList	0..1 AnalysisOutputList	AnalysisCategorization	class
analysisOutputList	0..1 AnalysisOutputList	The analyses defined for the reporting event.	class
analysisOutputList	0..1 AnalysisOutputList	The default method used to analyze any analysis results.	class
analysisOutputList	0..1 AnalysisOutputList	Output	class
analysisOutputList	0..1 AnalysisOutputList	Reference to a resource	class
analysisOutputList	0..1 AnalysisOutputList	Any operator defined extension to extension terminology.	class
analysisOutputList	1..1 String	URI	class

**Identifier and Mapping Information**

Schema Source

- From schema: <https://cdisc.org/ars/1-0>

**Mappings**

Mapping Type	Mapped Slot
uri	ars:ReportingEvent
uri	ars:ReportingEvent

**LinkML Source**

Direct

- Details

Induced

- Details



# Example Reporting Events

- Common Safety Displays
  - Summary of Demographics
  - Overall Summary of Treatment-Emergent Adverse Events
  - Summary of TEAE by System Organ Class and Preferred Term
  - Summary of Observed and Change from Baseline by Scheduled Visits - Vital Signs
  - Summary of Observed and Change from Baseline by Scheduled Visits - Vital Signs <Vertical Layout>
- FDA STF
  - Table 2: Baseline Demographic and Clinical Characteristics, Safety Population

# User Guide

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

## 1 INTRODUCTION

- 1.1 Background and Purpose
- 1.2 Relationship to Other CDISC Standards
- 1.3 How to Read this Document

## 2 ANALYSIS RESULTS MODEL

- 2.1 ReportingEvent
- 2.2 [Common Components](#)
  - 2.2.1 NestedList
  - 2.2.2 AnalysisOutputCategorization
  - 2.2.3 ReferenceDocument
    - 2.2.3.1 DocumentReference
  - 2.2.4 TerminologyExtension
    - 2.2.4.1 ExtensibleTerminologyTerm
  - 2.2.5 AnalysisOutputProgrammingCode
- 2.3 Analysis Components
  - 2.3.1 WhereClause
    - 2.3.1.1 WhereClauseCondition
    - 2.3.1.2 WhereClauseCompoundExpression
  - 2.3.2 AnalysisSet
  - 2.3.3 DataSubset
  - 2.3.4 GroupingFactor
    - 2.3.4.1 SubjectGroupingFactor
    - 2.3.4.2 DataGroupingFactor
  - 2.3.5 AnalysisMethod
    - 2.3.5.1 Operation
    - 2.3.5.2 AnalysisProgrammingCodeTemplate
  - 2.3.6 Analysis
    - 2.3.6.1 OperationResult

## 2.4 Output Components

- 2.4.1 GlobalDisplaySection
- 2.4.2 Output
  - 2.4.2.1 OutputDisplay

## 3 EXAMPLE REPORTING EVENTS

- 3.1 Common Safety Displays
  - 3.1.1 Summary of Demographics
  - 3.1.2 Overall Summary of Treatment-Emergent Adverse Events
  - 3.1.3 Summary of TEAE by System Organ Class and Preferred Term
  - 3.1.4 Summary of Observed and Change from Baseline by Scheduled Visits - Vital Signs
  - 3.1.5 Summary of Observed and Change from Baseline by Scheduled Visits - Vital Signs <Vertical Layout>
- 3.2 FDA STF
  - 3.2.1 Table 2: Baseline Demographic and Clinical Characteristics, Safety Population

## APPENDICES

- Appendix A: Glossary and Abbreviations
- Appendix B: Representations and Warranties, Limitations of Liability, and Disclaimers

# User Guide Page Layouts

## Class Descriptions

### WhereClauseCondition

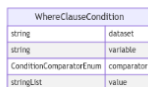
Created by Richard Marshall, last modified on Aug 14, 2023

#### Model Documentation

Class: `WhereClauseCondition`

The `WhereClauseCondition` class is used to define simple selection criteria conditions in the `condition` attribute of the `WhereClause` class (or any of its specializations).

#### ER Diagram: WhereClauseCondition



In each instance of the `WhereClauseCondition` class:

- The **dataset** attribute contains the name of the dataset in which the condition is being applied.
- The **variable** attribute contains the name of the variable (within the specified dataset) to which the condition applies.
- The **comparator** attribute contains a value from the `ConditionComparatorEnum` enumeration ("EQ" for "is equal to", "LT" for "is less than", etc) to indicate how values of the **variable** attribute are compared to the **value** attribute.
- The **value** attribute specifies the selection value, or values, for comparison with values in the specified dataset variable. If the value of the **comparator** attribute is:
  - "IN" or "NOTIN", there should be at least 2 values specified in the **value** attribute; otherwise there should be no more than 1 value specified in the **value** attribute.
  - "EQ" or "NE", the **value** attribute may contain no value to indicate a condition based on a missing value (i.e., "is missing" or "is not missing" respectively).

