

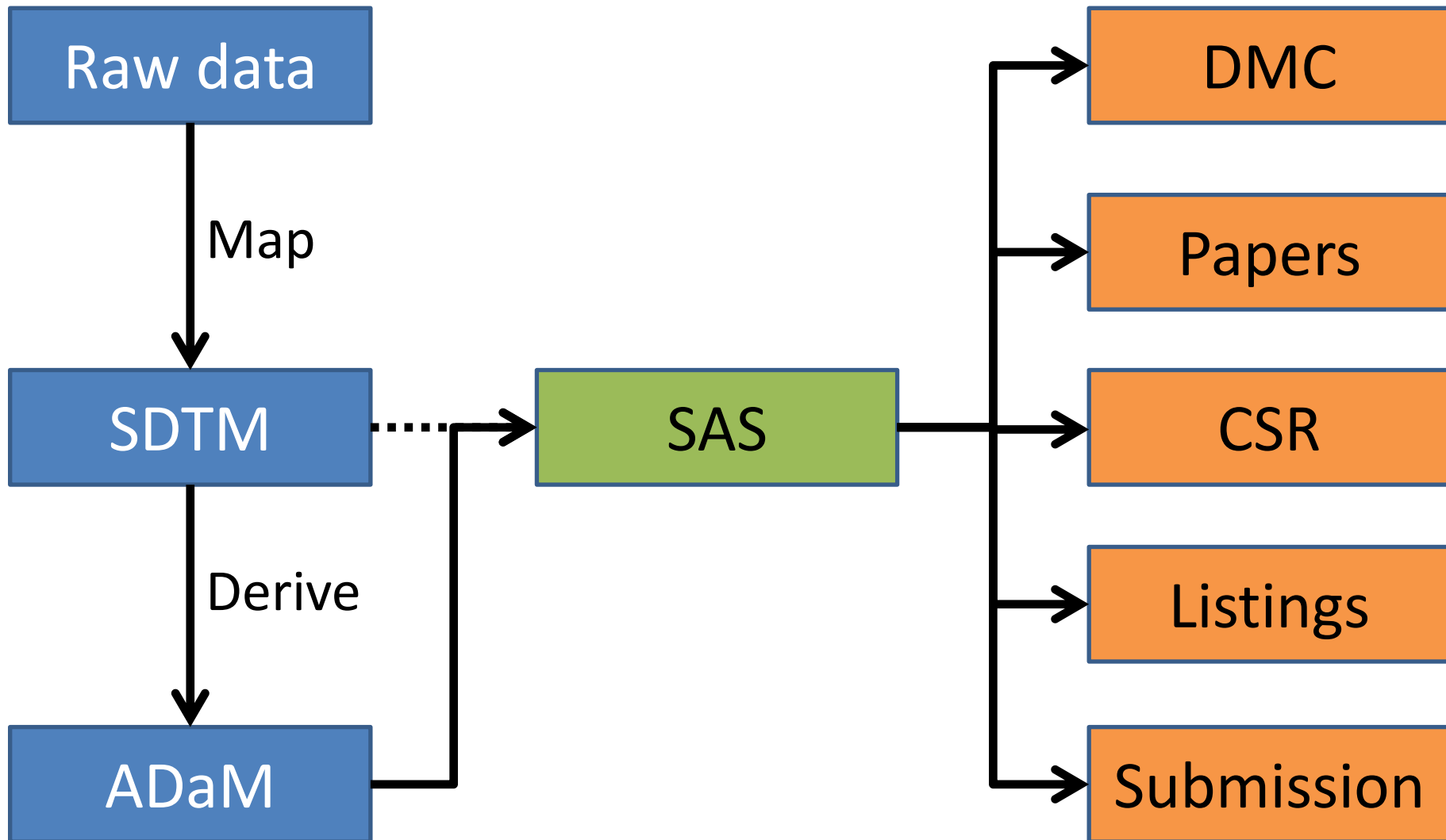
Implementing CDISC Standards in large academic clinical trials

Will Stevens
Clinical Trial Service Unit
Nuffield Department of Population Health
Oxford
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Context

- Several large clinical trials (>10000 participants)
- Some smaller trials
- Starting from about 2010 CDISC standards began to be adopted, beginning with SDTM
- This talk is about my work in implementing SDTM, ADaM and define.xml in three trials since 2013. Most of the work was done for the REVEAL trial (30,000 participants, 4.1 year median FU).

