



2024 CDISC + TMF
EUROPE INTERCHANGE

BERLIN

24-25 APRIL: CONFERENCE & EXPO | 22, 23, 26 APRIL: TRAININGS

**ARD Generator:
A SAS tool for metadata-driven analysis using the
Analysis Result Standard**

Karl Wallendszus
Senior Analyst/Programmer, Oxford Population Health



Meet the Speaker

Karl Wallendszus

Title: Senior Analyst/Programmer

Organization: Clinical Trial Service Unit, Oxford Population Health

Karl Wallendszus is a statistical programmer in the Oxford Population Health, a department of the University of Oxford. Over the last 30 years he has worked on a number of large clinical trials in cardiology and kidney disease as well as the RECOVERY trial of treatments for COVID-19. He has been using CDISC standards since 2010, which at the time was unusual for an academic institution. Karl was a volunteer in the CDISC team which developed the new Analysis Results Standard (ARS) and participated in the CDISC ARS Hackathon the summer of 2023.



Disclaimer and Disclosures

- *The views and opinions expressed in this presentation are those of the author(s) and do not necessarily reflect the official policy or position of CDISC.*
- *The author(s) have no real or apparent conflicts of interest to report.*



Agenda

1. The Analysis Result Standard
2. The ARD Generator tool
3. Usage example
4. Conclusions

The Analysis Result Standard



ARS and JSON

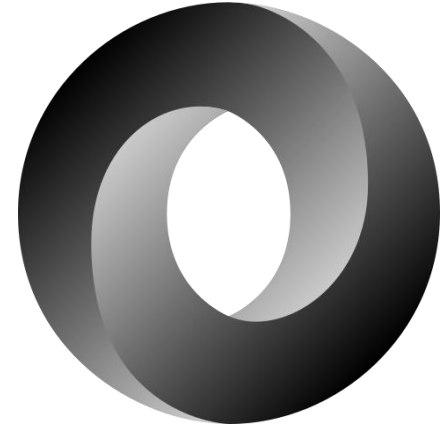


We use JSON as a representation of the ARS.

But JSON is not part of that standard.

And other representations of ARS are possible.

JSON



ARS: Separating results from displays



PhUSE US Connect 2019

Paper DH12

Analysis Results that Save Trees - #KillTFLs

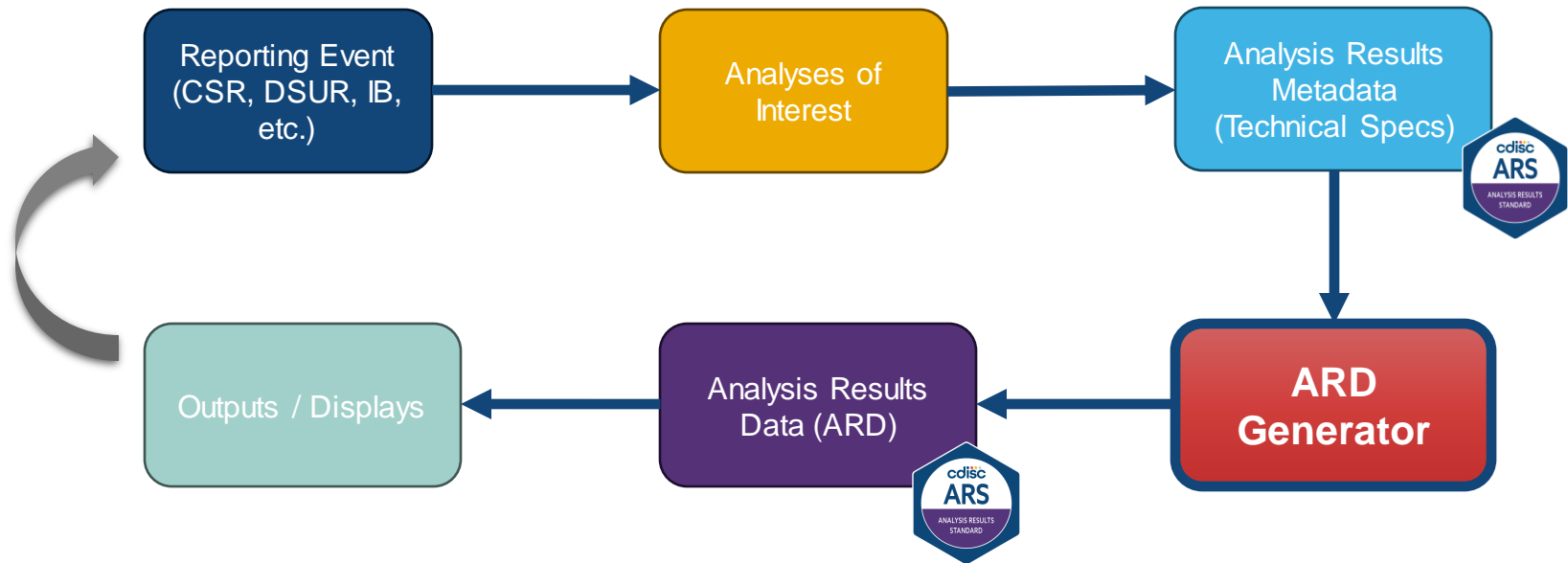
Chris Decker, d-wise, Morrisville, NC, USA

