DH12: Metadata-Driven Tool for Creation of a Dataset for Exploratory Analyses (AXDM)

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Alexey Kuznetsov
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1. Definitions

- **Data visualization**
  Data visualization is the visual and interactive exploration and graphic representation of data of any size, type (structured and unstructured) or origin.[1]

- **Data mining**
  Data mining is the computing process of discovering patterns in large data sets involving methods at the intersection of machine learning, statistics, and database systems.[2]

- **Metadata-driven program**
  Metadata is data (information) that provides information about other data.[3] And metadata-driven program is the program that decides on how to process the data based on the provided metadata.

REFERENCES
[3]. https://en.wikipedia.org/wiki/Metadata
2. Dataset for exploratory analyses (AXDM)

ADaM datasets

<table>
<thead>
<tr>
<th>Name</th>
<th>Size</th>
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<tbody>
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**AXDM dataset contents**

<table>
<thead>
<tr>
<th>NAME</th>
<th>LABEL</th>
</tr>
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<tbody>
<tr>
<td>ADEG_1_HRMEAN_2_30</td>
<td>ADEG_HOLTER ECG EXTRACT_Mean Heart Rate (beats/min)_VISIT 2_30 minutes after the start of infusion</td>
</tr>
<tr>
<td>ADEG_1_HRMEAN_2_45</td>
<td>ADEG_HOLTER ECG EXTRACT_Mean Heart Rate (beats/min)_VISIT 2_45 minutes after the end of infusion</td>
</tr>
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<td>ADEG_1_HRMEAN_2_80</td>
<td>ADEG_HOLTER ECG EXTRACT_Mean Heart Rate (beats/min)_VISIT 2_60 minutes after the start of infusion</td>
</tr>
<tr>
<td>ADEG_1_HRMEAN_2_90</td>
<td>ADEG_HOLTER ECG EXTRACT_Mean Heart Rate (beats/min)_VISIT 2_90 minutes after the end of infusion</td>
</tr>
<tr>
<td>ADLB_CREAT_1</td>
<td>ADLB_Creatinine (umol/l) VISIT 1</td>
</tr>
<tr>
<td>ADLB_CREAT_2</td>
<td>ADLB_Creatinine (umol/l) VISIT 2</td>
</tr>
<tr>
<td>ADLB_CREAT_3</td>
<td>ADLB_Creatinine (umol/l) VISIT 3</td>
</tr>
<tr>
<td>ADPI_WEEKLY_1</td>
<td>ADPI_Weekly average current pain intensity_Week 1</td>
</tr>
<tr>
<td>ADPI_WEEKLY_2</td>
<td>ADPI_Weekly average current pain intensity_Week 2</td>
</tr>
<tr>
<td>ADPI_WEEKLY_3</td>
<td>ADPI_Weekly average current pain intensity_Week 3</td>
</tr>
<tr>
<td>AGE</td>
<td></td>
</tr>
<tr>
<td>SITEGR1</td>
<td></td>
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<tr>
<td>STRATA</td>
<td></td>
</tr>
<tr>
<td>TRT01P</td>
<td></td>
</tr>
<tr>
<td>USUBJJ</td>
<td></td>
</tr>
</tbody>
</table>

AXDM is a dataset containing 1 line per subject with many variables from various domains (e.g. demographic, disposition, medical history characteristics as well as efficacy, laboratory, vital signs, procedures data for all or selected parameters at each period/visit/timepoint) The advantage is that you have all the trial data combined in one dataset and you can easily do plotting and modeling of the data.
3. Types of variables in AXDM

- **Subject-level**
  variables coming from the “1-line-per-subject” datasets e.g. ADSL

- **Transposed from BDS[^1]**
  variables created by transposing BDS datasets variable(s) AVAL/CHG/PCHG with id
  variables [PARCATx] <PARAMCD> [AVISIT(N)] [ATPT(N)] by subject

[^1] BDS – Basic Data Structure
3. Types of variables in AXDM (2)

- **Occurrence flags**
  Flag variables indicating occurrence of an event for a subject in a period e.g. occurrence of a particular treatment-emergent adverse event (AEDECOD) for a subject or presence of a particular medical history term in a specific system organ class (MHBODSYS) for a subject

- **Occurrence counts**
  Occurrence counts: numeric variables containing number of events that occurred for a subject e.g. number of occurrences of a particular treatment-emergent adverse event (AEDECOD)

- The list can be extended if required for specific exploratory analyses.
4. Variable naming conventions and rationale (Subject-level)

- Subject-level

  `<Dataset name>_<Orig. variable name>`

  Example: `ADSL_RACE`

  Rationale:
  1) Clear traceability to the original dataset,
  2) Ability to easily refer to all the variables coming from a particular domain as an array (e.g. `ADSL_:` in SAS).