Workflow



Tools and languages

- Study database and analysis databases hosted using Ingres SQL
- SQL is also used for data transformations
- But native language features aren't expressive enough
- SQL wrapped in a lightweight XML-based macro system
- PHP used on top of this to preprocess XML

Example

```
<!-- Assign a visit number based on SVSTDY -->  
<extremum-value min-or-max="min">  
  <src-table>tv join sv</src-table>  
  
  <src-key>sv.usubjid</src-key>  
  <src-key>sv.svrefid</src-key>  
  
  <test-exp>abs(tv.visitdy - svstdy)</test-exp>  
  
  <column>tv.visitnum</column>  
</extremum-value>
```

Documentation

Tables

- [adadj](#)
- [adadjo](#)
- [adae](#)
- [adaend](#)
- [adaendo](#)
- [adaeo](#)
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Added 

Changed 

Removed 

ADMEDDRA

Modified: 2017-06-09

This table contains time to event for all MedDRA categories as follows: NONFATAL and SAE. For NONFATAL and AE. For each combination of fatal and non-fatal the time to event for all events (i.e. all events with

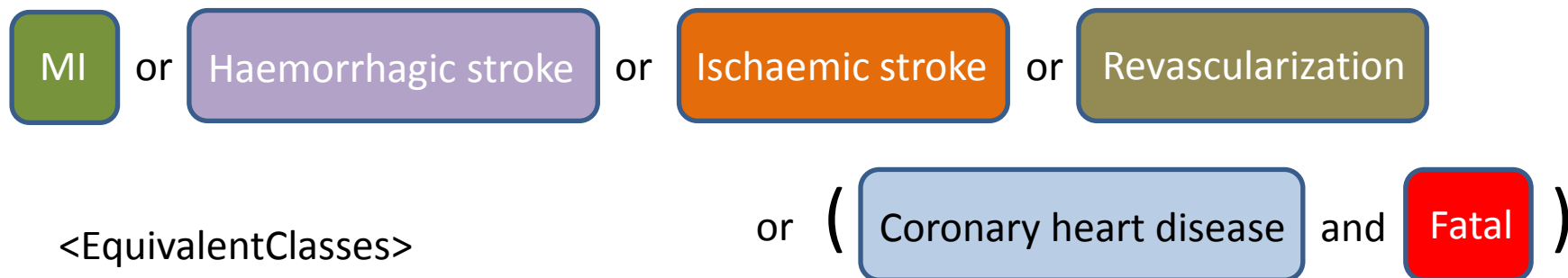
Column	Type	
usubjid	integer	Unique subject identifier for all studies
fasfl	char	Full analysis set population flag
siteid	varchar	Study site identifier
sitegr3	varchar	Site group for grouping 3 (coarse)
sitegr3n	integer	Site group ID for grouping 3 (coarse)
param	varchar	MedDRA term corresponding to the event
paramcd	integer	MedDRA SOC, HLGT or HLT code, corresponding to the event
parcat1	varchar	Level of the MedDRA code in PARCAT1
parcat1n	integer	Numeric code corresponding to PARCAT1
fatal	integer	Whether fatal or non-fatal event

Event classes

Type of Major Vascular Event	Niacin–Laropirant (N=12,838) <i>no. of participants with event (%)</i>	Placebo (N=12,835)
Major coronary event		
→ Nonfatal myocardial infarction	402 (3.1)	431 (3.4)
Death from coronary cause	302 (2.4)	291 (2.3)
→ Any major coronary event	668 (5.2)	694 (5.4)
Stroke		
Nonhemorrhagic stroke	389 (3.0)	415 (3.2)
Hemorrhagic stroke	114 (0.9)	89 (0.7)
Any stroke	498 (3.9)	499 (3.9)
Revascularization procedure		
Coronary revascularization	591 (4.6)	664 (5.2)
Noncoronary revascularization	236 (1.8)	258 (2.0)
Any revascularization procedure	807 (6.3)	897 (7.0)
→ Any major vascular event	1696 (13.2)	1758 (13.7)

Web Ontology Language (OWL)

