FROM CRF DATA TO DEFINE.XML: GOING “END TO END” WITH METADATA
Frank DiIorio, CodeCrafters, Inc., Chapel Hill NC
Jeff Abolafia, Rho, Inc., Durham NC

INTRODUCTION
The rapid, reliable, and cost-effective production of FDA deliverables is the Holy Grail of pharmaceutical companies and CROs. Difficult to achieve in a “calm” environment, they become even more problematic as statisticians, programmers, and project managers edge closer to the maelstrom that is the submission date. The potential for miscommunication within the project team increases, seemingly exponentially, thus making the need for robust programming tools and practices critical.

An earlier paper by the authors (see “References,” below) stated that well-constructed metadata and metadata access tools can significantly improve the creation of the datasets and documents that comprise an electronic submission. A year later, we even more emphatically believe this to be the case: metadata-driven applications and utilities can be integrated into standard business processes, speeding the production and improving the quality of deliverables.

The previous paper was heavy on theory and light on actual examples. We emphasized the benefits of metadata-driven processes over their earlier, mostly manual, counterparts. This paper switches the balance and describes the metadata tables and applications in detail. We demonstrate the ubiquity of the metadata by following its uses in the life cycle of one part of a submission package, namely, creation of CDISC SDTM-compliant Domain datasets and their accompanying documentation, define.XML.

The presentation is organized into four sections:
• A summary of the key features of metadata-driven applications;
• A description of key metadata tables and content currently used at Rho, Inc.;
• A detailed discussion describing the SDTM process flow and identifies how metadata and metadata access tools are utilized at each point; and
• A summary of key points and an outline of ideas we’re entertaining for further improving the scope and access to the metadata.

KEY FEATURES OF A METADATA-DRIVEN SYSTEM
Metadata is, classically, “data about data.” That is, it is data stored in a programmatically accessible manner, which contains characteristics of data stores. We expand the definition here so that it describes not only data but aspects of the work environment that uses the data. That is, we create metadata that describes the work flow that surrounds the data. The broader the definition, the wider the range of applications that can be automated.

Here are the key aspects of metadata usage, as described in the earlier paper and supplemented somewhat by our experience since its publication.
• Live by the Metadata, Die by the Metadata. Throughout the life cycle of a project, it’s key that the people entering and reviewing the metadata are well-trained (in terms of how to use the tools, what values are valid, etc.).
• Chose Storage Carefully. After considerable experimentation with SAS and other formats, we chose Microsoft Access as the database format for the metadata. It is a solid, reliable product that handles multiple users without complaint and is easily accessible by the all-important SAS applications that are used throughout the project life cycle.
• Develop Tools. The metadata organization, while not overly complex, should be presented to its users in a straightforward, well-documented manner. If the programmers chose to use the tables directly, they can do so, but they should also have utilities at their disposal that will reliably deliver information from the tables.
• Expect (and Welcome) Change. The amount of metadata tables and columns must be able to grow as the range of its uses grows. This, in turn, usually means growth of the tool set that accesses the metadata: existing tools are enhanced, and entirely new ones are created.
• Consider the Interface. The metadata must be easily entered. The interface must be easy to navigate, should be populated programmatically when possible, and drop-down boxes should be provided when appropriate to preempt entry of inappropriate values.
• Develop Complementary, Non-Metadata Tools. The development time of metadata tools is greatly shortened if a library of generalized, validated utilities is available.
• Utilize Complementary Metadata. The SAS dictionary tables and views are an invaluable adjunct to the project-specific metadata discussed in this paper. Familiarity with these resources is essential for developing robust metadata applications (see “References,” below, for examples).
• **Create Uniformity Across Projects.** Initial success created by metadata tools encourages their adoption in subsequent projects. Standardization of library names and directory structures greatly reduces, and often completely eliminates, the need for project-specific customizing.

• **Provide Documentation and Training.** The best tools in the world are of no use if they are poorly documented and not publicized. Programmer and end-user documentation must be in place for both the metadata and the tools that present the metadata to users.

**CURRENT METADATA ARCHITECTURE**

Figure 1, below, presents a high-level view of a project’s work flow. Many tasks need to be coordinated: data must be manually or electronically captured from the CRF; the statistical analysis plan must be reviewed to determine the analysis datasets’ contents; and Domain (SDTM) and analysis datasets must be programmed, validated, and prepared for submission to the FDA.

Figure 2 introduces two critical features into the work flow: metadata and metadata access tools. Due to space considerations, it focuses only on SDTM, but several points are made here that are applicable to any process described in Figure 1.

First, there is a considerable amount of logically separated metadata. Collectively, project configuration, SDTM and Analysis dataset, and Display (TFL) description metadata almost completely describe every piece of a project’s deliverables. Also notice the inclusion of SAS Dictionary Tables. They are included here because they are used extensively by metadata access tools.

The second key feature of Figure 2 is the macros and applications that appear at the top of the figure. Without tools that handle the metadata in a reliable and user-friendly fashion, the metadata would be rightly seen as a burden rather than as an asset. These tools, mostly macros, perform such tasks as creating fragments of ATTRIB and KEEP statements, displaying metadata contents in a PDF, and producing deliverables such as SAS transport files and define.XML. They will be discussed at length in
“Utilization in the SDTM Life Cycle,” below.

With out “Expect Change” caveat firmly in mind, here are some of the metadata databases that currently drive the systems at Rho. This section will give you a reasonably detailed view of the organization and contents of each database and most of the tables held in them. It is not our intent to get into the unnecessarily low-level details such as data types, lengths, keys, and the like.

**CONFIGURATION (Generic)**

Figure 3 describes the generic configuration metadata. It is a template, a generic version of a project’s structure. It is, essentially, a description of the recommended directory structure as described in SOPs. For directories containing data stores that will be accessed by SAS, the Structure table also contains recommended LIBNAMEs and engine options. See “CONFIGURATION (Project),” below, for more details.

**Tools Using the Metadata.** %buildConfig copies the generic configuration metadata to a new study. Then, after optional modification of the tables, %projectSetup creates the directory structure and configures programs to run various utilities. These macros are demonstrated in "Utilization in the SDTM Life Cycle," below.

