

解析結果メタデータの概要と SASによる作成方法の提案

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Overview and Creation of
Analysis Results Metadata with SAS

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要旨：

医薬品製造販売承認申請時の電子データ提出において、ADaMの Define.xmlに含めて提出することが推奨されている解析結果メタデータの概要と実装時の留意事項、ならびにSASによる作成方法を提案する。

For e-Data submission for Japan NDA, Analysis Results Metadata (ARM) which is recommended to be included in ADaM Define.xml and submitted to PMDA. In this presentation, overview and considerations of ARM will be summarized and the creation process using SAS will be proposed.

キーワード: CDISC, ADaM, Analysis Results Metadata, Define.xml

Outline

- **Background**
 - Analysis Results Metadata for e-Data submission in Japan
- **Overview of Analysis Results Metadata**
 - Analysis Results Metadata in ADaM
 - Structure of Analysis Results Metadata
 - Considerations on Analysis Results Metadata
- **Creation of Analysis Results Metadata with SAS**
 - Excel Metadata File and SAS program to generate Analysis Results Metadata as a part of Define.xml

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Background



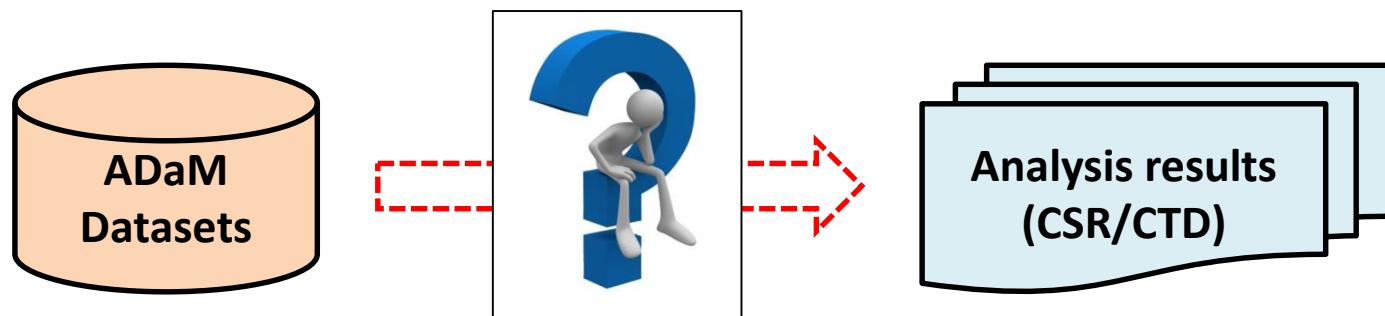
- New guidance and Technical Conformance Guide in Japan
 - PMDA will start to receive e-Data of clinical studies from Oct. 2016 (transition period is 3.5 years)
 - For mainly Ph-II, Ph-III and specific Ph-I/Clinical Pharmacology studies, sponsors have to submit the CDISC-compliant datasets (SDTM and ADaM), Define.xml, Reviewer's Guide and Annotated CRF
 - For Ph-I/CP studies, sponsors have to discuss the targeted studies and the contents of e-Data with PMDA at the consultation meetings
 - PMDA recommends sponsors to submit the Analysis Results Metadata (ARM) (Technical Conformance Guide).
 - Not for all analyses (e.g. primary and secondary analyses)
 - If it is difficult to prepare the ARM, sponsors need to prepare the other documents/specifications for the explanation of analyses

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What is the Analysis Results Metadata (ARM)?

- One of the Metadata in ADaM (V2.1)
 - Submitted to PMDA as a part of Define.xml for ADaM
 - Analysis Results Metadata V1.0 for Define.xml V2.0 was released in Jan., 2015
 - Provide traceability between analysis results and ADaM datasets (e.g. datasets, variables, selection criteria, programming code used in the analyses)
 - Reviewers can understand how the analysis results are created.



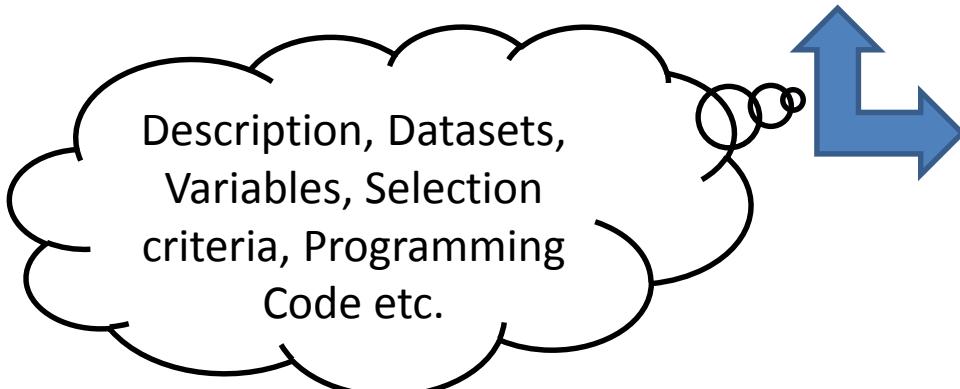
Traceability with ARM

- Between Analysis Results and ADaM datasets

Protocol: CDISCPILOT01 Population: Efficacy			Page 1 of 1
Table 14-3.11 ADAS Cog (11) - Repeated Measures Analysis of Change from Baseline to Week 24			
	Placebo (N=79)	Isoniazide Low Dose (N=81)	Isoniazide High Dose (N=74)
LS Means (SE)	1.6 (0.49)	1.5 (0.52)	1.1 (0.56)
p-value (Xan - Placebo)		0.555	0.556
Diff of LS Means (SE)		-0.0 (0.70)	-0.4 (0.72)
95% CI		(-1.4;1.3)	(-1.9;1.0)
p-value (Xan High - Xan Low)			0.606
Diff of LS Means (SE)			-0.4 (0.75)
95% CI			(-1.9;1.1)