Major vascular event =



```
<EquivalentClasses>
  <Class IRI="#MAJ_VASC_EV"/>
  <ObjectUnionOf>
    <Class IRI="#MI"/>
    <Class IRI="#STR_HAEM"/>
    <Class IRI="#STR_ISCH"/>
    <Class IRI="#REVASC"/>
  <ObjectIntersectionOf>
    <Class IRI="#CHD"/>
    <Class IRI="#Fatal"/>
  </ObjectIntersectionOf>
</ObjectUnionOf>
</EquivalentClasses>
```

```
<DisjointClasses>
  <Class IRI="#STR_HAEM"/>
  <Class IRI="#STR_ISCH"/>
</DisjointClasses>
```


Protégé : A free, open source user interface for OWL documents

The screenshot displays the Protégé OWL editor interface. The main window title is "ae (http://www.semanticweb.org/wills/ontologies/2015/8/ae) : [J:\ontologies\reveal\ae.owl]". The menu bar includes File, Edit, View, Reasoner, Tools, Refactor, Window, and Help. The toolbar shows navigation and search icons. The main workspace is divided into several panes:

- Class hierarchy:** Shows a tree view of classes. The class **INF_RESP** is selected and highlighted in blue. Other classes include AE_SOCIAL, AE_UROL, AE_VASC, AE_NEOP, AE_UNCLASSIFIED, CAN_ANY, ENTRY, EPS, INF_ANY, DTH_INFEC, INF_GI, INF_NERV, INF_OTH, INF_REPROD, INF_SKIN, INF_SYST, INF_URIN, RESP_ANY, DTH_ANY, Fatal, 'Full Hierarchy MedDRA', Hospitalised, 'Primary SOC Route MedDRA', 'Prioritised SOC Route MedDRA', and Uncoded.
- Annotations: INF_RESP:** Shows a list of annotations for the selected class. The annotations are:
 - comment:** Respiratory infection
 - filter [type: boolean]:** true
 - paramcd:** RIRS
- Description: INF_RESP:** Shows the logical description of the class:

Equivalent To **INF_ANY** and 'FH SOC 10038738 Respiratory, thoracic and mediastinal disorders' and (not ('Angina gangrenous (10002379)' or 'Paragonimiasis (10033794)' or 'Oropharyngeal candidiasis (10050346)' or 'Oro-pharyngeal aspergillosis (10053029)' or 'Oropharyngeal gonococcal infection (10066236)'))

The bottom status bar contains the text: "To use the reasoner click Reasoner Start reasoner Show Inferences".

Testing

- After producing SDTM domains, and after deriving ADaM datasets, tests are carried out on the data.
- Some tests apply across multiple domains/datasets. E.g. Does the domain contain at least some data? Are the --SEQ numbers correct?
- Some overlap with Pinnacle 21 tests.
- Some tests are written in response to errors found during development (to prevent reappearance of error, or recurrence of error elsewhere).

Testing - examples

- Phase/period/visit columns – check that they are consistent with dates obtained from another source.
- When ADaM datasets should only contain a subset of data from SDTM, check that nothing unexpected is present.
- Check that sums and totals of the same data represented in different places match.

Future developments

- Many of the submission components are also useful during the course of the trial: annotated CRF and define.xml should be made available early on in the trial.
- How does performance of SQL data transformations compare with SAS?
- Vector-based RDMS could potentially reduce storage requirements and speed up data transformation time.

Strengths and weakness of SDTM and ADaM

- Overall, CDISC standards seem to be beneficial for us. It makes it easier to transition between one trial and another than if each trial (perhaps developed and conducted by a different team of people) has its own way of representing data.
- There is a lack of specificity in SDTM and ADaM, and often scope for different interpretations.
- No (or little) entity-level modelling of data in the way that software developers are generally used to doing.

Strengths and weakness of SDTM and ADaM

- Standardised tabulation (SDTM), standardised analysis datasets (ADaM) and (with CDASH) standardised data at the point of collection. These are all at the edges of the trial: no standardised modelling of structured clinical trial data.
- Is this a strength or a weakness?

As standards develop over time, it is convenient if they are modular (well defined interfaces, but otherwise potentially independent of everything else), so that a change in one place doesn't require changes everywhere else.

Evolvability should be taken into account when putting together a system of standards. Design of the standards should consider: which parts of the standard can change without affecting other parts? What are the possible future changes?

