ABSTRACT
CDISC BRIDG (Biomedical Research Integrated Domain Group) is the canonical model in the biomedical / clinical research arena. On one end it serves as a communication model between domain users and the technical experts, but can be used further for enterprise interfacing for message-based communications and data exchange. Beside, BRIDG is on the way to get ISO approval and has reached an important step on this way in April 2015.

The challenge with BRIDG is, that is partial overwhelming and stays under the impression to be too complicated.

This article will give a simplified view on BRIDG to start up understanding and takes a simple example on how data from one system / standard can be / are mapped into BRIDG.
Finally I will highlight what’s new in Version 4 which is (at the time writing the abstract) out for comments.

INTRODUCTION
A LITTLE BIT OF HISTORY

2003-2004: CDISC AND HL7 RCRIM WG

The idea of building a Domain Analysis Model (DAM) which documented the shared semantics of the protocol-driven research domain began in September 2003. Following a discussion of the benefits, risks, and effort involved in pursuing the path of building a DAM, the CDISC Board of Directors (BoD) elected to begin funding the development of a CDISC DAM. The process officially started with a scope definition in March 2004 and the resulting DAM becoming the BRIDG Model around one year later.

2004-2005: THE NCI CABIG® (AND CDISC) STRUCTURED PROTOCOL REPRESENTATION TEAMS

In February 2004, NCI began a project to develop a structured model of a clinical trial protocol as part of the cabiG® project. The search of other interested groups naturally led to interactions with representatives from both CDISC and the HL7 RCRIM WG. The first joint modeling sessions for a “CDISC Domain Analysis Model project” involving both members of the CDISC DAM team and the NCI team were held at the end of 2004. This initial effort, the combination of the work that CDISC had done during the first 9 months of CDISC Domain Analysis Model Project and the first attempts to integrate the work of the NCI team, resulted in the first DAM of the domain of protocol-driven research.

2005-2006: MATURATION OF THE BRIDG PROJECT

The project was officially renamed “BRIDG” in the summer of 2005. BRIDG, which stands for “Biomedical Research Integrated Domain Group” was chosen to reflect the project’s contribution to “bridge” between the various stakeholder organizations. A more formal organizational governance structure must be defined. Interest Groups like HL7, W3C, and OASIS have been involved. An early version of the BRIDG Model was presented to the HL7 RCRIM WG in late-2005/early-2006 and was subsequently accepted as the single Domain Analysis Model for use by the HL7 RCRIM WG.

HL7 and CDISC decided to reverse engineer all of their standards (HL7: implementation-independent semantics from RCRIM WG specifications – CDISC: all existing CDISC specifications [e.g. SDTM, ODM, etc.]).

All four of the BRIDG stakeholder organizations are where involved with applying “BRIDG in context” in a number of application and/or data interchange projects focused on computable semantic interoperability and created the Board of Directors for additional strategic oversight and a Technical Harmonization Committee to manage the operational details in 2005.
2007 – 2012: BRIDG EVOLUTION

As BRIDG became more visible in the community, the BRIDG SCC started hearing feedback from both domain experts and technologists. Many domain experts said that the model was too large, too technical, and did not represent their semantics in language they could understand. The technologists said that the model was not clear enough to map cleanly to downstream models (e.g., design models, implementation models, etc.). As a result, the BRIDG BoD and SCC designed a BRIDG solution that would present each audience of BRIDG with its own perspective.

In addition, the BRIDG BoD decided in late 2008 to move BRIDG forward as an international standard through the ISO and has been forwarded through CDISC

2013 – PRESENT: BRIDG SCOPE EXPANSION – TRANSLATIONAL RESEARCH

Following a gap in BRIDG project in late 2012 to early 2013, during which the BRIDG BoD was reorganized into the BRIDG Advisory Panel, the leadership of BRIDG determined that an NCI model called the Life Sciences Domain Analysis Model (LS DAM) should be harmonized with BRIDG. The LS DAM harmonization introduced 3 new sub-domains covering concepts related to biospecimens, molecular biology and experiments.

In 2014, a new work group called the HL7 BRIDG Work Group was started. This work group provides a platform for BRIDG users and implementers to discuss BRIDG from an implementation view point and identify areas of enhancement.

Subsequently, in early 2015, the CDISC Pharmacogenetics and Pharmacogenomics (PGx) domains were harmonized with BRIDG to compliment the related genetic/genomic concepts added to the model for LS DAM. The PGx concepts leveraged and helped flesh out some concepts identified earlier by the LS DAM harmonization.