Metadata Field	Definition of field	Metadata
DISPLAY IDENTIFIER	Unique identifier for the specific analysis display	Table 14-3.11
DISPLAY NAME	Title of display	ADAS Cog (11) - Repeated Measures Analysis of Change from Baseline to Week 24
RESULT IDENTIFIER	Identifies the specific analysis result within a display	
PARAM	Analysis parameter	ADAS-Cog (11) Total Score
PARAMCD	Analysis parameter code	ACTOT11
ANALYSIS VARIABLE	Analysis variable being analyzed	CHG
REASON	Rationale for performing this analysis	Pre-specified in SAP
DATASET	Dataset(s) used in the analysis.	ADQSADAS
SELECTION CRITERIA	Specific and sufficient selection criteria for analysis subset and / or numerator	ITFL='Y' and AVISITN GT 0 AND DTYPE NE 'LOCF' AND PARAMCD='ACTOT11'
DOCUMENTATION	Textual description of the analysis performed	SAP Section 10.1.1. Adjusted means for the change from baseline at week 24 and pairwise comparisons between treatment groups at Week 24 using a repeated measures model with treatment group (as class variable); site (as class variable); time; treatment*time interaction; baseline score and baseline*time interaction terms; and an unstructured covariance matrix. Efficacy data, observed cases data.
PROGRAMMING STATEMENTS	The analysis syntax used to perform the analysis	PROC MIXED; CLASS USUBJID SITEGR1 AVISITN TRTP; MODEL CHG = TRTP SITEGR1 AVISITN TRTP*AVISITN BASE BASE*AVISITN / OUTP=PRED DDFM=KR; REPEATED AVISITN / SUBJECT=USUBJID TYPE=UN; LSMEANS TRTP / DIFF CL; RUN;

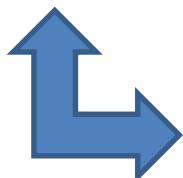
Description, Datasets,
Variables, Selection
criteria, Programming
Code etc.



ARM in the ADaM Define.xml

- Specific structure in Define.xml
 - Output Define.xml on a browser with XML stylesheet

Metadata Field	Definition of field	Metadata
DISPLAY IDENTIFIER	Unique identifier for the specific analysis display	Table 14-3.11
DISPLAY NAME	Title of display	ADAS Cog (11) - Repeated Measures Analysis of Change from Baseline to Week 24
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PARAM	Analysis parameter	ADAS-Cog (11) Total Score
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ANALYSIS VARIABLE	Analysis variable being analyzed	CHG
REASON	Rationale for performing this analysis	Pre-specified in SAP
DATASET	Dataset(s) used in the analysis.	ADQSADAS
SELECTION CRITERIA	Specific and sufficient selection criteria for analysis subset and / or numerator	ITTFL=Y and AVISITN GT 0 AND DTYPE NE 'LOCF' AND PARAMCD='ACTOT11'
DOCUMENTATION	Textual description of the analysis performed	SAP Section 10.1.1. Adjusted means for the change from baseline at week 24 and pairwise comparisons between treatment groups at Week 24 using a repeated measures model with treatment group (as class variable), site (as class variable), time, treatment*time interaction; baseline score and baseline*time interaction terms; and an unstructured covariance matrix. Efficacy data, observed cases data.
PROGRAMMING STATEMENTS	The analysis syntax used to perform the analysis	PROC MIXED; CLASS USUBJID SITEGR1 AVISITN TRTP; MODEL CHG = TRTP SITEGR1 AVISITN TRTP*AVISITN BASE BASE*AVISITN / OUTP=PRED DFDM=KR; REPEATED AVISITN / SUBJECT=USUBJID TYPE=UN; LSMEANS TRTP / DIFF CL; RUN;



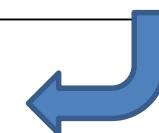
```

<arm:AnalysisResultDisplay>
  <arm:ResultDisplay OID="RD.Table_14-3.01" Name="Table 14-3.01">
    <Description>
      <TranslatedText xml:lang="en">Primary Endpoint Analysis: ADAS-Cog - Summary at Week 24 - LOCF (Efficacy Population)</TranslatedText>
    </Description>
    <Def:DocumentRef leafID="LF.Table_14-3.01">
      <Def:PDFPageRef PageRefs="2" Type="PhysicalRef"/>
    </Def:DocumentRef>
    <arm:AnalysisResult>
      <OID>AR.Table_14-3.01.R.1</OID>
      <ParameterOID>IT.ADQSADAS.PARAMOD</ParameterOID>
      <AnalysisReason>SPECIFIED IN SAP</AnalysisReason>
      <AnalysisPurpose>PRIMARY OUTCOME MEASURE</AnalysisPurpose>
      <Description>
        <TranslatedText xml:lang="en">Dose response analysis for ADAS-Cog changes from baseline</TranslatedText>
      </Description>
      <arm:AnalysisDataset>
        <arm:AnalysisDatasetItem ItemGroupOID="IG.ADQSADAS" />
        <Def:WhereClauseRef WhereClauseOID="WC.Table_14-3.01.R.1.ADQSADAS" />
        <arm:AnalysisVariable ItemOID="IT.ADQSADAS.CHG" />
      </arm:AnalysisDataset>
      <arm:Documentation>
        <Description>
          <TranslatedText xml:lang="en">Linear model analysis of CHG for dose response; using randomized dose (0 for placebo; 54 for low dose; 81 for high dose) and site group in model. Used PROC GLM in SAS to produce p-value (from Type III SS for treatment dose).</TranslatedText>
        </Description>
        <Def:DocumentRef leafID="LF.SAP-SEC-10.1.1">
          <Def:PDFPageRef PageRefs="4" Type="PhysicalRef"/>
        </Def:DocumentRef>
      </arm:Documentation>
      <arm:ProgrammingCode Context="SAS version 9.2">