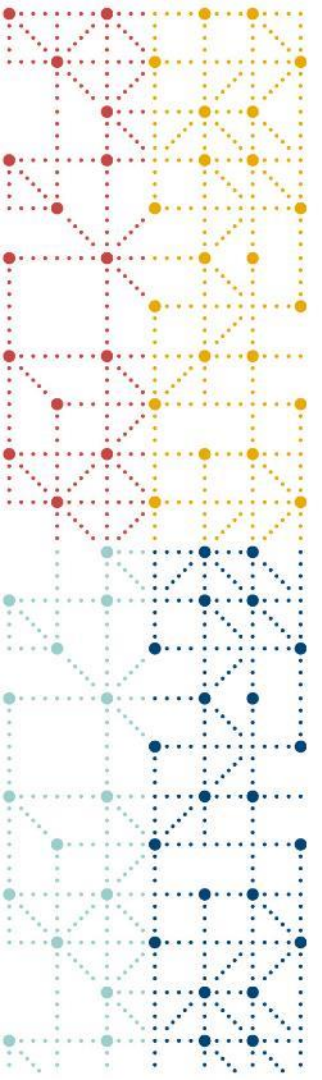
ABSTRACT

In today's clinical programming environment, we spend large amounts of time (and money) producing Tables, Figures, and Listings – single use, throw-away artifacts that I would challenge focus more on the cosmetics than the content. Artifacts that, by their own admission, are rarely looked at by regulatory reviewers. We build these reports for one narrowly focused outcome instead of making that result, and many more, available for many different uses. What if we could capture and store analysis results that gave the results context and allowed a clinician or reviewer to dynamically search the results within and across studies without sifting through 300 pages of paper? We can do this! This presentation will provide an overview of a project where a graph-based analysis results data store was designed and populated to capture and make analysis results available to many different stakeholders reducing their workload by 50%. We can leverage new technology and/or methodologies to optimize our archaic processes and #killTFLs.

https://phuse.s3.eu-central-1.amazonaws.com/Archive/2019/Connect/US/Baltimore/PAP_DH12.pdf

ARS Model Supported Workflow and Entry Points





The ARD Generator tool



ARD Generator

- Written in SAS
- Ingests ARS metadata from JSON → SAS datasets
- Processes analyses specified in that metadata
- Uses a macro library to automate analysis
- Output to Analysis Result Dataset (ARD)