A WORD ABOUT UML

The BRIDG model is documented using the UML: Unified Modeling Language. By definition through the Object Management Group (OMG), UML is “...is a graphical language for visualizing, specifying, constructing, and documenting the artifacts of a software-intensive system. The UML offers a standard way to write a system’s blueprints, including conceptual things such as business processes and system functions as well as concrete things such as programming language statements, database schemas, and reusable software components.”

UML contains 13 different diagram types with a package diagram on top to divide the model into logical container. From a simple Use case diagram to a complex class diagram - UML is just a ‘language’, to ‘translate’ between different ‘interpretations and views’ of certain domain topics. Technically it can be used to generate further documentation as well as software – but it still is a communication tool between domain and IT experts.

The BRIDG model uses mainly class and activity diagrams.

For a more detailed introduction into UML see the reference section.

LET'S LOOK INTO THE MODEL

If you open up the model the first time, you get overwhelmed with (class) boxes.
And if you look into the documentation, which can be downloaded as part of the package, you will find 1365 pages. Not the best way to start to understand the model.

But if you look into the legend, you will see some color code information about the meaning of certain classes. This is the starting point:

Also the advice, that …
So, means, the model contains sub-views describing, actually 9, sub-domains. One example domain is the Adverse Event (AE) Sub-Domain. This AE sub-view contains all elements and objects defined as 'playing a role in the world of clinical adverse events'. Let's use this sub-view as our beginner's example.

As we can see here, this AE sub-view includes objects (classes). These classes are directly related to the nature of Adverse Events, but include classes and relationships from other models and sub-domains as well. The Adverse Event (AE) sub-domain contains objects from the Common Domain like “Person” and “Animal”, from the protocol representation “DefinedAdverseEvent” and “DefinedObservation” for example and from the Study Conduct domain “PerformedActivity” and “PerformedSubstanceAdministration”. The Adverse Event sub-domain itself defines the elements of an AE: sure, the class “AdverseEvent” itself but also classes like “AdverseEventSeriousness” and “AdverseEventOutcomeResult”. All class names are nearly self-explaining the class meaning.

Every class definition in BRIDG contains the list of attributes. An attribute in a class is defined by its name and the data type. The class “AdverseEvent” contains, for example, an attribute called “reportedDate”, which is defined as data type TS.DATETIME (TS=time stamp).
The possible data types are defined by HL7 Abstract Data Types. Examples are:

- **ST** = String, **BL** = Boolean, **INT** = Integer, **TS.DATETIME** = timestamp / date time
- **EN** = Entity Name, **AD** = Address
- **CD** = Concept Descriptor (code list)
- **PQ** = Physical Quantity

A Concept Descriptor (CD) is a reference to a concept defined in an external code system, terminology or ontology.

An example of a CD can be seen in the “StudyProtocolVersion” class and the “phaseCode” attribute:

The phaseCode is defined by clinicaltrials.gov and contains some of the following values:

- **N/A**: for trials without phases
- **Phase 0**: exploratory trials, involving very limited human exposure, with no therapeutic or diagnostic intent (e.g., screening studies, microdose studies).
- **Phase 1**: includes initial studies to determine the metabolism and pharmacologic actions of drugs in humans, the side effects associated with increasing doses, and to gain early evidence of effectiveness; may include healthy participants and/or patients
- **Phase 1/Phase 2**: for trials that are a combination of phases 1 and 2
- Etc.

Now, this should be stored in systems as
Another key concept in BRIDG is the concept of ‘ACTIVITY’.

An Activity is an action that can, in the context of a study or a post-marketing investigation, be defined, planned, scheduled or performed.

There are 4 activity subtypes:

- Defined activity
  - An activity that frequently occurs in studies (e.g. more than one time in more than one arm) and therefore is called out as a reusable template in a global library of activities outside the context of any particular study, and may be used in the composition of a defined study subject activity group. A defined activity is a "kind of" activity rather than an "instance of" an activity.

- Planned activity
  - An activity that is intended to occur or start at some point in the context of a particular study.
  - For example, Pregnancy tests are planned for StudySubjects who are females of childbearing potential.

- Scheduled activity
  - An activity that is anticipated to occur at some time in the future and has been assigned a time or date when that activity is to be performed.
  - EXAMPLE(S): An X-Ray scheduled for February 15 is in state "Scheduled." If John is unable to have the X-Ray on that date, the X-Ray would either be rescheduled (remain in "Scheduled" state, but "date" attribute would change) or moved to state "Canceled."

- Performed activity
  - An activity that is successfully or unsuccessfully completed.
  - EXAMPLE(S): CBC performed on a specific StudySubject on a given day.
If you are familiar with HL7 'moodCode' then BRDIG activities are following the same idea.