```

Table 14-3.01

Display	Table 14-3.01 Primary Endpoint Analysis: ADAS-Cog - Summary at Week 24 - LOCF (Efficacy Population)
Analysis Parameter(s)	Dose response analysis for ADAS-Cog changes from baseline
Analysis Variable(s)	CHG (Change from Baseline)
Analysis Reason	Pre-specified in SAP
Analysis Purpose	Primary Outcome Measure
Data References (incl. Selection Criteria)	ADQSADAS [PARAMCD = "ACTOT" and AVISIT = "Week 24" and EFFFL = "Y" and ANL01FL = "Y"]
Documentation	Linear model analysis of CHG for dose response; using randomized dose (0 for placebo; 54 for low dose; 81 for high dose) and site group in model. Used PROC GLM in SAS to produce p-value (from Type III SS for treatment dose). SAP Section 10.1.1
Programming Statements	[SAS version 9.2 or later] <pre> proc glm data = ADQSADAS; where EFFFL='Y' and ANL01FL='Y' and AVISIT='Week 24' and PARAMCD="ACTOT"; class SITEGR1; model CHG = TRTPN SITEGR1; run; </pre>



Browser
(with Stylesheet)

Structure of Analysis Results Metadata

- Analysis Results Metadata V1.0 for Define.xml V2.0



Analysis Results Metadata Specification Version 1.0 for Define-XML Version 2

Prepared by
CDISC ADaM Metadata Sub-Team

Notes to Readers

- This is the specification for the Analysis Results Metadata extension of Version 2.0 of the CDISC Define-XML standard.
- This document is intended for companies and individuals involved in the analysis of clinical data that will be submitted to regulatory authorities.

Revision History

Date	Version	Summary of Changes
2015-01-27	1.0	Final version 1.0 incorporating all changes identified during the public comment period
2014-09-10	1.0 DRAFT	Draft for public review

Warranties Notice: see Appendix C

```

<?xml version="1.0" encoding="UTF-8"?>
<ODM xmlns="http://www.cdisc.org/ns/odm/v1.3"
      xmlns:def="http://www.cdisc.org/ns/def/v2.0"
      xmlns:xlink="http://www.w3.org/1999/xlink"
      xmlns:arm="http://www.cdisc.org/ns/arm/v1.0"
      ODMVersion="1.3.2" fileType="Snapshot" fileOID="CDISC-Sample"
      CreationDate="2014-03-28T11:07:23:00"
      originator="CDISC ADaM Metadata Team">
  <Study OID="cdisc01">
    <GlobalVariables>
      <StudyName>CDISC Sample</StudyName>
      <StudyDescription>CDISC-Sample Data Definitions</StudyDescription>
      <ProtocolName>CDISC-Sample</ProtocolName>
    </GlobalVariables>
    <MetaDataVersion OID="MDV.CDISC01.ADaMIG.1.0.ADaM.2.1">
      Name="Study CDISC-Sample, Data Definitions"
      Description="Study CDISC01, Data Definitions"
      def:DefineVersion="2.0.0"
      def:StandardName="ADaM-IG"
      def:StandardVersion="1.0">
      <Supplemental Data Definitions (def:SupplementalDoc) >
      <Value Level Metadata (def:ValueListDef) >
      <Where Clause Definitions (def:WhereClauseDef) >
      <Domain Level Metadata (ItemGroupDef) >
      <Variable Level Metadata (ItemDef) >
      <Controlled Terminology Metadata (CodeList) >
      <Computational Algorithms (MethodDef) >
      <Comments (def:CommentDef) >
      <Referenced Documents (def:leaf) >
      <Analysis Results Metadata (arm:AnalysisResultDisplays) >
    </MetaDataVersion>
  </Study>
</ODM>

```

Structure of Analysis Results Metadata

- Tagsets under the <arm:AnalysisResultDisplays>

```
<arm:AnalysisResultDisplays>
  <arm:ResultDisplay>
    <Description>
      <TranslatedText>
    <def:DocumentRef>
      <def:PDFPageRef>
    <arm:AnalysisResult>
      <Description>
        <TranslatedText>
      <arm:AnalysisDatasets>
        <arm:AnalysisDataset>
          <def:WhereClauseRef>
          <arm:AnalysisVariable>
        <arm:Documentation>
          <Description>
            <TranslatedText>
          <def:DocumentRef>
            <def:PDFPageRef>
        <arm:ProgrammingCode>
          <arm:Code>
          <def:DocumentRef>
            <def:PDFPageRef>
```