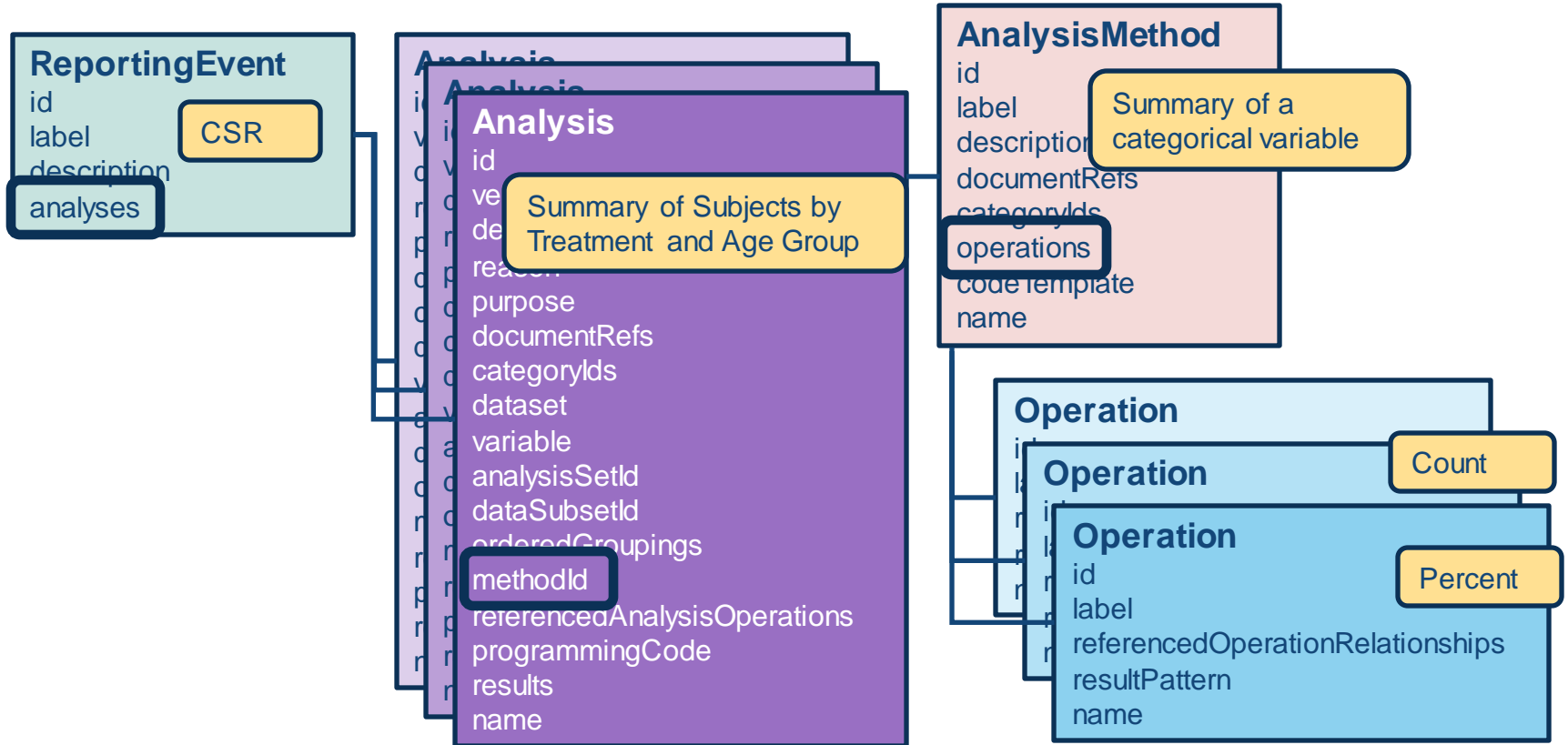
Ingesting JSON

```
"analyses": [  
  {  
    "id": "An03_02_AgeGrp_Comp_ByTrt",  
    "reason": {  
      "controlledTerm": "SPECIFIED IN SAP"  
    },  
    "purpose": {  
      "controlledTerm": "PRIMARY OUTCOME MEASURE"  
    },  
    "methodId": "Mth03_CatVar_Comp_PChiSq",  
    "version": 1,  
    "categoryIds": [  
      "Catn_01_Grp_1_Pop",  
      "Catn_02_Dclass_1_Sbj",  
      "Catn_03_SbjDType_1_Dm"  
    ],  
    "documentRefs": [  
      {  
        "referenceDocumentId": "CDISCPIL0T01_SAP",  
        "pageRefs": [  
          {  
            "refType": "PhysicalRef",  
            "label": "9.2. Demographic and Baseline Characteristics",  
            "pageNumbers": [  
              13  
            ]  
          }  
        ]  
      }  
    ],  
    "analysisSetId": "AnalysisSet_02_SAF",  
    "dependentGroupings": [  
      {  
        "id": "An03_01_Age_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Age by Treatment"  
      },  
      {  
        "id": "An03_01_Age_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Age by Treatment"  
      },  
      {  
        "id": "An03_02_AgeGrp_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects by Treatment and Age Group"  
      },  
      {  
        "id": "An03_02_AgeGrp_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Age Group by Treatment"  
      },  
      {  
        "id": "An03_03_Sex_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects by Treatment and Sex"  
      },  
      {  
        "id": "An03_03_Sex_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Sex by Treatment"  
      },  
      {  
        "id": "An03_04_Ethnic_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects by Treatment and Ethnicity"  
      },  
      {  
        "id": "An03_04_Ethnic_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Ethnicity by Treatment"  
      },  
      {  
        "id": "An03_05_Race_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects by Treatment and Race"  
      },  
      {  
        "id": "An03_05_Race_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Race by Treatment"  
      },  
      {  
        "id": "An03_06_Height_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Height by Treatment"  
      },  
      {  
        "id": "An03_06_Height_Comp_ByTrt",  
        "version": 1,  
        "name": "Comparison of Height by Treatment"  
      },  
      {  
        "id": "An07_01_TEAE_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One TEAE, by Treatment"  
      },  
      {  
        "id": "An07_01_TEAE_Comp_ByTrt_PlacLow",  
        "version": 1,  
        "name": "Comparison of Subjects with TEAEs by Treatment - Placebo vs Low Dose"  
      },  
      {  
        "id": "An07_01_TEAE_Comp_ByTrt_PlacHigh",  
        "version": 1,  
        "name": "Comparison of Subjects with TEAEs by Treatment - Placebo vs High Dose"  
      },  
      {  
        "id": "An07_02_RelTEAE_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One Related TEAE, by Treatment"  
      },  
      {  
        "id": "An07_03_SerTEAE_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One Serious TEAE, by Treatment"  
      },  
      {  
        "id": "An07_04_RelSerTEAE_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One Related Serious TEAE, by Treatment"  
      },  
      {  
        "id": "An07_05_TEAEld2Dth_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One TEAE Leading to Death, by Treatment"  
      },  
      {  
        "id": "An07_06_RelTEAEld2Dth_Summ_ByTrt",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One Related TEAE Leading to Death, by Treatment"  
      },  
      {  
        "id": "An07_07_TEAEld2DoseMod_Summ_By",  
        "version": 1,  
        "name": "Summary of Subjects with At Least One TEAE Leading to Dose Modification, by Treatment"  
      }  
    ]  
  }  
]
```