![Figure 3](#)

**Generic Configuration**

<table>
<thead>
<tr>
<th>Macros</th>
</tr>
</thead>
<tbody>
<tr>
<td>%ISO %ISOdur</td>
</tr>
<tr>
<td>%setup %listConfig %distinct %libViewer et al.</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>structure</th>
<th>One observation per directory</th>
</tr>
</thead>
<tbody>
<tr>
<td>var</td>
<td>description</td>
</tr>
<tr>
<td>location</td>
<td>Directory name. Complete or relative path. May include MDB file name</td>
</tr>
<tr>
<td>libname</td>
<td>LIBNAME to allocate for this location</td>
</tr>
<tr>
<td>CDISC</td>
<td>Check box: is this a CDISC location?</td>
</tr>
<tr>
<td>submission</td>
<td>Check box: is this location part of an NDA?</td>
</tr>
<tr>
<td>formatSeq</td>
<td>Location’s order in format search path</td>
</tr>
<tr>
<td>macroSeq</td>
<td>Location’s order in macro autocall path</td>
</tr>
<tr>
<td>readOnly</td>
<td>Check box: is this location part of an NDA?</td>
</tr>
<tr>
<td>engineOpts</td>
<td>Options to use when allocating this location (DBMAX_TEXT, etc.)</td>
</tr>
</tbody>
</table>

**CONFIGURATION (Project)**

This database, shown in Figure 4 (next page), describes the directory structure and other key features of a project. During project setup, the generic configuration file is copied to a project-specific directory. The file may be customized to accommodate specific project needs (additional autocall or format libraries, data directories, etc.). Whatever the level of customization, the only hard and fast requirement is that the Structure table correctly identifies the project’s root directory, since all other directories for the project are entered as relative paths.

**Tools Using the Metadata.** %setup is run at the beginning of every program in the system. Using the Structure table, it creates LIBNAMEs, macro variables, and autocall and format search paths. Its function and output are described in detail in "Utilization in the SDTM Life Cycle," below.
SDTM (Generic, “Gold Standard”)

Figure 5 (next page) describes the metadata containing the CDISC standards for SDTM domains. Collectively, these tables identify the variables needed in each domain, their data type, descriptive label, permitted values, and the like. They do not, however, contain a complete description of how to create the domains, nor do they have what an FDA reviewer requires to understand how the fields were derived. This is handled by the project-specific SDTM metadata, described below.

Tools Using the Metadata. The %projectSetup macro, described above, copies the Gold Standard SDTM MDB to the project.

%printGoldstd displays all or selected Domain information in a PDF.

SDTM (Project)

Just as the generic configuration database is customized for a project, so too is the SDTM “Gold Standard” database (Figure 6, next page). It contains the fields from the Gold Standard as well as fields needed for programming and other purposes. Note that some of the Gold Standard fields (Domain and variable names, label) are not editable for standard, predefined Domains (AE, CM, etc.) but are subject to change for user-defined Domains.

Many of the SDTM domains are “vertical” representations of data that is often collected in a more “horizontal” manner. For example, a traditional Vital Signs dataset may have separate variables for weight, supine systolic blood pressure, and sitting systolic blood pressure. The SDTM data is a more relational design. Rather than repeating test name, value, and so on, making the dataset “wide,” the SDTM domains use variables that identify the name, value, unit, and other characteristics of a particular measurement, thus making the Domain “tall” and “skinny” (more observations, fewer variables). This verticality introduces the need for value-level metadata, in this case that describes each possible value of VSTESTCD. Thus we create the VALUE table, shown in Figure 6.
Figure 5

SDTM “Gold Standard”

Macros

%ISO %ISOdur
%setup %printGoldstd
%libViewer et al.

domains One observation per Domain

var  description

domain The name of the SDTM domain
description A description of the SDTM domain
structure The data structure of the domain (e.g., “One record per subject”) 
keyFields The fields identifying a unique record for a domain

variables One observation per domain per variable

var  description

domain The name of the SDTM domain
CDname The variable name
label The variable label
type The variable type
core Is the variable required, expected, or permitted

Figure 6

Project SDTM

Macros

%ISO %ISOdur
%setup %quoteList %distinct %libViewer et al.

value One observation per Domain per Variable per variable value

var  description

domain The Domain name
CDname The variable name
value The value for a specified level of CDNAME (i.e. SYSBP)
label A label for “VALUE” (e.g., “Systolic blood pressure”)
orsresU The programming specification for to how to create orresU (original result unit) for a specified level of CDNAME
length The variable length
controlledTerms Project-specific controlled terms (may be different than those found in the Gold Standard)
definition The programming specification for to how to create a given variable
FDAdefinition The definition of “VALUE” that will be presented in the define file
CRFloc The CRF page name and number, if the variable came from a CRF.
submit A flag indicating if the variables to be included in the submission database

domains One observation per Domain

var  description

domain The Domain name
inputNotes Notes about the input datasets used to create a given domain
outputNotes Notes about the domain to be created
submit A flag indicating if a given domain is to be included in the submission database

SAS Dictionary Tables

tables columns formats options
Pre-SDTM domains were often non-relational. The SDTM/relational approach for Vital Signs data, for example, looks like:

<table>
<thead>
<tr>
<th>CDname</th>
<th>VStestcd</th>
<th>orres</th>
<th>orresu</th>
</tr>
</thead>
<tbody>
<tr>
<td>VSTESTCD</td>
<td>Systolic BP</td>
<td>90</td>
<td>mg/hg</td>
</tr>
<tr>
<td>VSTESTCD</td>
<td>Diastolic BP</td>
<td>110</td>
<td>mg/hg</td>
</tr>
</tbody>
</table>

A non-relational representation of this data combines everything into one observation:

<table>
<thead>
<tr>
<th>sysbp</th>
<th>sysbpunit</th>
<th>diasBP</th>
<th>diasBPunit</th>
</tr>
</thead>
<tbody>
<tr>
<td>90</td>
<td>mg/hg</td>
<td>110</td>
<td>mg/hg</td>
</tr>
</tbody>
</table>

This is significant because the SDTM approach introduces the need for value-level metadata. Simply displaying `VSTESTCD` in the define file is inadequate. Its values and their meaning require additional metadata and, in turn, additional programming so that `Define.XML` can not only describe the variables' attributes, but also the possible values of the variable that have been, in effect, transposed.