Structure of Analysis Results Metadata

- <arm:ResultDisplay>
 - Corresponds to a table/figure in CSR

```
<arm:AnalysisResultDisplays>
  <arm:ResultDisplay OID="RD.Table_14-3.01" Name="Table 14-3.01"> Analysis result display
    <Description>
      <TranslatedText xml:lang="en">Primary Endpoint Analysis: ADAS-Cog – Summary at
      Week 24 – LOCF (Efficacy Population)
      </TranslatedText>
    </Description>
    <def:DocumentRef leafID="LF.Table-14-3.01"> Link to tables in CSR
      <def:PDFPageRef PageRefs="49" Type="PhysicalRef"/>
    </def:DocumentRef>
    <arm:AnalysisResult>
      ...
    </arm:AnalysisResult> Analysis result (multiple results can be
    included in one table/figure)
    <arm:AnalysisResult>
      ...
    </arm:AnalysisResult>
  </arm:ResultDisplay>
  ...
</arm:AnalysisResultDisplays>
```

Structure of Analysis Results Metadata

- <arm:AnalysisResult>
 - Corresponds to an analysis result

<arm:ResultDisplay OID="RD.Table_14-3.01" Name="Table 14-3.01">

...

<arm:AnalysisResult OID="AR.Table_14-3.01.R.1" ParameterOID="IT.ADQSADAS.PARAMCD" AnalysisReason="SPECIFIED IN SAP" AnalysisPurpose="PRIMARY OUTCOME MEASURE">

<Description>

<TranslatedText xml:lang="en">Dose response analysis for ADAS-Cog changes from baseline</TranslatedText>

</Description>

<arm:AnalysisDatasets>

...

</arm:AnalysisDatasets>

<arm:Documentation>

...

</arm:Documentation>

<arm:ProgrammingCode>

...

</arm:ProgrammingCode>

</arm:AnalysisResult>

<arm:AnalysisResult>

...

</arm:AnalysisResult>

</arm:ResultDisplay>

Analysis Result Display

Analysis result (multiple results can be included in one table/figure)

Datasets (multiple datasets can be described)

Link to the specification of an analysis

Analysis Program (code or link to program file)

Analysis Results Metadata (Summary) for Study CDISC-Sample

Table 14-3.01 Primary Endpoint Analysis: ADAS-Cog - Summary at Week 24 - LOCF (Efficacy Population)

Dose response analysis for ADAS-Cog changes from baseline

Pairwise comparisons to placebo for ADAS-Cog changes from baseline

Adverse Events by Treatment Group

As Adverse Events by Treatment Group

Detail) for Study CDISC-Sample

Table 14-3.01 Primary Endpoint Analysis: ADAS-Cog - Summary at Week 24 - LOCF (Efficacy Population)

Dose response analysis for ADAS-Cog changes from baseline

e) PARAMCD = "ACTOT" (Adas-Cog(11) Subscore)

CHG (Change from Baseline)

Analysis Reason

SPECIFIED IN SAP

MEASURE

PARAMCD = "ACTOT" and AVISIT = "Week 24" and EFFFL = "Y" and ANL01FL = "Y"]

ysis of CHG for dose response; using randomized dose (0 for placebo; 54 for low dose; 81 for high dose) and site group in model. Used to produce p-value (from Type III SS for treatment dose).

SAP Section 10.1.1

Programming Statements

```
[SAS version 9.2]
proc glm data = ADQSADAS;
  where EFFFL='Y' and ANL01FL='Y' and AVISIT='Week 24' and PARAMCD="ACTOT";
  class SITEGR1;
  model CHG = TRTPN SITEGR1;
run;
```

Structure of Analysis Results Metadata

- <arm:AnalysisDatasets> and <arm:AnalysisDataset>
 - Corresponds to datasets, variables, selection criteria

<arm:AnalysisResultDisplays>

...

<arm:AnalysisDatasets def:CommentOID="COM. JOIN-ADSL-ADAE">

<arm:AnalysisDataset ItemGroupOID="IG. ADAE">

<def:WhereClauseRef WhereClauseOID="WC. Table_14-5. 02. R. 1. ADAE" />

<arm:AnalysisVariable ItemOID="IT. ADAE. AEBODSYS"/>

<arm:AnalysisVariable ItemOID="IT. ADAE. AEDECOD"/>

</arm:AnalysisDataset>

...

<arm:AnalysisDataset ItemGroupOID="IG. ADSL">

...

</arm:AnalysisDataset>

</arm:AnalysisDatasets>

...

</arm:AnalysisResultDisplays>

If multiple def:WhereclauseRef are described,
each criteria is combined using "or" operator

Adverse Events by Treatment Group by Treatment Group	
Analysis Parameter(s)	
Analysis Variable(s)	AEBODSYS (Body System or Organ Class) AEDECOD (Dictionary-Derived Term)
Analysis Reason	SPECIFIED IN SAP
Analysis Purpose	PRIMARY OUTCOME MEASURE
Data References (incl. Selection Criteria)	ADAE [TRTEML = "Y" and AESER = "Y"] ADSL [SAFFL = "Y"] Get denominators for percentages from ADSL and counts and numerators from ADAE. Join ADAE with ADSL based on the unique subject identifier (USUBID) keeping only records in ADAE for the numerator.
Documentation	Unique count of subjects that experienced an Adverse Event by Preferred Term, System Organ Class, and Treatment Group and percentages based on the number of subjects in the safety population within each treatment group. The total number of times an event occurred was recorded by Preferred Term, System Organ Class, and Treatment Group. Fisher's exact test was used for treatment comparison of event rates. SAP Section 11.2
Programming Statements	[SAS version 9.2] at14-5-02.sas