VIEWTABLE: Jsonmd.Analyses

| | id | version | name |
|----|----------------------------------|---------|---|
| 1 | An01_05_SAF_Summ_ByTrt | 1 | Summary of Subjects by Treatment |
| 2 | An03_01_Age_Summ_ByTrt | 1 | Summary of Age by Treatment |
| 3 | An03_01_Age_Comp_ByTrt | 1 | Comparison of Age by Treatment |
| 4 | An03_02_AgeGrp_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Age Group |
| 5 | An03_02_AgeGrp_Comp_ByTrt | 1 | Comparison of Age Group by Treatment |
| 6 | An03_03_Sex_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Sex |
| 7 | An03_03_Sex_Comp_ByTrt | 1 | Comparison of Sex by Treatment |
| 8 | An03_04_Ethnic_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Ethnicity |
| 9 | An03_04_Ethnic_Comp_ByTrt | 1 | Comparison of Ethnicity by Treatment |
| 10 | An03_05_Race_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Race |
| 11 | An03_05_Race_Comp_ByTrt | 1 | Comparison of Race by Treatment |
| 12 | An03_06_Height_Summ_ByTrt | 1 | Summary of Height by Treatment |
| 13 | An03_06_Height_Comp_ByTrt | 1 | Comparison of Height by Treatment |
| 14 | An07_01_TEAE_Summ_ByTrt | 1 | Summary of Subjects with At Least One TEAE, by Treatment |
| 15 | An07_01_TEAE_Comp_ByTrt_PlacLow | 1 | Comparison of Subjects with TEAEs by Treatment - Placebo vs Low Dose |
| 16 | An07_01_TEAE_Comp_ByTrt_PlacHigh | 1 | Comparison of Subjects with TEAEs by Treatment - Placebo vs High Dose |
| 17 | An07_02_RelTEAE_Summ_ByTrt | 1 | Summary of Subjects with At Least One Related TEAE, by Treatment |
| 18 | An07_03_SerTEAE_Summ_ByTrt | 1 | Summary of Subjects with At Least One Serious TEAE, by Treatment |
| 19 | An07_04_RelSerTEAE_Summ_ByTrt | 1 | Summary of Subjects with At Least One Related Serious TEAE, by Treatment |
| 20 | An07_05_TEAEld2Dth_Summ_ByTrt | 1 | Summary of Subjects with At Least One TEAE Leading to Death, by Treatment |
| 21 | An07_06_RelTEAEld2Dth_Summ_ByTrt | 1 | Summary of Subjects with At Least One Related TEAE Leading to Death, by Treatment |
| 22 | An07_07_TEAEld2DoseMod_Summ_By | 1 | Summary of Subjects with At Least One TEAE Leading to Dose Modification, by Treatment |

How analyses are modelled



A macro library allows automation

ard-generator / macros / source /

KarlWallendzusz Added operations for this

Name

..

append_addcols.sas

build_expression.sas

build_work_dataset.sas

define_analset.sas

op_catvar_comp_pchisq.sas

op_catvar_count_bygrp_n.sas

op_catvar_summ_bygrp_pct.sas

op_contvar_summ_bygrp.sas

outline_ard.sas

run_analysis.sas

run_method.sas

run_operation.sas

standardize_grouping.sas

standardize_groupings.sas

```
* Loop through analyses;
%local ianal analid;
%let ianal = 1;
%do %while(%scan(&analids., &ianal., '|') ne );
    %let analid = %scan(&analids., &ianal., '|');

    * Run this analysis;
    %run_analysis(mdlib=&mdlib., datalib=&datalib., ardlib=&ardlib.,
        analid=&analid., debugfl=&debugfl.);
```

```
* Run the analysis method;
%run_method(mdlib=&mdlib., datalib=&datalib., ardlib=&ardlib.,
```

```
    %* Loop through analysis operations;
    %do iop = 1 %to &noperations.;

        %* Execute this operation;
        %run_operation(mdlib=&mdlib., datalib=&datalib.,
            opid=&&opid&iop., opseq=&iop., nop=&noperations., methid=&methid.,
            analid=&analid., analsetid=&analsetid., datasubsetid=&datasubsetid.,
            groupingids=&groupingids., analds=workds, analvar=&analvar.,
            ard=&ardlib..ard, debugfl=&debugfl.);

    %end;
```

Operation macros

ard-generator / macros / source /

KarlWallendzusz Added operations for chisq a

Name

..

append_addcols.sas

build_expression.sas

build_work_dataset.sas

define_analset.sas

op_catvar_comp_pchisq.sas

op_catvar_count_bygrp_n.sas

op_catvar_summ_bygrp_pct.sas

op_contvar_summ_bygrp.sas

outline_ard.sas

run_analysis.sas

run_method.sas

run_operation.sas

standardize_grouping.sas

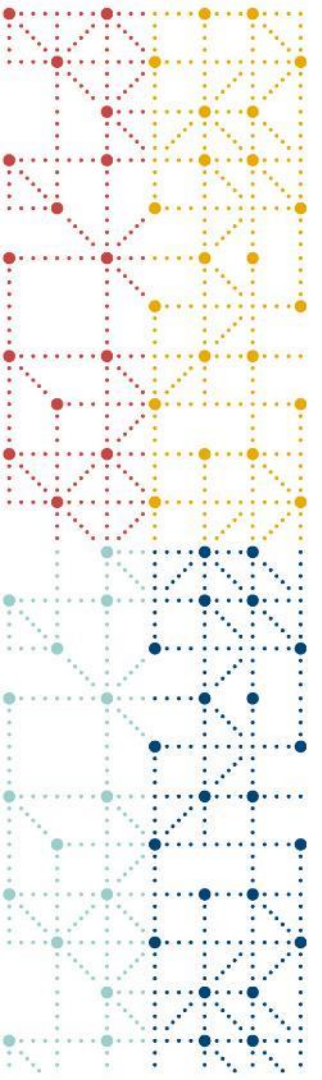
standardize_groupings.sas

```
proc freq data = &dsin.;  
  tables &tabreq. / chisq;  
  output out = rawres chisq pchi;  
run;
```

VIEWTABLE: Work.Rawres

| | N | _PCHI_ | DF_PCHI | P_PCHI | |
|---|-----|-----------|---------|-----------|--|
| 1 | 254 | 1.7166152 | 2 | 0.4238788 | |

```
/* Set the result column to use;  
%local rescol;  
%if %bquote(&oplabel.) = chisq %then %let rescol = _PCHI_;  
%else %if %bquote(&oplabel.) = df %then %let rescol = DF_PCHI;  
%else %if %bquote(&oplabel.) = %str(p-value) %then %let rescol = P_PCHI;
```