**Tools Using the Metadata.** %printCDspecs displays all related metadata for a Domain in a PDF.

%attrib uses the VARIABLES table to create KEEP list and ATTRIB statements for a Domain. Since the macro reads from the metadata, the programmer is assured that the generated code is accurate.

%checkDomain evaluates a created Domain for consistency with the variables described in the project-level tables and similar tables in the Gold Standard.

These tools are described in more detail in "Utilization in the SDTM Life Cycle," below.

**ANALYSIS**

Figure 7 describes the datasets and variables in the Analysis metadata. The contents of these data stores differs from their SDTM counterparts. They contain not only CRF variables found in the SDTM Domains, but also have derived variables (e.g., change from baseline). This difference, plus the absence of a "gold standard" to mandate data type and order, results in metadata that bears only some resemblance to the SDTM tables.

**Figure 7**

![Analysis Metadata](image)

**Tools.** %printSpecs creates a programming spec PDF. Macro options give the user control over the sort order of the variables, whether to display only recently edited entries, and so on.
%attrib reads the variables table and creates KEEP and ATTRIB statements for a dataset. This relieves the programmer of the need to enter this information manually. It also ensures that the metadata and the output datasets will be in sync (provided, of course, that the programmer reruns the program creating the dataset once the metadata has been changed).

DISPLAYS

The creation of data displays – tables, figures, and listings – is well-suited for improvement using well-designed metadata and tools. The metadata can exploit similarities among a group of related displays. For example, the same table design might be used in 12 distinct displays, varying only by subpopulations (safety population, gender, etc.). Rather than write essentially the same program 12 times with only minor variations, we can create metadata, use macros to extract the metadata for a given display, and write a single, generalized program. This is a loaded statement, for it implies the need not only for new metadata tables and macros, but also for a change in the way displays are programmed. There is a learning curve, to be sure, just as there are there also huge productivity benefits to be gained.

UTILIZATION IN THE SDTM LIFE CYCLE

The power and elegance of a metadata-driven system are best appreciated when seen in action. In this section, we’ll illustrate the use of metadata during the creation of SDTM Domains and their accompanying documentation, Define.XML. For each phase, we will:

• Describe the work being performed, comparing “before and after” metadata usage efficiencies and issues;
• Identify the metadata tables that are read, written, or updated; and
• Describe some of the metadata tools that facilitate use of the metadata and then present program and output examples.

Overview

Figure 8 presents a highly simplified view of the SDTM life cycle. There is little here that is surprising to any one who has been involved in the production of clinical or analysis datasets and their documentation.

1. The project is initialized. A directory structure is created that complies with internal SOPs, and databases that will house configuration and other metadata are seeded with corporate-wide default values.

2. An in-house or outsourced data management system collects data from case report forms (CRFs) and other sources such as laboratory measurements.

3. Statisticians and support groups develop specifications for statistical programmers. Despite the presence of many predefined Domains, there are still many questions to be addressed. The spec writers need to consider: mapping of CRF variables to Domains; variables to use for conversion to ISO 8601 date-time values; what to put in supplemental Domains versus entirely new Domains; and so on.

4. Programming and validation for a Domain begin once the specifications are complete. The process is iterative: the datasets created by the primary and validation programmers may not be identical. The lack of agreement could be due to programming errors; it could also be caused by vague or incorrect specifications. Eventually, exchanges between the statistician, primary programmer, and validation programmer result in a valid, SDTM-compliant dataset.

5. The dataset is converted into a format acceptable to the FDA: either a SAS Transport file or an XML file.

6. The last stop in this simplified tour is the creation of the Domain dataset documentation. We create the “define” file as a PDF or, more typically, an XML file.

With this somewhat idealized view of the process in mind, let’s take a detailed look at the process.

Project Initialization

One of the great advantages of a table-driven, metadata-oriented environment is that it promotes standardization. Perhaps nowhere in the system is this better seen than during project initialization, depicted in Figure 9 (next page). This is a two-step
The user invokes a SAS macro (**buildConfig**) to copy a generic configuration file to the project root directory. The user passes the project root directory location to the macro and, if creating a multiple-study project, specifies a list of study names and descriptions. A simple invocation looks like:

```
%buildConfig(root=d:\comm\bigPharma\cholBgone)
```

The macro copies the generic configuration MDB to the project directory, using the root directory to change path values in the project database. The macro also creates a PDF that shows what is essentially the directory structure, LIBNAME assignment, and macro and format path sequence that will be used each time the metadata tools will be used. Once the output is reviewed and the user edits the MDB, the directory structure can be populated by the **projectSetup** macro.

```
%projectSetup(root=d:\comm\bigPharma\cholBgone)
```

**projectSetup** reads the configuration MDB created by **buildConfig** and:

- Creates directories underneath the project root directory;
- Copies generic SDTM and other metadata to the newly-created directories;
- Copies generic **RUN_xxx.SAS** programs to project directories. These programs run some of the metadata tools, and are populated with typical default macro parameters suitable for the study. The following is an example of the generated program **Run_printSpecs.sas**. Note that the **setup** call is prefilled with the correct drive and project parameters.

```
/* The program header has a copy of the source macro’s header. It contains a description
 of input and output datasets, macro parameters, examples of use, and other usage
 notes. */
%inc 's:\submissions\macros\util\setup.sas' / nosource2 ;
%setup(drive=s,drive=S, project=RHO\Sponsor\Project\PR01)
%printSpecs(order=alpha, use=_all_, skip=)
```

- Writes a PDF showing the directory tree that was created. Other diagnostics in the SAS Log identify the metadata tables (SDTM and others) that were seeded and the sample programs that were copied to the project directory.

The advantages of this approach compared to a manual one are significant.
• LIBNAMEs and directory locations for common data stores (raw data, derived data, macro libraries, etc.) are stored as data that will be accessed by a setup macro. Thus there is no need for a manually-created initialization program and, assuming no LIBNAMEs were changed, the new project uses the standard, corporate-wide names found in the SOPs.

• The sample programs don’t require any editing except for parameter changing. The documentation header for each macro called is included with each RUN_program.