Structure of Analysis Results Metadata

- <arm:ProgrammingCode>
 - 1. Actual Code using <arm:Code>

```
<arm:ProgrammingCode Context=" SAS version 9.2 or later">
```

```
<arm:Code>
```

```
proc glm data = ADQSADAS;
  where EFFF1='Y' and ANL01FL='Y' and AVISIT='Week 24' and PARAMCD="ACTOT";
  class TRTPN SITEGR1;
  model CHG = TRTPN SITEGR1 BASE;
  means TRTPN;
  lsmeans TRTPN / OM STDERR PDIFF CL;
run;
```

```
</arm:Code>
```

```
</arm:ProgrammingCode>
```

Actual Code

Program Code/File

Programming Statements

[SAS version 9.2]

```
proc glm data = ADQSADAS;
  where EFFF1='Y' and ANL01FL='Y' and AVISIT='Week 24' and PARAMCD="ACTOT";
  class TRTPN SITEGR1;
  model CHG = TRTPN SITEGR1 BASE;
  means TRTPN;
  lsmeans TRTPN / OM STDERR PDIFF CL;
run;
```

- 2. Link to Program file using <def:DocumentRef>

Definition of external file

```
<def:leaf ID="LF.at14-5-02.sas" xlink:href="..../programs/at14-5-02-sas.txt">
```

```
  <def:title>at14-5-02.sas</def:title>
```

```
</def:leaf>
```

```
...
```

```
<arm:ProgrammingCode Context=" SAS version 9.2 or later">
```

```
  <def:DocumentRef leafID="LF.at14-5-02.sas" />
```

```
</arm:ProgrammingCode>
```

Program Code/File

Link to external file

Considerations on Analysis Results Metadata

- **Creation** of ARM in Define.xml
 - Need basic knowledge for XML structure
 - [XML schema, well-formed /valid XML, XML stylesheet](#)
 - Difficult to create ARM automatically without standardization
 - Several tagset patterns
 - e.g. multiple results/datasets/variables, multiple Whereclause elements
 - [Define.xml Generator \(Fujitsu\)](#) can generate Define.xml including ARM with Excel metadata
- **Validation**
 - At this point (Jun., 2015), [OpenCDISC](#) cannot validate the ARM
 - Need [schema validation](#)
- **Legacy Data Conversion**
 - If TLFs in CSR are created using [legacy data](#), there will be no analysis programs using ADaM datasets
 - Sponsors may re-create the [analysis programs/results to include them as ARM in Define.xml](#)
 - [For the validation, sponsors may compare the results with TLFs in CSR.](#)



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Creation of Analysis Results Metadata

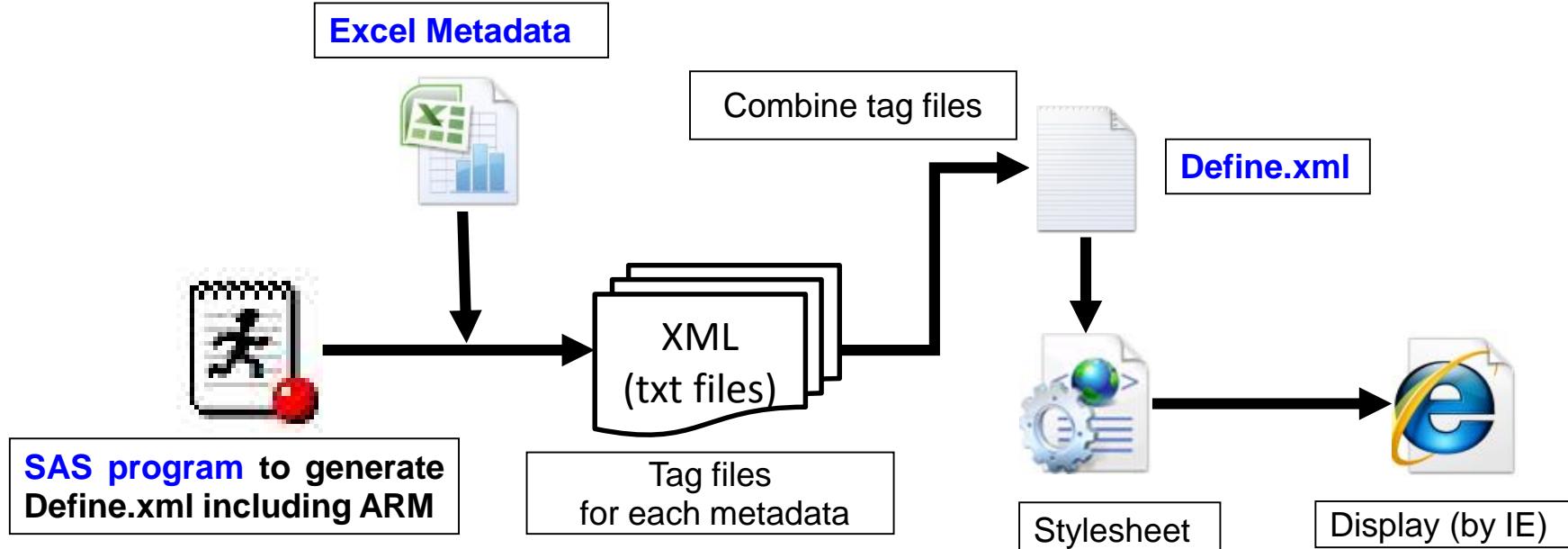
- How to create Analysis Results Metadata
 - Tools
 - OpenCDISC, [Define.xml Generator \(Fujitsu\)](#), Metadata Repository (MDR)
 - Latest Version of Define.xml Generator (Fujitsu) can generate ARM using Excel metadata and validate the Define.xml file using schema file
 - At this point (June 2015), OpenCDISC cannot generate and validate ARM
 - Programs
 - [SAS](#) or other programming languages
 - Read Excel metadata and generate using program
 - » Similar flow to the tools above (e.g. Define.xml Generator)
 - Need programming skill and understanding of the Define.xml structure including ARM