Usage example

Comparison of Age Group by Treatment

| VIEWTABLE: Jsonmd.Analyses | | | | | | | | |
|----------------------------|---------------------------|---------|--|-------------|--|------------------|-------------------------|----------|
| | id | version | name | description | categorylds | reason | purpose | analysis |
| 1 | An01_05_SAF_Summ_ByTrt | 1 | Summary of Subjects by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 2 | An03_01_Age_Summ_ByTrt | 1 | Summary of Age by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 3 | An03_01_Age_Comp_ByTrt | 1 | Comparison of Age by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 4 | An03_02_AgeGrp_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Age Group | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 5 | An03_02_AgeGrp_Comp_ByTrt | 1 | Comparison of Age Group by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 6 | An03_03_Sex_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Sex | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 7 | An03_03_Sex_Comp_ByTrt | 1 | Comparison of Sex by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 8 | An03_04_Ethnic_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Ethnicity | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 9 | An03_04_Ethnic_Comp_ByTrt | 1 | Comparison of Ethnicity by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 10 | An03_05_Race_Summ_ByTrt | 1 | Summary of Subjects by Treatment and Race | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 11 | An03_05_Race_Comp_ByTrt | 1 | Comparison of Race by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 12 | An03_06_Height_Summ_ByTrt | 1 | Summary of Height by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 13 | An03_06_Height_Comp_ByTrt | 1 | Comparison of Height by Treatment | | Catn_01_Grp_1_Pop Catn_02_Dclass_1_Sbj Catn_03_SbjDType_1_Dm | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |
| 14 | An07_01_TEAE_Summ_ByTrt | 1 | Summary of Subjects with At Least One TEAE, by Treatment | | Catn_01_Grp_2_Saf Catn_02_Dclass_2_Evt Catn_04_EvtDType_1_Ae Catn_05_PatType_1_Co | SPECIFIED IN SAP | PRIMARY OUTCOME MEASURE | Analysis |

Comparison of Age Group by Treatment

| Variable | Value |
|-----------------|--------------------------------------|
| Id | An03_02_AgeGrp_Comp_ByTrt |
| Name | Comparison of Age Group by Treatment |
| Reason | SPECIFIED IN SAP |
| Purpose | PRIMARY OUTCOME MEASURE |
| analysisSetId | AnalysisSet_02_SAF |
| groupingId1 | AnlsGrouping_01_Trt |
| resultsByGroup1 | FALSE |
| groupingId2 | AnlsGrouping_03_AgeGp |
| resultsByGroup2 | FALSE |
| Dataset | ADSL |
| Variable | USUBJID |
| Method_id | Mth03_CatVar_Comp_PChiSq |

Analysis sets

VIEWTABLE: Jsonmd.Analysissets

| | id | label | compndExpression_logi | condition_dataset | condition_variable | condition_comparator | condition_value | level | order |
|---|--------------------|----------------------------|-----------------------|-------------------|--------------------|----------------------|-----------------|-------|-------|
| 1 | AnalysisSet_01_ITT | Intent-to-Treat Population | | ADSL | ITTFL | EQ | Y | 1 | 1 |
| 2 | AnalysisSet_02_SAF | Safety Population | | ADSL | SAFFL | EQ | Y | 1 | 1 |

```
* Derive expressions in SAS syntax for conditions;|
%build_expression_all(dsin=jsonmd.analysissets, datalib=adam,
  dsout=jsonmd.expressions);
%build_expression_all(dsin=jsonmd.datasubsets, datalib=adam,
  dsout=jsonmd.expressions);
%build_expression_all(dsin=jsonmd.analysisgroupings, idvar=group_id,
  labelvar=group_label, levelvar=group_level,
  ordervar=order, logopvar=group_logicalOperator,
  dsvar=group_condition_dataset, varvar=group_condition_variable,
  compvar=group_condition_comparator, valvar=condition_value,
```

VIEWTABLE: Jsonmd.Expressions

| | id | label | dsconds | expression |
|---|--------------------|----------------------------|---------|--------------|
| 1 | AnalysisSet_01_ITT | Intent-to-Treat Population | ADSL | ITTFL EQ "Y" |
| 2 | AnalysisSet_02_SAF | Safety Population | ADSL | SAFFL EQ "Y" |