Once the directory structure is created and the metadata databases are seeded, the domain creation process can begin. First, though, let’s take a look at the macro that is called at the beginning of every program in the system. %setup, as the name suggests, reads the configuration MDB and assigns LIBNAMEs, creates option and format search paths, and performs other activities based primarily on the STRUCTURE table. The call is simple:

```sas
%setup(project=root_directory)
```

Diagnostic output is written to the SAS Log and to setup.txt. Typical Log output follows:

```
SETUP-> Begin ----------------------------------------------------------------------
SETUP-> SETUP Ver 3. For help, rerun with HELP=y, or see s:\submissions\docs\setupMacro.doc
SETUP-> Project [RHO\Sponsor\Project\PR01] config [metadata\config\config.mdb]
SETUP-> Project drive [S]  Prefix []  Generic drive [S]
SETUP-> Configuration database [S:\RHO\Sponsor\Project\PR01\metadata\config\config.mdb]
SETUP-> LIBNAME AMETAANL had this engine option value: dbmax_text=10000
SETUP-> LIBNAME AMETACD had this engine option value: dbmax_text=10000
SETUP-> LIBNAME ACONFIG had this engine option value: dbmax_text=500
SETUP-> ODS path: template.templates sasuser.templat(update) sashelp.tmplmst(read)
SETUP-> See S:\RHO\Sponsor\Project\PR01\Prog\Derive\setup.txt for diagnostics
SETUP-> Processing required 1.3 seconds.
SETUP-> End ----------------------------------------------------------------------
```

The setup.txt file noted in the Log is a thorough description of all the LIBNAMEs, macro variables, and other settings that were created by the macro. Here, and throughout the system, the benefits of having clear, thorough diagnostics are two-fold.

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**Figure 10**

Create Programming Spec

```sas
%build %project %printCD %check %attrib %quoteList %distinct %libVieweret
```

First, they provides information to the user in an easily readable format. Input and run-time errors are caught and reported by the macros so that the program fails gracefully (that is, we don’t let SAS generate pages of run-time errors). This leads to the
next benefit of the system design: ease of use and good diagnostics encourage both continued usage and suggestions for enhancements and new tools. This is a back door approach, if you will, to acceptance of the standardized library names and directory structures encouraged by the SOPs.

Programming Specs

For years, the specifications for creating datasets were held in word processing formats such as Microsoft Word. Although this was a familiar and easy-to-use tool, it required considerable transcription by programmers. Dataset and variables specifics such as lengths, labels, and formats had to be copied into SAS programs. The process was tedious and error-prone. Moving the specifications out of a document and into a database (Figure 10, previous page) created significant opportunities. Now not only could a spec document be produced (duplicating and improving on the look and feel of the Word documents), but the document, since it is a table conceptually identical to a SAS dataset, could be used to generate SAS FORMAT, LENGTH, and other statements. This section describes generation of the spec document. We’ll see other programming tools later.

Recall that one of our points in the “Lessons Learned” section was that metadata are complex and so require a user-friendly interface. Figure 11, below, shows one of the data entry prototypes. The screens are simple and intuitive. This makes data entry easier and results in higher initial metadata quality (and, not to be discounted, happier statisticians).

Figure 11 CDISC Metadata Entry

A statistician or programmer can use the SDTM metadata to produce a spec document at any point during the metadata entry process. The %printCDspecs macro processes the metadata and produces a PDF that not only displays the metadata but also performs some metadata quality checks (note that some checks were already performed by the GUI and that other, more thorough checking is done later, by the %domainCheck macro). The macro displays dataset-level metadata for a Domain and, if found, its supplemental qualifier (“suppqual”) dataset, followed by variable-level and value-level data for the Domain.

Suppose a programmer wants to review the spec for Domain AE. The complete program follows:

```
%include 'utility_directory\setup.sas' / nosource2;
%setup(project=project_root_directory)
```
Messages to the Log describe the macro’s activity, noting among other items the complete path of the output PDF. Other parameters to the macro control the order of the variables (alphabetical within Domain, position within Domain), Domains not to process, and so on. Figure 12, below, shows partial screen capture for a Domain. The output displays not only the metadata information but also other features that are sometimes helpful. Among these: location of the Gold Standard metadata; key variables from the Domains table; location of the program generating the display; and highlighting of Required or Expected variables.

Figure 12 %printSpecs Output

<table>
<thead>
<tr>
<th>Variable Specs for Domain AE</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gold Standard: SDTM [321]</td>
</tr>
<tr>
<td>Created: 2008-01-12 [321]</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>Key variables from Domains table: STUDIED, AE_DOC, AESEQ</td>
</tr>
<tr>
<td>Variables are arranged by their position in the output Domain. Only Required, Expected and SUBMIT&lt;-&gt;Tes Variables are Displayed</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>STUDIED</td>
<td>Study Identifier</td>
<td>Char</td>
<td>12</td>
<td>Req</td>
<td>Yes</td>
<td>CRF</td>
<td>DEMOGRAPHICS</td>
<td>1</td>
<td>[ProDef] =E7/17/4/001/11/30</td>
</tr>
<tr>
<td>DOMAIN</td>
<td>Domain Abbreviation</td>
<td>Char</td>
<td>2</td>
<td>Req</td>
<td>Yes</td>
<td>DER</td>
<td></td>
<td></td>
<td>[ProDef] =AE</td>
</tr>
<tr>
<td>SUBJID</td>
<td>Unique Subject Identifier</td>
<td>Char</td>
<td>24</td>
<td>Req</td>
<td>Yes</td>
<td>DER</td>
<td></td>
<td></td>
<td>[ProDef] Concatenate RAW.DOCS.PROTOCOL and three-digit RAW.DOCS ID separated by '-'</td>
</tr>
<tr>
<td>ASEQ</td>
<td>Sequence Number</td>
<td>Num</td>
<td>4</td>
<td>Req</td>
<td>Yes</td>
<td>MC</td>
<td></td>
<td></td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE TERMA</td>
<td>Medically Terminated</td>
<td>Char</td>
<td>70</td>
<td>Req</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE EDCOD</td>
<td>Dictionary Defined Term</td>
<td>Char</td>
<td>250</td>
<td>Req</td>
<td>Yes</td>
<td>DER</td>
<td></td>
<td></td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE FREEQ</td>
<td>Body System of Organ Class</td>
<td>Char</td>
<td>250</td>
<td>Req</td>
<td>Yes</td>
<td>DER</td>
<td></td>
<td></td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE PRTY</td>
<td>Severity Probability</td>
<td>Char</td>
<td>8</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =p amat(RAW.AEDOCIE SEVEREVENT, AEDOCIEVENT)</td>
</tr>
<tr>
<td>AE TIMM</td>
<td>Duration</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =p amat(RAW.AEDOCIE ACTION, AEDOCIEATTERM)</td>
</tr>
<tr>
<td>AE ACONC</td>
<td>Other Action Taken</td>
<td>Char</td>
<td>100</td>
<td>Perm</td>
<td>Yes</td>
<td>ORF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE EQTY</td>
<td>Causality</td>
<td>Char</td>
<td>18</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =p amat(RAW.AESEQ, AEDOCIEVENT)</td>
</tr>
<tr>
<td>AE OUTF</td>
<td>Outcome of Adverse Event</td>
<td>Char</td>
<td>25</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =p amat(RAW.AESEQ.AESEQOUTCOMER, AEDOCIEEVENT)</td>
</tr>
<tr>
<td>AE ACONC</td>
<td>Congenital Anomaly or Birth Defect</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>ORF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE SADD</td>
<td>Restless or Sleep Disorder</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>ORF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE EMTS</td>
<td>Malignant Disease</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE ACONC</td>
<td>Allergic Reaction</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>CRF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
<tr>
<td>AE ACONC</td>
<td>Other</td>
<td>Char</td>
<td>1</td>
<td>Perm</td>
<td>Yes</td>
<td>ORF</td>
<td>ADVERSE EXPERIENCES</td>
<td>0</td>
<td>[ProDef] =RAW.AESEQ</td>
</tr>
</tbody>
</table>