Creation of Analysis Results Metadata **with SAS**

- Excel metadata and SAS program to generate ARM
 - SAS program read the Excel Metadata and Generate the ADaM Define.xml including ARM

***Only the ARM generation part is covered in this presentation and the SAS program to generate the ARM is available on the web site**



Creation of Analysis Results Metadata with SAS

- Excel metadata for ARM (1)
 - Description of analyses and **programming code** or link to the program file

ID	Name	ResultDisplay	Analysis ID	Parameter OID	Reason	Purpose	Description	Documentation
Table_1.1	Table 1.1	Disease XXX Healing Rate at week 12 (FAS)	Table_1.1_1	ADEFF.PARAMCD	SPECIFIED IN SAP	PRIMARY OUTCOME MEASURE	Disease XXX Healing Rate at week 12 (FAS)	Wald test to compare the Disease XXX Healing Rate between treatment groups. FREQ procedure was used for this analysis.
Table_1.1	Table 1.1	Disease XXX Healing Rate at week 4, 8 (FAS)	Table_1.1_2	ADEFF.PARAMCD	SPECIFIED IN SAP	SECONDARY OUTCOME MEASURE	Disease XXX Healing Rate at week 4, 8 (FAS)	Calculate the Disease XXX Healing Rate at Week 4 and Week 8 using FREQ procedure
Table_2.1	Table 2.1	Summary of AE (Safety Analysis Set)	Table_2.1		SPECIFIED IN SAP	SAFETY OUTCOME MEASURE	Summary of AE (Safety Analysis Set)	Summary of TEAE

DocumentRef	ProgrammingCode	Version	ProgramID	href	Title
SAP	<pre> proc format ; value _TRTPF 1 = "Treat A" 2 = "Treat B"; run; proc freq data=ADEF; where FASFL = "Y" and AVISIT = "Week 12" and ANL01FL = "Y"; table TRTPN*AVALC / riskdiff(equal var=sample) nocol nopercent; format TRTPN _TRTPF.; run;</pre>	SAS version 9.2 or later			
SAP	<pre> proc format ; value _TRTPF 1 = "Treat A" 2 = "Treat B"; run; proc freq data=ADEF; where FASFL = "Y" and AVISIT in ("Week 4", "Week 8") and ANL01FL = "Y"; table TRTPN*AVALC / riskdiff(equal var=sample) nocol nopercent; format TRTPN _TRTPF.; run;</pre>	SAS version 9.2 or later			
SAP		SAS version 9.2 or later	AEsas	AE.sas	AE.sas

Creation of Analysis Results Metadata with SAS

- Excel metadata for ARM (2)
 - Description of datasets, variables and selection criteria
 - Multiple datasets and variables can be specified

Analysis ID	WhereClauseOID	Analysis Variables	Dataset	Variable	Comparator	Value
Table_1.1_1	ADEFF.AVAL.HEAL	AVAL	ADSCOP	PARAMCD	EQ	SCHEAL
Table_1.1_1	ADEFF.AVAL.HEAL	AVAL	ADSCOP	FASFL	EQ	Y
Table_1.1_1	ADEFF.AVAL.HEAL	AVAL	ADSCOP	ANL01FL	EQ	Y
Table_1.1_1	ADEFF.AVAL.HEAL	AVAL	ADSCOP	AVISIT	EQ	Week 8
Table_1.1_2	ADEFF.AVAL.HEAL2	AVAL	ADSCOP	PARAMCD	EQ	SCHEAL
Table_1.1_2	ADEFF.AVAL.HEAL2	AVAL	ADSCOP	FASFL	EQ	Y
Table_1.1_2	ADEFF.AVAL.HEAL2	AVAL	ADSCOP	ANL01FL	EQ	Y
Table_1.1_2	ADEFF.AVAL.HEAL2	AVAL	ADSCOP	AVISIT	IN	Week2, Week 4
Table_2.1	ADSL.SAFFL		ADSL	SAFFL	EQ	Y
Table_2.1	ADAE.PT	AEDECOD,AEBODSYS	ADAE	AOCCPFL	EQ	Y
Table_2.1	ADAE.PT	AEDECOD,AEBODSYS	ADAE	SAFFL	EQ	Y

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Preprocessing and Read the Excel metadata using Excel libname engine
 - _MPATH and _CPATH are the full path for Excel metadata file and output text files (xml tagsets)

```
libname META excel "&_MPATH." SCAN_TEXTSIZE=YES ;
data RESULTS_METADATA ;
  set META.'Analysis Results Metadata'$ n ;
  RESNO = _N_ ;
  if ID ne "" ;
run ;
data ARESM_WHR ;
  set META.'AResM Where'$ n ;
  if WhereClauseOID ne "" ;
  WHRNO = _N_ ;
run ;
filename _AR      "&_CPATH.¶temp¥_AR.txt" ;
filename _ANPGMLF "&_CPATH.¶temp¥_ARPGM.txt" ;
filename _ARWHR   "&_CPATH.¶temp¥_ARWHR.txt" ;
proc sort data=RESULTS_METADATA ; by RESNO ID ; run ;
```

Metadata (1)

Metadata (2)

Output files:

1. Tagsets for main part of ARM
2. Link to Program files
3. WhereClause Definition, respectively

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Link to SAS programs

```
data _null_ ;
  set RESULTS_METADATA end=_EOF ;
  file _ANPGMLF ;
  if ProgramID ne "" then do ;
    put '      <def:leaf ID="LF.' ProgramID + (-1) '" xlink:href="" href + (-1) '">' ;
    put '          <def:title>' Title + (-1) '</def:title>' ;
    put '      </def:leaf>' ;
  end ;
run ;
```

Program ID and Path

File name



```
<def:leaf ID="LF.AEsas" xlink:href="AE.sas">
  <def:title>AE.sas</def:title>
</def:leaf>
```

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Generation of WhereClause Definition for ARM

```
proc sort data=ARESM_WHR ;
  by Analysis_ID WhereClauseOID WHRNO ;
run ;

data _null_ ;
  set ARESM_WHR end=_EOF ;
  by Analysis_ID WhereClauseOID WHRNO ;
  file _ARWHR ;

  if first.WhereClauseOID then put '      <def:WhereClauseDef OID="WC.' WhereClauseOID + (-1) '">' ;
  put '          <RangeCheck Comparator=' Comparator + (-1)
    ' SoftHard="Soft" def:ItemOID="IT.' Dataset + (-1) '.' Variable + (-1) '">' ;
  VAL_C = count(Value, ",") + 1 ;
  do I = 1 to VAL_C ;
    VAL = scan(Value, I, ",") ;
    put '          <CheckValue>' VAL + (-1) '</CheckValue>' ;
  end ;
  put '      </RangeCheck>' ;

  if last.WhereClauseOID then put '      </def:WhereClauseDef>' ;
run ;
```

Variables and Operator for WhereClause

For "IN" operator

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Generation of WhereClause Definition for ARM

```
<def:WhereClauseDef OID="WC.ADEFF.AVAL.HEAL">
  <RangeCheck Comparator="EQ" SoftHard="Soft" def:ItemOID="IT.ADEFF.PARAMCD">
    <CheckValue>HEAL</CheckValue>
  </RangeCheck>
  <RangeCheck Comparator="EQ" SoftHard="Soft" def:ItemOID="IT.ADEFF.FASFL">
    <CheckValue>Y</CheckValue>
  </RangeCheck>
  <RangeCheck Comparator="EQ" SoftHard="Soft" def:ItemOID="IT.ADEFF.ANL01FL">
    <CheckValue>Y</CheckValue>
  </RangeCheck>
  <RangeCheck Comparator="EQ" SoftHard="Soft" def:ItemOID="IT.ADEFF.AVISIT">
    <CheckValue>Week 12</CheckValue>
  </RangeCheck>
</def:WhereClauseDef>
....
```

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Main part of ARM

```
proc sort data=RESULTS_METADATA ;
  by Analysis_ID ;
run ;
data _AResM ;
  merge RESULTS_METADATA ARESM_WHR ;
  by Analysis_ID ;
run ;
proc sort data=_AResM ;
  where ID ne "";
  by ID Analysis_ID Dataset WhereClauseOID ;
run ;
data _null_;
  set _AResM end=_EOF ;
  by ID Analysis_ID Dataset WhereClauseOID ;
  file _AR lrecl=1000 ;

  ProgrammingCode = htmlencode(ProgrammingCode) ;

  if _N_ = 1 then put '<arm:AnalysisResultDisplays>' ;
```

Merge the metadata (1) and (2)

Main part of ARM

Creation of Analysis Results Metadata with SAS

```

if first.ID then do ;
put '      <arm:ResultDisplay OID="RD.' ID + (-1) '' '' Name="" Name + (-1) '">' ;
put '          <Description>' ;
put '          <TranslatedText xml:lang="en">' ResultDisplay + (-1) '</TranslatedText>' ;
put '          </Description>' ;
end ;
if first.Analysis_ID then do ; Start of ARM
put '      <arm:AnalysisResult OID="AR.' Analysis_ID + (-1) '' ;
put '                      ParameterOID="IT.' Parameter_OID + (-1) '' ;
put '                      AnalysisReason="" Reason        + (-1) '' ;
put '                      AnalysisPurpose="" Purpose       + (-1) '">' ;
put '          <Description>' ;
put '          <TranslatedText xml:lang="en">' Description + (-1) '</TranslatedText>' ;
put '          </Description>' ;
put '          <arm:AnalysisDatasets>' ;
end ;
if first.Dataset then do ; Dataset, WhereClause and Analysis Variables
put '      <arm:AnalysisDataset ItemGroupOID="IG.' Dataset + (-1) '' >' ;
put '          <def:WhereClauseRef WhereClauseOID="WC.' WhereClauseOID + (-1) ''/>' ;
if Analysis_Variables ne "" then do ;
_NVAL = count(Analysis_Variables, ",") + 1 ;
do I = 1 to _NVAL ;
VARIABLE = strip(scan(Analysis_Variables, I, ","));
put '          <arm:AnalysisVariable ItemOID="IT.' Dataset + (-1) '.' VARIABLE + (-1) ''/>' ;
end ;
end ;
put '      </arm:AnalysisDataset>' ;
end ;