Analysis groupings

| VIEWTABLE: Jsonmd.Analysisgroupings | | | | | | | | | | | | | |
|-------------------------------------|---------------|------------------------|--------------------|---------------|------------|--------------------------|---|-------------|---------------|-------------|------------|---|------|
| | grouping_type | id | label | groupingVaria | dataDriven | group_id | group_label | group_logic | group_condtio | group_condi | group_cond | condition_value | gr ^ |
| 1 | Subject | AnlsGrouping_01_Trt | Treatment | TRT01A | FALSE | AnlsGrouping_01_Trt_1 | Placebo | | ADSL | TRT01A | EQ | Placebo | 1 |
| 2 | Subject | AnlsGrouping_01_Trt | Treatment | TRT01A | FALSE | AnlsGrouping_01_Trt_2 | Xanomeline Low Dose | | ADSL | TRT01A | EQ | Xanomeline Low Dose | 1 |
| 3 | Subject | AnlsGrouping_01_Trt | Treatment | TRT01A | FALSE | AnlsGrouping_01_Trt_3 | Xanomeline High Dose | | ADSL | TRT01A | EQ | Xanomeline High Dose | 1 |
| 4 | Subject | AnlsGrouping_02_Sex | Gender | SEX | FALSE | AnlsGrouping_02_Sex_1 | Male | | ADSL | SEX | EQ | M | 1 |
| 5 | Subject | AnlsGrouping_02_Sex | Gender | SEX | FALSE | AnlsGrouping_02_Sex_2 | Female | | ADSL | SEX | EQ | F | 1 |
| 6 | Subject | AnlsGrouping_03_AgeGp | Age Group | AGEGR1 | FALSE | AnlsGrouping_03_AgeGp_1 | < 65 years | | ADSL | AGEGR1 | EQ | <65 | 1 |
| 7 | Subject | AnlsGrouping_03_AgeGp | Age Group | AGEGR1 | FALSE | AnlsGrouping_03_AgeGp_2 | ≥ 65 years | | ADSL | AGEGR1 | IN | 65-80 >80 | 1 |
| 8 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_1 | American Indian or Alaska Native | | ADSL | RACE | EQ | AMERICAN INDIAN OR ALASKA NATIVE | 1 |
| 9 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_2 | Asian | | ADSL | RACE | EQ | ASIAN | 1 |
| 10 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_3 | Black or African American | | ADSL | RACE | EQ | BLACK OR AFRICAN AMERICAN | 1 |
| 11 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_4 | Native Hawaiian or Other Pacific Islander | | ADSL | RACE | EQ | NATIVE HAWAIIAN OR OTHER PACIFIC ISLANDER | 1 |
| 12 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_5 | White | | ADSL | RACE | EQ | WHITE | 1 |
| 13 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_6 | Multiple | | ADSL | RACE | EQ | MULTIPLE | 1 |
| 14 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_7 | Not Reported | | ADSL | RACE | EQ | NOT REPORTED | 1 |
| 15 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_8 | Unknown | | ADSL | RACE | EQ | UNKNOWN | 1 |
| 16 | Subject | AnlsGrouping_04_Race | Primary Race | RACE | FALSE | AnlsGrouping_04_Race_9 | Other | | ADSL | RACE | EQ | OTHER | 1 |
| 17 | Subject | AnlsGrouping_05_Ethnic | Ethnicity | ETHNIC | FALSE | AnlsGrouping_05_Ethnic_1 | Hispanic or Latino | | ADSL | ETHNIC | EQ | HISPANIC OR LATINO | 1 |
| 18 | Subject | AnlsGrouping_05_Ethnic | Ethnicity | ETHNIC | FALSE | AnlsGrouping_05_Ethnic_2 | Not Hispanic or Latino | | ADSL | ETHNIC | EQ | NOT HISPANIC OR LATINO | 1 |
| 19 | Data | AnlsGrouping_06_Soc | System Organ Class | AESOC | TRUE | AnlsGrouping_06_Soc | System Organ Class | | | | | | |
| 20 | Data | AnlsGrouping_07_Pt | Preferred Term | AEDECOD | TRUE | AnlsGrouping_07_Pt | Preferred Term | | | | | | |
| 21 | Data | AnlsGrouping_08_Param | Parameter | PARAMCD | FALSE | AnlsGrouping_08_Param_1 | Systolic Blood Pressure (mmHg) | | ADVS | PARAMCD | EQ | SYSBP | 1 |
| 22 | Data | AnlsGrouping_08_Param | Parameter | PARAMCD | FALSE | AnlsGrouping_08_Param_2 | Diastolic Blood Pressure (mmHg) | | ADVS | PARAMCD | EQ | DIABP | 1 |
| 23 | Data | AnlsGrouping_08_Param | Parameter | PARAMCD | FALSE | AnlsGrouping_08_Param_3 | Pulse Rate (beats/min) | | ADVS | PARAMCD | EQ | PULSE | 1 |
| 24 | Data | AnlsGrouping_08_Param | Parameter | PARAMCD | FALSE | AnlsGrouping_08_Param_4 | Temperature (C) | | ADVS | PARAMCD | EQ | TEMP | 1 |

Output in ARD

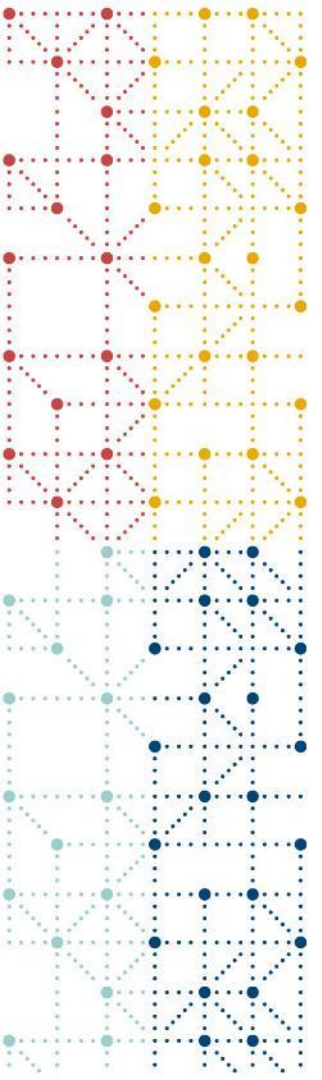
| VIEWTABLE: Ard.Ard | | | | | | | | |
|--------------------|---------------------------|--------------------------------------|--------------------|-------------------|--------------|-----------------|-----------|------------------|
| | analysisId | analysisName | analysisSetId | analysisSetLabel | dataSubsetId | dataSubsetLabel | analysisD | analysisVariable |
| 29 | An03_02_AgeGrp_Comp_ByTrt | Comparison of Age Group by Treatment | AnalysisSet_02_SAF | Safety Population | | | ADSL | USUBJID |
| 30 | An03_02_AgeGrp_Comp_ByTrt | Comparison of Age Group by Treatment | AnalysisSet_02_SAF | Safety Population | | | ADSL | USUBJID |
| 31 | An03_02_AgeGrp_Comp_ByTrt | Comparison of Age Group by Treatment | AnalysisSet_02_SAF | Safety Population | | | ADSL | USUBJID |
| 32 | An03_02_AgeGrp_Summ_ByTrt | Summary of Subjects by Treatment and | AnalysisSet_02_SAF | Safety Population | | | ADSL | USUBJID |