(1) Red background indicates a Gold Standard that was changed during the study phase. The value displayed is the Study value.
(2) Blue background indicates a Key variable (in the Domains variable KEY).
Table consists of [1] Required, Expected variables and/or [2] CNAMED (used for project metadata)
Shaded row indicates Required or Expected variables (CNAMED="Key" or "Rep")

Programming Tools

The creation of the Domains, as we noted earlier, requires interplay among the primary programmer, validation programmer, and the statistician who wrote the specification. The Domain dataset is FDA-ready only when all three agree that the specification is correct and that the two programmers independently created a dataset that represents the spec’s intentions. Figure 13 (next page) presents an overview of the process.

Conforming to ISO 8601: %ISO and %ISOdur

Date and time-related values and durations are represented in SDTM using the ISO 8601 standard. The ISO specification is thoroughly documented, a tad arcane, and for most programmers, unfamiliar. We created several macros that offload the programming burden. (The spec writer does not necessarily have to know anything about ISO standards other than entering “ISO 8601” in the definition, along with the variables containing the date or date-time components.)

%ISO The programmer passes the name of the variable to be created, whether it is building a date, datet ime, or time variable, and the component/input variables. The output variable is an ISO 8601-compliant date-time in extended format (as required by the SDTM guidance). The macro call depends on the type of input variable being passed. In the simplest case, where a SAS date-time variable is being converted, the call looks like:
data cdisc.AE;
set AEtemp;
%iso(dateTime=AEstdtm, outVar=AEstdtc)

The macro looks for parameter conflicts, writes messages to the SAS Log as needed, and creates character variable AEstdtc as an ISO 8601-compliant date.

%ISOdur Conforming to the ISO 8601 representation of event duration is, to put it mildly, a challenge. Values can be represented in a number of forms: duration, duration/end date-time, start date-time/duration, or start date-time/end date-time.

Here, as with %ISO, we developed a tool that allows the Domain dataset programmer to pass a minimal set of parameters, knowing that errors and inconsistencies will be identified and that the duration value that is created will be ISO 8601-compliant. A call to the macro looks like:

%ISOdur(unit=dt, start=AEstdtc, end=AEendtc, out=AEdur, outType=d)

This call to the macro creates AEdur, which holds the duration between date-time variables AEstdtc and AEendtc. The value stored in AEdur contains duration in the form PyyYmmMddD. For example, if the duration between the two events was 1 year, 5 months, and 2 days, AEdur would equal P1Y5M2D.

Using Metadata to Build Code: %ATTRIB

One of the most onerous tasks of the pre-metadata programming days was manually moving the dataset specification’s variable names, formats, and other attributes into the text of a SAS program. Once the metadata were moved into a format that was programmatically accessible, issues of accuracy and timeliness disappeared, and programmers were happier. The %ATTRIB macro reads variable-level metadata for a Domain and builds portions of KEEP and ATTRIB statements. Since the macro reads directly from the metadata, there is no possibility of specs and programs being out of sync – if a label changes in the metadata for table X, the next time BUILD_X.SAS runs, it will pick up the change. The key point here is that the
programmer does not have to change the program. Each time the program is rerun, %ATTRIB picks up the variable attributes (variable name, label, length, type, and format) currently specified in the metadata.

The macro is simple to use. It creates macro variables _ATTRIB and _KEEPLIST. In the simplest case, all the programmer needs to know to use this tool are the names of the input metadata and Domain.

```sas
%attrib(in=meta.variables, domain=DM)
data CDISC.DM;
%unquote(& ATTRIB)
other SAS statements
keep & _keeplist;
run;
```

**Making Sure We Have What We Need: %checkDomain**

Two programmers producing the same results only validates that they understood the specification the same way. The %checkDomain macro performs additional QA by comparing what was produced (the Domain dataset) to the SDTM standard for the Domain. It creates a report summarizing errors and possible inconsistencies. The macro says, in effect, “Let’s compare the attributes of the Domain datasets to those in the project SDTM and the Gold Standard metadata.” Just what constitutes an error is sometimes a matter of judgment, but many of the criteria are taken from the current guidance from CDISC. A complete list of checks currently being used is found in Appendix B.

**Creating Deliverables: %export and %define**

The final phase of the SDTM life cycle is the creation of data and documentation deliverables. This is depicted in Figure 14, below.
Conversion of the Domain datasets from the proprietary SAS dataset format (Windows files with a SAS7BDAT extension) to a public domain format acceptable by the FDA (transport or XML) is straightforward. The %export macro utilizes the SAS dictionary tables to create a list of datasets in the CDISC domain directory, then reads the configuration metadata to identify the target directory. The macro call is as simple as its functionality. A complete exporting program follows:

```sas
%include 'utility_directory\setup.sas' / nosource2;
%setup(project=project_root_directory)
%export(type=xpt)
```

Things get more complicated when creating define.XML. This happens for several reasons. First, recall the discussion of value-level metadata ("SDTM (Project)," above). Simply displaying the name of a transposed variable in the define file is inadequate. Its values and their meaning require additional metadata and, in turn, additional programming. Define.XML must not only describe the variables' attributes, but also the possible values of the variable that have been transposed.