  Note: CDISC restriction to maximum 8 characters for a variable name length cannot be applied to AXDM.
4. Variable naming conventions and rationale (Transposed BDS)

- Transposed from BDS

<Dataset name>[_PARCATx][_PARCATy][_PARAMCD][_AVISIT(N)][_ATPT(N)][_ANLzzFL]

Example: ADEF_RESP_RED30P_5

Rationale:
1) Same as for Subject-level variables +
2) Easy code for creation of the dataset
(e.g.

```
proc transpose data = ADEF delimiter=_;
   by USUBJID;
   var AVAL;
   id PARAMCD AVISITN;
run;
```
4. Variable naming conventions and rationale (Occurrence)

- **Occurrence flags**
  
  \(<\text{Dataset name}\>_FL[\_\text{Standardized term variable name}]<\_\text{Standardized term code}>\)

  **Examples:**
  
  ADAE\_FL\_AEDECOD\_HEADACHE
  
  ADMH\_FL\_MHBODSYS\_1

  (Standardized term code here is a number since the text within the variable may be too long. The complete description of the term may be provided in the variable label)

- **Occurrence counts**
  
  \(<\text{Dataset name}\>_N[\_\text{Standardized term variable name}]<\_\text{Standardized term code}>\)

  **Example:**
  
  ADAE\_N\_AEDECOD\_HEADACHE

  The same concept as for the flags, but the variable is numeric and represents the counts of occurrences of the AE.
Example of plotting distributions of baseline laboratory values from AXDM
5. Data visualization using AXDM and SAS JMP®

- Example of plotting primary endpoint on y-axis versus baseline lab value on x-axis by medical history subgroup
Example:
The following code applies a model with effect selection in the framework of general linear models which is supposed to identify if age, sex, race, medical history event involving any particular body system or baseline creatinine level can explain change from baseline to visit 8 in quality of life score:

```sas
proc glmselect data=AXDM(where = (RANDFL = "Y" and TRT01P ne "Placebo"));
   class ADMH_FL_MHBODSYS_;;
   model ADEF_EQ5DTOT_V8_CHG = AGE SEX RACE ADLB_CREAT_BSL ADMH_FL_MHBODSYS_: / details=all;
run;
```
7. Creation of AXDM

Metadata-driven approach for creation of AXDM consists of 3 steps:

- creation of metadata for a trial
- automatic generation of a SAS program based on the metadata
- manual fine-tuning the SAS program if required and running it
7. Creation of AXDM (Metadata table)

An example of a metadata table for AXDM:

<table>
<thead>
<tr>
<th>MEMNAME</th>
<th>BY</th>
<th>ID</th>
<th>IDLABEL</th>
<th>OCCUR</th>
<th>WHERE</th>
<th>VAR</th>
</tr>
</thead>
<tbody>
<tr>
<td>ADSL</td>
<td>USUBJID</td>
<td></td>
<td></td>
<td></td>
<td>RANDFL = 'Y'</td>
<td></td>
</tr>
<tr>
<td>ADQS</td>
<td>USUBJID</td>
<td>PARAMCD AVISITN</td>
<td>PARAM AVISIT</td>
<td></td>
<td>RANDFL = 'Y' &amp; PARAMCD = 'PAINNOW' &amp; ANL01FL='Y'</td>
<td>CHG</td>
</tr>
<tr>
<td>ADVS</td>
<td>USUBJID</td>
<td>PARAMCD AVISITN</td>
<td>PARAM AVISIT</td>
<td></td>
<td>RANDFL = 'Y' &amp; not missing (AVAL)</td>
<td>AVAL</td>
</tr>
<tr>
<td>ADMH</td>
<td>USUBJID</td>
<td></td>
<td></td>
<td></td>
<td>PARCAT2 = 'MEDICAL HISTORY' &amp; not missing (MHBODSYS)</td>
<td></td>
</tr>
<tr>
<td>ADLB</td>
<td>USUBJID</td>
<td>PARAMCD AVISITN</td>
<td>PARAM AVISIT</td>
<td></td>
<td>RANDFL = 'Y' &amp; not missing (AVAL) PARCAT2 = 'CENTRAL LABORATORY'</td>
<td>AVAL</td>
</tr>
</tbody>
</table>

Note:
Parts of the metadata table in black font can be generated automatically based on the contents the SAS library.
The part in red most likely is added by the user accounting for the needs of the analysis.
7. Creation of AXDM (Generated SAS code)

```sas
data ADSL;
  set temp.ADSL(where=(RANDFL='Y'));
run;
%get_dups(ds=ADSL, by= , id=%str());

data ADQS;
  set temp.ADQS(where=(RANDFL='Y' & PARAMCD='PAINNOW' & ANL01FL='Y'));
  attrib all_label=" ";
  length id $40  idlabel $200;
  id=catx("_",trim("ADQS"), "CHG", trim(PARAMCD), trim(AVISIT));
  idlabel=catx("_",trim("ADQS"), "CHG", trim(PARAM), trim(AVISIT));
run;
%get_dups(ds=ADQS, dsout=ADQS_CHG, by= , id=PARAMCD AVISIT, bdupsby=PARAMCD);

proc transpose data = ADQS CHG dups (where=(bdups=0)) out=t_ADQS_CHG(drop= _name_);
  by USUBJID;
  id id;
  idlabel idlabel;
  var CHG;
run;
...
```
8. Conclusion

Pros:
- AXDM is convenient to do easily customizable data visualization and high level data mining
- AXDM is more understandable for people with minimal or no CDISC experience
- Semi-automated metadata file generation from SAS library contents eases the creation of such a dataset

Cons:
- The data structure does not cover all the data that could be stored in ADaM/SDTM
- The metadata file has to be prepared by a statistical programmer or a biostatistician with good knowledge of the particular trial data
Questions?

Don’t hesitate to contact me after PhUSE 2017 at Alexey.Kuznetsov@grunenthal.com