#### Example 1

This example shows an instance of the `WhereClause` class (or one of its specializations) that defines a simple condition based on a single specified selection value. The example shows:

- `ADSL.SAFFL EQ 'Y'`

Note that some attributes (such as the expected **level** and **order** attributes) have been excluded from this example.

#### YAML Example

```
...
condition:
  dataset: ADSL
  variable: SAFFL
  comparator: EQ
  value:
  - Y
```

This condition could be represented in tabular form as:

dataset	variable	comparator	value
ADSL	SAFFL	EQ	Y

Other conditions based on a single value (i.e., with a **comparator** value of "EQ", "NE", "LT", "LE", "GT", or "GE") would be represented in a similar way.

## Example Reporting Events

### Overall Summary of Treatment-Emergent Adverse Events

Created by Richard Marshall, last modified on Aug 16, 2023

Study - CD40C 349 Page x of y

Table 14.3.1-1.1  
Overall Summary of Treatment-Emergent Adverse Events  
Safety Population

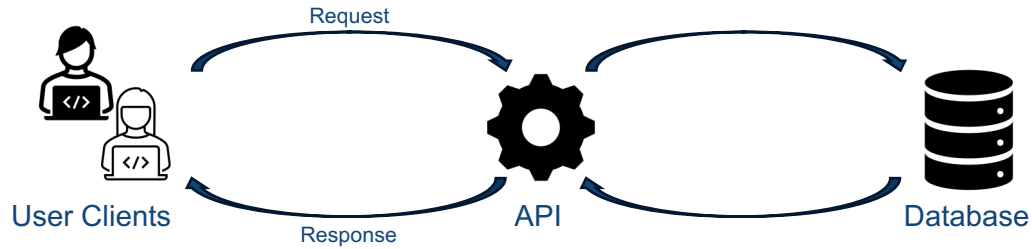
Categories, n (%)	Placebo (N=92)	Xanomeline Low Dose (N=92)	Xanomeline High Dose (N=92)
Number of subjects with at least one event			
TEAE	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
Related TEAE	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
Serious TEAE	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
Related Serious TEAE	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
TEAE Leading to Death	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
Related TEAE Leading to Death	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
TEAE Leading to New Modification [a]	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)
TEAE Leading to Treatment Discontinuation	XX ( XX.X)	XX ( XX.X)	XX ( XX.X)

Note: TEAE=Treatment-Emergent Adverse Events  
[a] Dose Modification Includes Dose Reduction; Drug Interrupted in the AE section taken with study treatment.

Source dataset: adev, Generated on: 20230816T11:11:10M  
Report: CD40C\_349, Output: CD40C\_349\_P11, Generated on: 20230816T11:11:10M

- Overall Summary of Treatment-Emergent Adverse Events  
Output: Out14-3-1-1 (Overall Summary of Treatment-Emergent Adverse Events)  
Documentation:  
> AE Summary Table Shell (/AE\_Summary\_Table\_Shell.pdf)  
Categories:  
> Safety  
> Adverse Events  
Output File(s):  
> RTF Format: t14-3-1-1-teae-summ (.)  
> PDF Format: t14-3-1-1-teae-summ (.)  
Displays:  
1. Disp14-3-1-1 (AE\_Summ)  
Display Title: Overall Summary of Treatment-Emergent Adverse Events  
Sections:  
> Title:

# Analysis Results Standard Application Programming Interface (API)



GET	/mdr/ars/packages/	Get All Ars Packages
GET	/mdr/ars/packages/{package_id}/reportingevents/	Get All Package Reporting Events
GET	/mdr/ars/reportingevents/{reportingevent_id}/	Get Reporting Event
GET	/mdr/ars/reportingevents/{reportingevent_id}/methods/	Get All Reportingevents Methods
GET	/mdr/ars/methods/{method_id}/	Get Method
GET	/mdr/ars/methods/{method_id}/operations/	Get All Methods Operations