The second complexity introduced by Define.XML is the additional hyperlinking. Continuing the VSTESTCD example, there must be a link between the variable-level VSTESTCD and the display of its value-level metadata. The sample CDISC define.xml and define.xslt files demonstrate this and other internal hyperlinking. These are only the starting point for truly robust and usable files – modifications are needed to the XSLT and, to a lesser degree, the XML file so that the screen and printed versions of the files will be as well as aesthetically pleasing.

The %define macro heavily utilizes the project CDISC and configuration metadata to create Define.xml. It also uses numerous general-purpose macros (discussed in “General Purpose Tools,” below) to unload user-written format libraries, create variable and dataset lists, and so on. The call, however, can be deceptively simple:

```
%defineXML
```

The macro adheres to an important design principle: as the number of “moving parts” of a macro increases, so should its default diagnostic output. The macro’s Log messages tell the story of its execution. If all goes well, the story is a routine list of tasks attempted and successfully completed. If there is a problem, the macro should fail gracefully, explaining the problem and shutting down without letting SAS create needless and, frequently, puzzling error messages. The %defineXML Log excerpt in Appendix A illustrates this principle. The “Log as execution narrative” approach requires more coding, but is worth the effort. Consider, too, that if a general-purpose macro utility library is available, the coding effort can be significantly reduced.

General-Purpose Tools

If metadata without tools to access them is difficult, then access tools built without the benefit of a general-purpose utility library is almost impossible. We have developed a collection of well-documented macros that perform tasks ranging from the small-scoped and tightly defined to complete, standalone applications. The use of this library accrues several benefits:

- **Reliability.** The utilities only provide well-defined outputs (reports, macro variables, *et al.*). Additionally, the processing performed by the macro is likely more thorough than an application program’s “quick and dirty” coding.
- **Code Reduction.** A one-line call to a macro, possibly followed by another line that tests a return code or similar status check, is much more compact than writing code in-line. The fewer “moving parts” in a program, the better.

In the remainder of this section we discuss how some of these tools are used, either in conjunction with metadata tools or as individual applications.

Case 1: Use In a Metadata Application

This code fragment demonstrates how a metadata access application can use generalized tools from a macro library. Don’t focus here on how the macro is called and what it produces. Rather, consider how the application is not burdened with tedious SQL coding, quoting, and the like. That burden is offloaded to the macro, allowing the application programmer can focus on the program's core functionality.

The macro uses variable-level metadata to create a list of datasets marked for FDA submission. For each of dataset, it looks for identifier variables UNIQUEID, PATNO, and SCREENNO (not all variables are necessarily in each Domain). Then it lists any of the located observations from the Domain.

```sas
%macro listIDs;
%distinct(data=domMeta.variables, var=domain, where=%str(submit=1))
%do i = 1 %to &nLevels.;
%let token = %scan(&levels., &i.)
%varCheck(data=domain.&token., varList=uniqueid patno screenno)
%if &found. ^= %then %do;
proc report data=domain.&token. panels=5;
   columns &found.;
title “Identifier Variables for Domain &token.”
%end;
%end;
```

```
run;
%end;
%else %put No identifier variables were found for domain &token.;
%end;
%mend listIDs;

The code is compact and reliable, and will fail gracefully if identifier variables cannot be located. %distinct and %varCheck create lists and macro variables, held in &nLevels, &levels, and &found. The program is made both shorter and more reliable by their use. All that is required on the part of the programmer is to know that they are available and how to use them.

**Case 2: Standalone Application**

Some generalized, recurring requirements can become standalone applications that are useful diagnostic or display tools for a project. %libViewer is an example of this class of support tool (Figure 15, below). Unlike %distinct and %varCheck, shown in the previous example, %libViewer is a full-fledged, standalone application. It heavily utilizes the macro library to read the SAS Dictionary tables for an input library and create an HTML frameset. The application displays dataset and variable-level information about the library’s datasets. It also unloads any user-written formats currently allocated and builds links from the variable-level tables to a frame containing the contents of the individual formats. The HTML is useful when questions about variable types, formats, and the like arise. The program is relatively short (489 lines, 280 statements). Its design points out the power of a good utility library: it would be dramatically longer and likely less bulletproofed if it did not effectively use the utilities. The invocation is simple:

`%libViewer(lib=domain)`

**Ad Hoc Programming**

It would be foolish to believe that we could ever have a comprehensive, all-inclusive set of metadata and metadata tools that could address all data, display, and submission needs of all our projects. This is especially true in the CRO environment,
where corporate agility is key; often data may come from in-house systems or one or more EDC vendors. Add to this the diverse data, documentation and other standards of multiple clients, and you can easily see why flexibility (and "hand me the Maalox") becomes part of the programming mantra.

The unpredictable demands of this environment can be handled to some degree by knowing the metadata and generic utility resources that are available. We present two scenarios here, and then comment on the solutions.

**Case 1: Use Project Metadata**

**Scenario.** Analysis dataset metadata for a project checked out correctly – variable types, formats, lengths, and definitions were scanned by our generic metadata validation tools. The sharp-eyed statistician assigned to the project noticed a discrepancy in the wording of variable labels between two variables residing in different datasets but with different names.

Rather than manually review the metadata in Access, we used the analysis metadata as input to a program that read the metadata, built a string containing various attributes, then compared the string for like-named variables, reporting instances where there were multiple values (i.e., differing characteristics). The core section of the program is shown below:

```sas
proc sql noprint;
create table attribProblems as
  select varName, dsName, type, length, format, label,
  catx('#', type, put(length,5.), format, label) as compare
  from metaANL.variables(keep=dsName varName type length format label)
  group by varName
  having count(distinct compare) > 1
  order by varName, datasetName
;
quit;
```

Once dataset `attribProblems` dataset is created, it's a simple matter to use a reporting procedure such as `PRINT` or `REPORT` to display the results.

**Case 2: Use Generic Utilities**

**Scenario.** Visual review of early drafts of CRT datasets that were to be input to SDTM domains identified several variables that contained only missing values. The reviewer wanted a way to identify all missing variables that wasn’t as error-prone and tedious as manual inspection. It was a fairly simple matter to use several general-purpose macros to quickly produce the desired output. The program is shown in its entirety below (comments have been removed for the sake of brevity):