```

Creation of Analysis Results Metadata with SAS

```
if last.Analysis_ID then do ;
  put '                                </arm:AnalysisDatasets>' ;
  put '                                <arm:Documentation>' ;
  put '                                <Description>' ;
  put '                                <TranslatedText xml:lang="en">' Documentation + (-1) '</TranslatedText>' ;
  put '                                </Description>' ;
  if DocumentRef ne "" then do ;
    put '                                <def:DocumentRef leafID="LF.' DocumentRef + (-1) '">' ;
    put '                                </def:DocumentRef>' ;
  end ;
  put '                                </arm:Documentation>' ;
  if ProgrammingCode ne "" then do ;
    put '                                <arm:ProgrammingCode Context="" Version + (-1) '">' ;
    put '                                <arm:Code>' ;
    put ProgrammingCode ;
    put '                                </arm:Code>' ;
    put '                                </arm:ProgrammingCode>' ;
  end ;
  else if ProgramID ne "" then do ;
    put '                                <arm:ProgrammingCode Context="" Version + (-1) '">' ;
    put '                                <def:DocumentRef leafID="LF.' ProgramID + (-1) '">' ;
    put '                                </arm:ProgrammingCode>' ;
  end ;
  put '                                </arm:AnalysisResult>' ;
end ;
```

Description and Documentation

Programming code or Link to program files

Creation of Analysis Results Metadata with SAS

- SAS program for ARM creation
 - Main part of ARM

```
<arm:AnalysisResultDisplays>
  <arm:ResultDisplay OID="RD.Table_1.1" Name="Table 1.1">
    <Description>
      <TranslatedText xml:lang="en">Disease XXX Healing Rate (FAS)</TranslatedText>
    </Description>
    <arm:AnalysisResult OID="AR.Table_1.1_1"
      ParameterOID="IT.ADEFF.PARAMCD"
      AnalysisReason="SPECIFIED IN SAP"
      AnalysisPurpose="PRIMARY OUTCOME MEASURE">
      <Description>
        <TranslatedText xml:lang="en">Disease XXX Healing Rate at week 12 (FAS)</TranslatedText>
      </Description>
      <arm:AnalysisDatasets>
        <arm:AnalysisDataset ItemGroupOID="IG.ADEFF">
          <def:WhereClauseRef WhereClauseOID="WC.ADEFF.AVAL.HEAL"/>
          <arm:AnalysisVariable ItemOID="IT.ADEFF.AVAL"/>
        </arm:AnalysisDataset>
      </arm:AnalysisDatasets>
    ....
```

Creation of Analysis Results Metadata with SAS

- Output on a browser with a stylesheet

Analysis Results Metadata (Summary) for Study DRUG-XXX/STUDY-YYY	
Table 1.1 Disease XXX Healing Rate (FAS) Disease XXX Healing Rate at week 12 (FAS) Disease XXX Healing Rate at week 4, 8 (FAS)	Analysis Results Metadata (Detail) for Study DRUG-XXX/STUDY-YYY
Table 1.1	
Analysis Result	Disease XXX Healing Rate at week 12 (FAS)
Analysis Parameter(s)	PARAMCD = "HEAL" (Healed/Unhealed of Disease XXX)
Analysis Variable(s)	AVAL (Analysis Value)
Analysis Reason	SPECIFIED IN SAP
Analysis Purpose	PRIMARY OUTCOME MEASURE
Data References (incl. Selection Criteria)	ADEFF [PARAMCD = "HEAL" and FASFL = "Y" and ANL01FL = "Y" and AVISIT = "Week 12"]
Documentation	Calculate the Disease XXX Healing Rate, proportion differance and their 95% CI between treatment groups and perform the Wald test for the comparison of treatment at Week 12. FREQ procedure was used for this analysis. SAP
Programming Statements	[SAS version 9.2 or later] <pre>proc format ; value _TRTPF 1 = "Treat A" 2 = "Treat B" ; run ; proc freq data=ADEFF ; where FASFL = "Y" and AVISIT = "Week 12" and ANL01FL = "Y" ; table TRTPN*AVALC / riskdiff(equal var=sample) nocol nopercents ; format TRTPN _TRTPF. ;</pre>

Summary

- ARM in the e-Data submission package
 - PMDA recommends to submit the ARM as a part of ADaM Define.xml
 - Reviewers can understand how the analysis results are created
 - Need standardization to some extent
 - Several patterns (e.g. multiple datasets, variables, selection criteria, programming code)
 - Validation issue
 - OpenCDISC (V2.0) cannot validate ARM
 - Need re-creation/validation of programs/results for Legacy data conversion
 - Targeted analyses to be included in the ARM
 - Need the discussion/agreement with PMDA
- SAS is one of good tools for ARM generation
 - Need the standardization of input parameters (e.g. Excel metadata)
 - Need the basic knowledge of XML and Define.xml structure, but the programming code is not so complicated



References

- PMDA HP (e-Data submission)
<https://www.pmda.go.jp/review-services/drug-reviews/about-reviews/p-drugs/0003.html>
- CDISC Define-XML Specification Version 2.0
<http://www.cdisc.org/define-xml>
- CDISC ADaM (ADaM, ADaM IG and ARM are available)
<http://www.cdisc.org/adam>
- tsClinical Define.xml Generator (Fujitsu)
<http://www.fujitsu.com/jp/solutions/industry/life-sciences/cdisctool/>
- OpenCDISC
<http://www.opencdisc.org/>
- Yohei Takanami (2013). Simple Tool for Creating ADaM Define.xml for Statisticians in Pharmaceutical Companies Using SAS and HTML Application with Excel Metadata File. CDISC 2013 Japan Interchange
- Yohei Takanami, Nobuo Funao (2012, Kohgakusha). Statistical Analysis Software - SAS.
- Chris Holland, Jack Shostak (2012). Implementing CDISC Using SAS: An End-to-End Guide. Sas Inst

Thank you for your attention