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

```
Curl
curl -X 'GET' \
'http://127.0.0.1:8000/mdr/ars/reportingevents/0/methods/' \
-H 'accept: application/json'

Request URL
http://127.0.0.1:8000/mdr/ars/reportingevents/0/methods/

Server response
Code      Details
200

Response body
[
  {
    "name": "Summary by group of a categorical variable",
    "id": "Mth01_CatVar_Summ_ByGrp",
    "operations": [
      {
        "name": "Count of subjects",
        "id": "Mth01_CatVar_Summ_ByGrp_1_n",
        "label": "n",
        "resultPattern": "XX"
      },
      {
        "name": "Percent of subjects",
        "id": "Mth01_CatVar_Summ_ByGrp_2_pct",
        "label": "%",
        "referencedOperationRelationships": [
          {
            "id": "Mth01_CatVar_Summ_ByGrp_2_pct_NUM",
            "referencedOperationRole": "NUMERATOR",
            "operationId": "Mth01_CatVar_Summ_ByGrp_1_n",
            "description": "The count operation whose result is operation."
          },
          {
            "id": "Mth01_CatVar_Summ_ByGrp_2_pct_DEN",
            "referencedOperationRole": "DENOMINATOR",
            "operationId": "Mth01_CatVar_Summ_ByGrp_1_n",
            "description": "The count operation whose result is operation."
          }
        ]
      }
    ]
  }
]
```