| VIEWTABLE: Ard.Ard | | | | | |
|--------------------|--------------------------|---|----------------------------------|--------------------|---|
| | methodId | methodName | operationId | operationName | r |
| 29 | Mth03_CatVar_Comp_PChiSq | Pearson's chi-square test group comparison for a categorical variable | Mth03_CatVar_Comp_PChiSq_1_chisq | Chi-squared | / |
| 30 | Mth03_CatVar_Comp_PChiSq | Pearson's chi-square test group comparison for a categorical variable | Mth03_CatVar_Comp_PChiSq_2_df | Degrees of freedom | / |
| 31 | Mth03_CatVar_Comp_PChiSq | Pearson's chi-square test group comparison for a categorical variable | Mth03_CatVar_Comp_PChiSq_3_pval | P-value | / |

| VIEWTABLE: Ard.Ard | |
|--------------------|-------------------------|
| | resultGroup1_groupingId |
| 29 | AnlsGrouping_01_Tr |
| 30 | AnlsGrouping_01_Tr |
| 31 | AnlsGrouping_01_Tr |

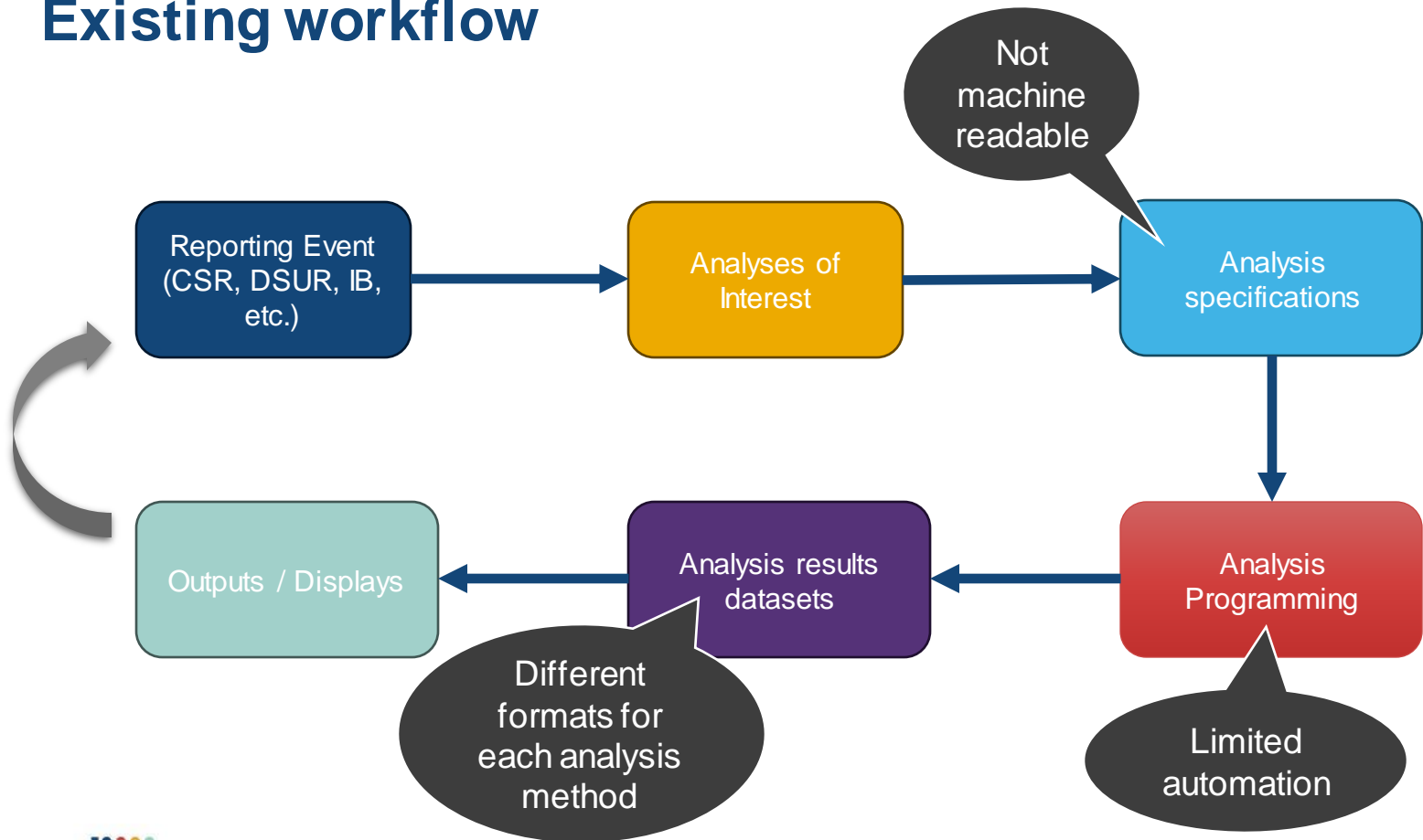
| VIEWTABLE: Ard.Ard | |
|--------------------|-------------------------|
| | resultGroup2_groupingId |
| 29 | AnlsGrouping_03_AgeGp |
| 30 | AnlsGrouping_03_AgeGp |
| 31 | AnlsGrouping_03_AgeGp |