Finally, the relation between classes is classified by constraints and multiplicities (0..1:1, 1:n, m:n etc). Also, some classes are defined as 'abstract', where another class inherits from. An example from the real world would be the abstract class 'car' with its real implementation as a 'passenger car' or a 'lorry'. It inherits all attributes from the abstract class (example: coachwork, steering wheel) and can add their own implementation attributes, an attribute which is just important for this class (example for a passenger car: backdoors)

**MAPPING EXAMPLE**

To understand how the classes and attributes are mapped you have 2 possibilities: Start in the model to see to which target models the BRDIG classes and attributes has been mapped to, or start in the mapping spreadsheet to start with a target model and get the classes and attributes from BRIDG. The mapping spreadsheet is part of the BRIDG package.

BRIDG is mapped against the following models or targets, parts of the 'Biomedical Research' domain:

- LSDAMv2.2.3Plus
- SDTM IGv3.1.3x
- PGxIG r1.0
- CTRPv3.8
- HCTv1.0 2012
- FDA HL7 SD SD DSTU2012
- Statisticsv1.0
- CTRv1.0
- CTRRr3
- NCI CRF Round 3
- NCI CRF Rounds 1&2
- CTSA-HSDBv1.0
- HCTv1.0 Apr2011
- SDTM IGv3.1.2
- CDASHv1.1
- ICSRr2
- C3PRv2.9
- PScv2.6
- caAERSv2.2
- LabViewer2.2
- CTR&R R2
- COPPA
- Lab
- PSC
- AE
- CTR&R R1 Comprehensive
- SDTM IG
- TDM
- Study Design RMIM
- HL7SP
- RPS1
- C3PR
- CTOM

For every target a single worksheet exists and defines the single classes and attributes.
To understand the syntax used in the sheet, a Mapping Path definition is also available.

The opposite way is to start in the BRIDG model. For elements which are mapped a TAG has been created in Enterprise Architect. As an example, multiple TAGs have been created for the “severityCode” in Adverse Events.
to define the mapping to the target model. In the following screenshot you can see that the “severityCode” has been mapped to:

CDASHv1.1
AE
SDTMIGv3.1.1
PSCv2.6
SDTMIGv3.1.2
CTRv1.0

However, one problem is that some mapping targets are missing. In this particular example we see in the mapping spreadsheet, that “severityCode” is mapped against SDTM IGv3.1.3, but this target version is not listed in the TAG session in Enterprise Architect:

WHAT’S NEW IN VERSION 4
With the release of BRIDG 4.0, the scope of BRIDG is now officially expressed as "basic, pre-clinical, clinical, and translational research and associated regulatory artifacts. 4.0 is the first release of BRIDG as a translational research model and therefore some of the newer sub-domains are not fully fleshed out as yet.

In Version 4 the LS DAM (Life Sciences Domain Analysis Model) has been added. It is a Hypothesis driven basic and pre-clinical research model and supports discovery sciences as well. Over years it has been developed in parallel to BRIDG. The main stakeholder was the NCI, working closely with the HL7 Clinical Genomics workgroup.

Further new in Version 4 are 3 domains:
- Biospecimen
- Experiment
- Molecular Biology

Version 4 added also related SDTM changes: trial design changes, inclusion of the therapeutic area Oncology and the content published in the SDTM amendment 1. Also, SDTM PGx has been added – Pharmacogenomics and Pharmacogenetics.

CONCLUSION / SUMMARY
BRIDG is the canonical model for the Life Sciences Domain. It uses UML as the ‘language’ to define the model.

For a newbie it can be overwhelming the first time, but diving deeper it is not so complicated to start and understand the model. Once the concept of classes and attributes and the grouping into sub-domains has been understood it can be simple to understand the mappings to different target models like CDISC SDTM and ICSR r2 as well.

With Version 4 the NCI LS DAM has been added and BRIDG is awaiting fully ISO approval and publication over the next few months.

REFERENCES
BRIDG home: http://bridgmodel.nci.nih.gov/
BRDIG history: http://bridgmodel.nci.nih.gov/about_bridg/background
HL7: http://www.hl7.org/special/committees/bridg/index.cfm
Definition of CD – Study Phases on clinicaltrials.gov: https://prsinfo.clinicaltrials.gov/definitions.html
Enterprise Architect, Sparxsystems: http://www.sparxsystems.com/

GLOSSARY
BRIDG - Biomedical Research Integrated Domain Group
DAM – Domain Analysis Model
UML – Unified Modelling Language
NCI – National Cancer Institute
FDA – US Food and Drug Administration
HL7 – Health Level 7
CDISC – Clinical Data Interchange Standards Consortium

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http://www.cdisc.org/bridg
http://bridgmodel.nci.nih.gov/
http://www.hl7.org/special/committees/bridg/index.cfm
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