## **ARS model will drive automation and open-source tool development**



# CDISC ARS: Hackathon Objectives



Drive adoption of  
CDISC Analysis  
Results Standard



Foster open-source  
software tools for  
operationalization

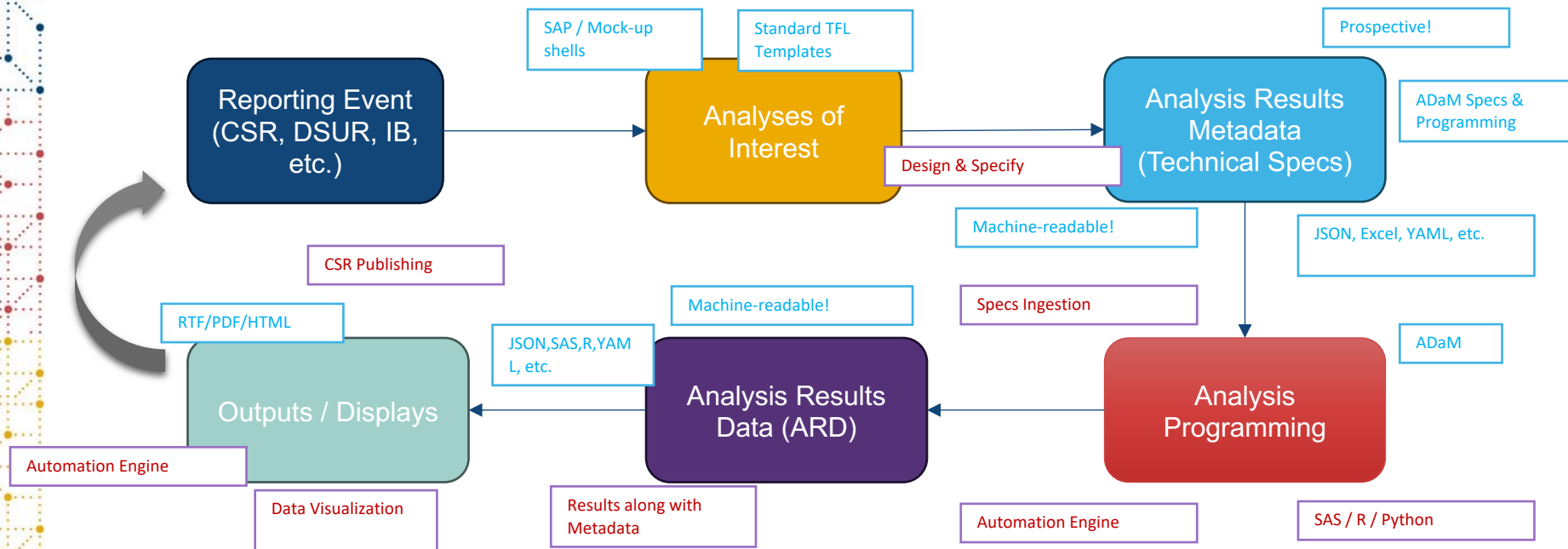


Leveraging hackathon  
learnings to enhance  
the standards

# Hackathon Timeline



# ARS Model Supported Workflow and Entry Points



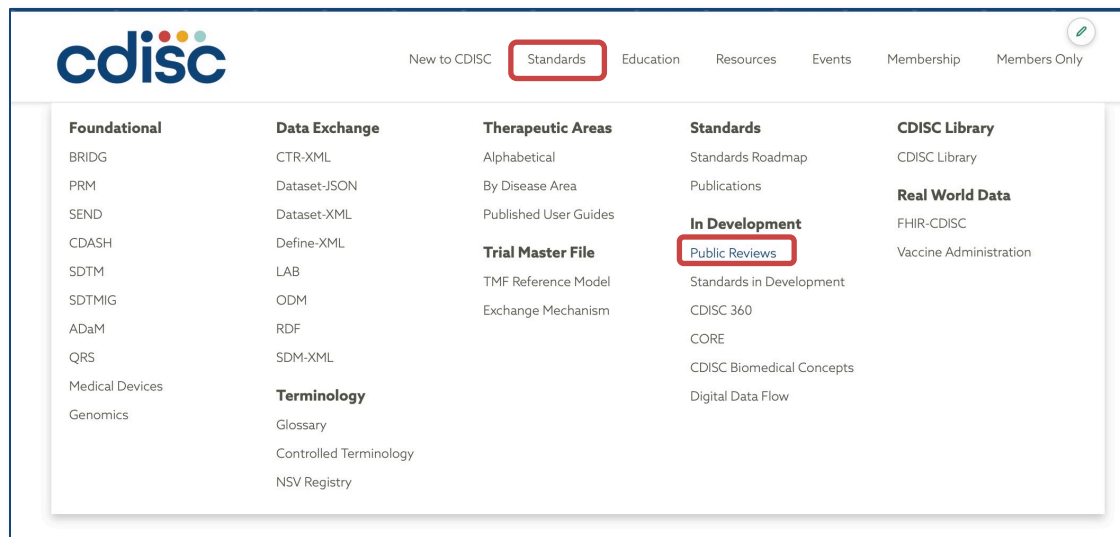
# Release Plan

## Version 1.0

- Logical Model
- User Guide
- Common safety examples based on team and FDA developed tables
  - Demographics
  - Adverse Events
  - Vital signs
- CDISC ARS Hackathon: July 12th, 2023
- CDISC Internal Review: August 18th, 2023
- CDISC Public Review: Through January 15th, 2023
- US Interchange Workshop: October 2023
- Anticipated Final Release: Q1 2024

# Provide Public Review Comments by January 15th!

- <https://www.cdisc.org/public-review/analysis-results-standard-v1-0>

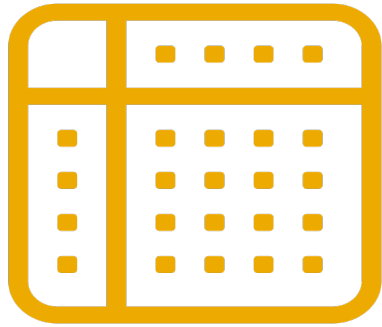


The screenshot shows the CDISC website's navigation menu. The 'Standards' link is highlighted with a red box. Under the 'Standards' section, the 'Public Reviews' link is also highlighted with a red box. The website layout includes the CDISC logo, a navigation bar with links for 'New to CDISC', 'Standards', 'Education', 'Resources', 'Events', 'Membership', and 'Members Only', and a main content area with five columns of links.

Foundational	Data Exchange	Therapeutic Areas	Standards	CDISC Library
BRIDG	CTR-XML	Alphabetical	Standards Roadmap	CDISC Library
PRM	Dataset-JSON	By Disease Area	Publications	<b>Real World Data</b>
SEND	Dataset-XML	Published User Guides	<b>In Development</b>	FHIR-CDISC
CDASH	Define-XML	<b>Trial Master File</b>	<b>Public Reviews</b>	Vaccine Administration
SDTM	LAB	TMF Reference Model	Standards in Development	
SDTMIG	ODM	Exchange Mechanism	CDISC 360	
ADaM	RDF		CORE	
QRS	SDM-XML		CDISC Biomedical Concepts	
Medical Devices	<b>Terminology</b>		Digital Data Flow	
Genomics	Glossary			
	Controlled Terminology			
	NSV Registry			



# What Is Next?: Adding Informative Content



ADaM Dataset



Analysis Results and  
Associated Metadata



TFL Example

Planned Collaboration with  
PHUSE Safety Analytics  
Working Group



# Contact Details

## **Bhavin Busa**

ARS Product Owner & Co-Lead

[bhavin@clymbclinical.com](mailto:bhavin@clymbclinical.com)

## **Richard Marshall**

Principal Data Modeler

[rmarshall@accuratesystems.co.uk](mailto:rmarshall@accuratesystems.co.uk)

## **Bess LeRoy**

Head of Standards Innovation, CDISC

[bleroy@cdisc.org](mailto:bleroy@cdisc.org)