| rawValue |
|--------------|
| 1.7166151981 |
| 2 |
| 0.4238788486 |

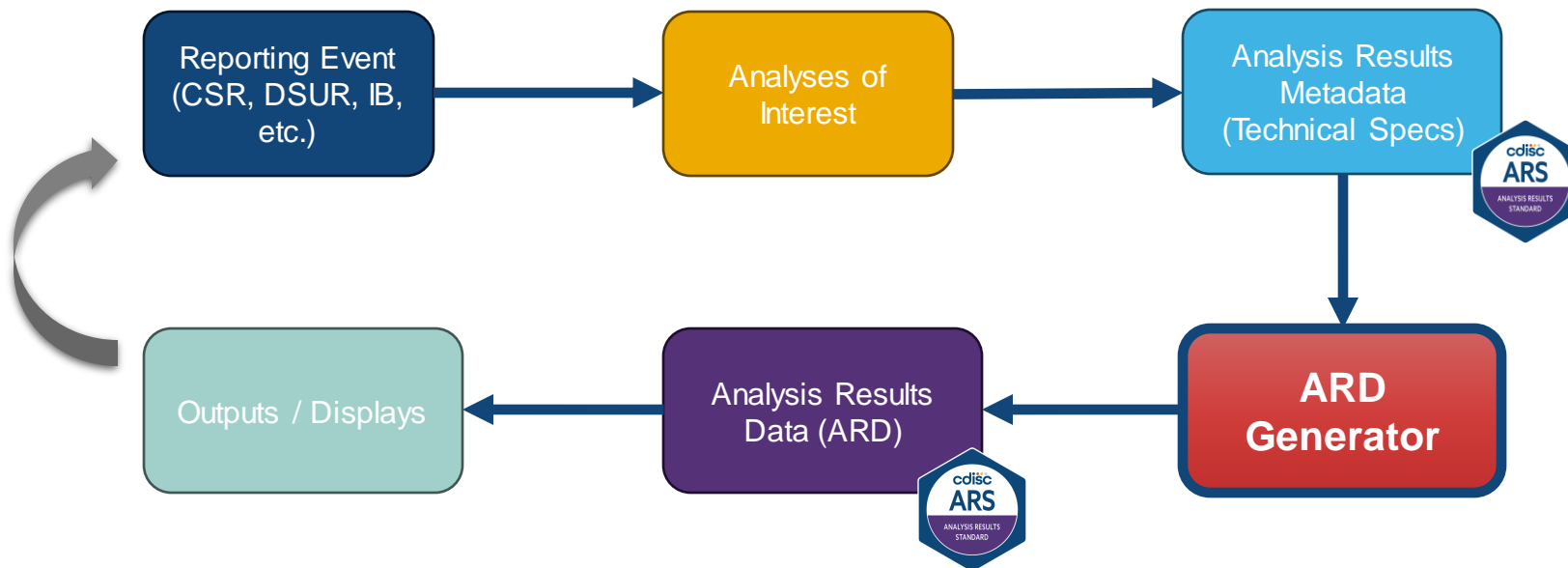


Conclusions

Existing workflow



Future workflow



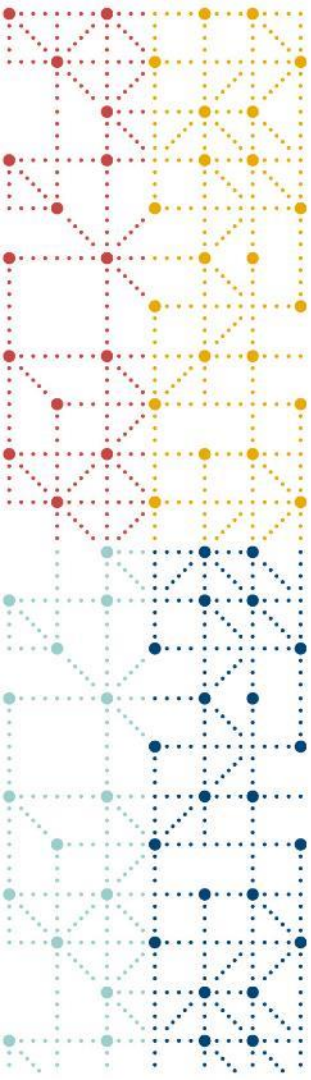


Future development

- Handle more analysis operations and methods.
- Implement display patterns.
- Output ARD as JSON
- Complement ARD Generator with other tools

Summary

- The Analysis Results Standard (ARS) extends the coverage of CDISC standards to the representation of analysis results in their full context.
- Our SAS tool, ARD Generator, makes use of the ARS to automate the generation of results.
- The tool ingests ARM in JSON format and maps its deep hierarchical structure to SAS datasets. It then uses a macro library to perform the specified analyses and writes results to an ARS-compliant Analysis Result Dataset (ARD).
- The tool is highly modular with much re-use of code. It largely automates the analysis process, as once the relevant macros exist, all the information needed to apply them can be taken from the ARM.
- ARD Generator came out of work for a CDISC Hackathon.



Thank You!

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<https://github.com/KarlWallendszus/ard-generator>

cdisc