```sas
%inc 'directory\setup.sas' / nosource2 ;
%setup(project=d:\comm\bigPharma\cholBgone) /* Allocates LIBNAME DOMAIN */
%macro allMiss(libname=);
  %memlist(lib=&libname.)
  %let path = %sysfunc(pathname(&libname.));
  %put; %put LIBNAME=&libname. Path=&path.~Datetime: &dateTime.)
  %do i = 1 %to &_nmemList_.;
    %let temp = %scan(&_memList_., &i.);
    %nLevels(data=&libname..&temp, out=temp_&temp.)
    %dsetAttribs(data=&libname..&temp)
    %put; %put 
    %put # obs: &nobs.  # vars: &nVar. (&nchar. char, &nnum. num)  Updated: &modC.;
    %put &nMissing. variables have all missing values: ;
    %put &missing.;
  %end;
%mend;
%allMiss(libname=domain)
```

A partial listing of the Log output is shown below:

```
LIBNAME=sderive Path=S:\Client\Project\XX-01\Data\Derived
Run date-time: Friday, February 23, 2007 11:09 AM
Datasets [AE DEMOG PROTDEV RANDLIST]
[1 of 4] Dataset [sderive.AE]
  # obs: 203 # vars: 31 (20 character, 11 numeric) Last updated: 23FEB07:11:01
  No variables had all missing values
[2 of 4] Dataset [sderive.DEMOG]
  # obs: 82 # vars: 16 (9 character, 7 numeric) Last updated: 23FEB07:11:01
  No variables had all missing values
[3 of 4] Dataset [sderive.PROTDEV]
```
A review of how these ad hoc programs work, rather than what they produced, reveals the following important points:

- The programs will always supplement, not replace, manual processes. Some parts of the project life cycle are best done manually. The role of the ad hoc programs shown here is to help team members exercise their good judgment without forcing them to stay up through the wee hours of the night.

- The programs are compact and were coded quickly because we were aware of the resources at our disposal, namely, the metadata databases and utility library macros. The attribute-checking program required knowledge of the metadata tables. The missing value program required knowledge of the workings of several general-purpose macros (%nLevels, %memList, and %dsetAttribs).

- They started as ad hoc, but these programs can be integrated into larger, more formal and validated programs. The attribute inconsistency program could, for example, become a standard Analysis metadata quality assurance check. Its output would be enhanced in a production-quality tool (for example, we could highlight or display only the conflicting attributes). A formal, production-quality version of the program would also implement parameter and run-time error trapping (test for valid formation of the passed LIBNAME, see if the library had been allocated, and so on).

CONCLUSION

Pharmaceutical projects are typically complex, involving organizing hundreds of files in wide variety of formats. Furthermore, as the submissions date nears, there is an increased need for quick production of deliverables without sacrificing quality. Moving to a metadata-driven system at Rho has lead to more efficient processes at each phase in the project life cycle, and has improved the overall quality of deliverables. By utilizing a metadata based system:

1) Studies are now set up and configured by a standard program. This ensures a uniform setup across projects and allows studies to be setup instantaneously by a wide variety of personnel. Also by storing setup information in a database, it can be utilized by all subsequent programs.

2) Specifications for datasets as well as displays are now held in a database instead of in a document. This has lead to considerable savings in the time to produce datasets and displays and has led to higher quality datasets and displays. Furthermore, the data comprising these specifications can be repurposed downstream when producing transport and define files.

3) Once the dataset and display specifications have been entered, standard push button programs can be created to produce submission deliverables such as the SAS transport files and the Define file.

4) Quality control can be automated and made more robust. Programs can be developed to check specifications and to check the consistency between the metadata and the data. Also, by using metadata as input to programs, programmer error can be decreased significantly.

5) Metadata created at the early stages of a project can be utilized throughout the life cycle of a project, thereby increasing efficiency and decreasing the amount of time and resources needed for project deliverables. For example, metadata created at study setup time can be used at all subsequent stages of a project.

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Russ Helms and Ron Helms of Rho, Inc. provided the impetus for metadata use at Rho. They have also created an environment at Rho that encourages experimentation, rewards success, and is sympathetic when ideas don’t pan out.

April Sansom provided her typically thorough copyediting input, and is probably wondering why, with all those years of Catholic school education, the authors punctuate in a way that can only be described as whimsical.

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REFERENCES

Previous papers by the authors, focusing on metadata and tool development:

Appendix A: %Define Log Excerpt

See “Creating Deliverables,” page 14, for details. This Log excerpt is included as an example of the “Log as program narrative” approach to application diagnostics.

Begin Execution ---------------------------------------------------------------------------------------------------------------------------------
Error-trapping ---------------------------------------------------------------------------------------------------------------------------------
All initial checks passed.
Set options and variables for DOCTYPE=PDF -----------------------------------------------------------------------------------------------
Input options and file names -----------------------------------------------------------------------------------------------
Run time [Monday, February 26, 2007 10:02 AM] Type [ANL]
USE list [AEXP1 XRAY1] SKIP list []
SAS style template [style.ndarpt] Link color [RED]
Handling of user-written formats [Convert to LINKS]
From ACONFIG.STUDIES: name/description [XX-01/Miracle Drug]
Orientation [LAND] Doc type [PDF]
File locations:                                                                                                       ...
... Program [S:\RHO\Client\Project\XX-01\prog\define\Run_define.sas]
... Input data dir. [S:\RHO\Client\Project\XX-01\data\aSub]
... Metadata dir. [S:\RHO\Client\Project\XX-01\MetaData]
... Distiller watched dir. [I:\PDF]
... Define file [S:\RHO\Client\Project\XX-01\Submission\definetest.pdf]
... SETUP diagnostics [S:\RHO\Client\Project\XX-01\prog\define\setup.txt]
Create filtered subsets of META.VARIABLES. Also make ANCHORS list, counter. -------
Use this filter: (where=(dataSource='A' & (root in ('AEXP1' 'XRAY1'))))
2 datasets in DATASETS (has possible ANCHOR / ROOT duplicates due to dataset splitting)
2 datasets in DATASETSDUPS (unique ANCHOR / ROOT values)
2 distinct data sets: AEXP1 XRAY1
2 anchors from WORK.DATASETSFORDEFINENODUPS: A_AEXP1_X A_XRAY1_X
203 variables in VARIABLESFORDEFINE
Unload user formats --------------------------------------------------------------------------------------------------------------------------------
Report set up ---------------------------------------------------------------------------------------------------------------------------------
Define column widths and set page orientation, margin options
Delete S:\Submissions\macros\development\define\NDAtestdata\Analysis\defineFCDTEST.pdf
Build ODS statements for doctype=PDF, links=YES
ODS type resolves to [ps]
Write report 1: Datasets, TOC --------------------------------------------------------------------------------------------------------------------------------
Iteration [1 of 1] Subset for DOCTYPE=A (Statistical Analysis) Anchor: A_Datasets
Write report 2: Variable-level --------------------------------------------------------------------------------------------------------------------------------
[1 of 2] Anchor=A_AEXP1_X
# of vars [135]
Title 1 [Cerezyme: Variables for Table AEXP1]
Title 3 [Description: Adverse Events (Anly]
Title 4 [Structure: Multiple records per patient]
[2 of 2] Anchor=A_XRAY1_X

Also see http://www.CodeCraftersInc.com for other papers and resources not directly related to the current paper.

CDISC resources:
SDTM Version 3.1 http://www.cdisc.org/models/sds/v3.1/
# of vars [68]
Title 1 [Cerezyme: Variables for Table XRAY1]
Title 3 [Description: X-Ray Assessment and Findings of the Target Elbow (Anly)]
Title 4 [Structure: multiple records per patient.]

Write report for unloaded user formats ---------------------------------------------------
Create PDF with links - send to Distiller
Input ["S:\Submissions\macros\development\define\NDAtestdata\Analysis\defineFCDTEST.tmp"]
Output/Distill ["I:\Rho_Pub\PDF\in\defineFCDTEST.ps"]
Copy PDF to Submissions directory
Delete temp files from watched Distiller directory
Terminate  -------------------------------------------------------------------------------

APPENDIX B: DOMAIN CHECKING

See “Making Sure We Have What We Need: %checkDomain,” page 13, for more information. The format and presentation of the list is noteworthy. It is taken directly from the %checkDomain program header. This is, of course, a matter of preference and style (the list could have been located in a separate document, for example). We chose to place the documentation "close to the source."

[1] Metadata
   [a] Compare Gold Standard, Project
      [1] Variable labels don't match
      [2] Variable type does not match
      [3] Core value does not match
      [4] Project metadata is missing a Required or Expected variable.
      [5] Project metadata contains a variable that is not in a Gold Standard domain (this test ignores custom Domains).
   [b] Compare Project SUPPQUAL metadata, Domain data header:
      [1] RELVAR (CDISC field name IDVAR) is blank
      [2] RELVAR (CDISC field name IDVAR) is non-blank, but the variable is not found in the related Domain
      [3] RELVALUE (CDISC field name IDVARVAL) is blank
      [4] QNAM (CDISC field name VALUE) is blank
      [5] QNAM (CDISC field name VALUE) is non-blank, but the value does not conform to SAS naming conventions
      [6] LABEL (CDISC field name QLABEL) is blank
      [7] LABEL (CDISC field name QLABEL) is not-blank, but is longer than 40-characters
      [8] ORRES (CDISC field QVAL) is blank
   [c] Compare Project metadata, Domain data header:
      [1] Variable labels don't match
      [2] Variable types don't match
      [3] Key variable (in DOMAINS table, variable KEY_VARS) is missing in VARIABLES table

   [a] Compare Project SUPPQUAL metadata, Domain data header:
      [1] RELVAR (CDISC field name IDVAR) is blank
      [2] RELVAR (CDISC field name IDVAR) is non-blank, but the variable is not found in the related Domain
      [3] RELVALUE (CDISC field name IDVARVAL) is blank
      [4] QNAM (CDISC field name VALUE) is blank
      [5] QNAM (CDISC field name VALUE) is non-blank, but the value does not conform to SAS naming conventions
      [6] LABEL (CDISC field name QLABEL) is blank
      [7] LABEL (CDISC field name QLABEL) is not-blank, but is longer than 40-characters
      [8] ORRES (CDISC field QVAL) is blank
   [b] Compare Gold Standard metadata, Project metadata, Domain data header:
      [1] Variable labels don't match
      [2] Variable types don't match
      [3] Key variable (in DOMAINS table, variable KEY_VARS) is missing in VARIABLES table
   [c] Compare Project metadata, Domain data header:
      [1] Lengths are not equal
      [2] Formats don't match
      [3] Data types don't match
      [4] Variable labels don't match
      [5] xxTESTCD must conform to short SAS variable names
      [6] xxTEST must <= 40 characters
      [7] DM.ARMCD must conform to short SAS variable names

[3] Data
   [a] Non-SUPPQUAL Domains:
      [1] Missing value found for a required variable
      [2] Expected variable does not have at least one non-missing value
      [3] Improperly-formed ISO 8601 date or date-time value
      [4] DOMAIN variable corresponds to expected Domain (e.g., all values equal EX for Domain data set EX)
      [5] Variable is in the data, but not in either Gold Standard or Project metadata
[6] xxTESTCD must conform to short SAS variable name
[7] If <domain>SEQ is present, test for unique values within subject (USUBJID).
[8] USUBJID contains one or more blanks
[9] xxTEST must be <= 40 characters
[10] DM.ARMCD must conform to short SAS variable name

[b] SUPPQUAL Domains:
[1] RDOMAIN, if non-blank, is in list of Domain datasets
[3] QNAM does not conform to SAS short-variable naming
[4] QLABEL longer than 40 characters
[5] Observation is not uniquely identified by STUDYID, RDOMAIN, USUBJID, IDVAR, IDVARVAL, and QNAM
[6] Domain contains non-standard variable(s)
[7] Domain contains numeric variable(s)
[8] Domain and metadata QNAM values do not reconcile (Domain has values not in metadata and/or is missing values defined in the metadata)
[9] Metadata.IDVAR not found in Domain dataset
[10] Expected SUPPQUAL variable not in Domain dataset
[11] Expected/actual QNAM values don't reconcile