



Allotrope
Foundation
Building an Open Framework for Laboratory Data

Pharmaceutical companies collaborate on building a Framework to manage Analytical Data more efficiently

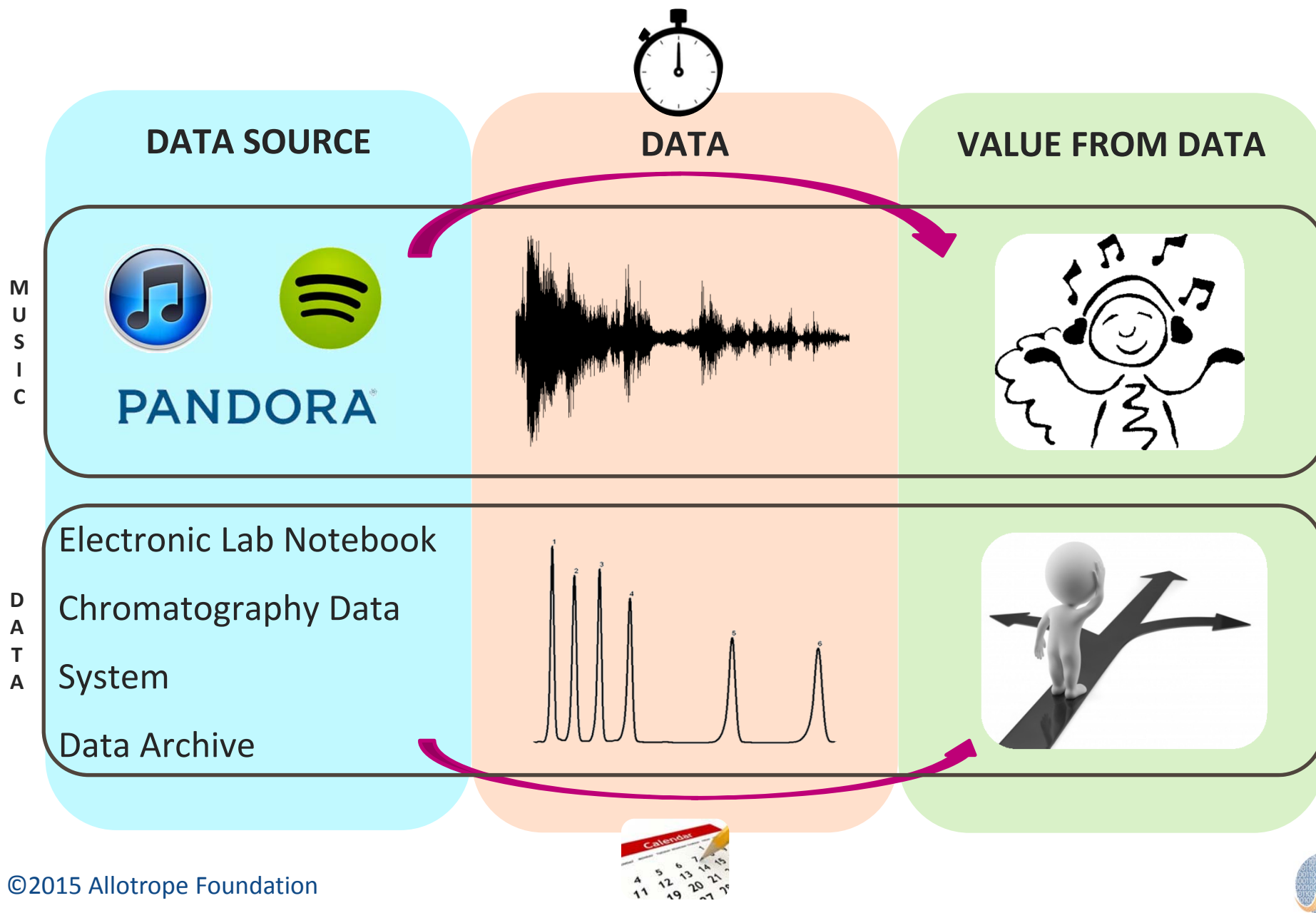
Dr. Gerhard Noelken
Allotrope BoD member, Pfizer Allotrope Liaison

GINAS Symposium, Uppsala
7 September 2015



- **MOTIVATION**
- THEORY
- REDUCING IT TO PRACTICE

What is the problem we are trying to solve?



Why is access to music so much easier than access to scientific data?

Think about music...

Music is typically stored in a small number of *standard, non-proprietary* formats...



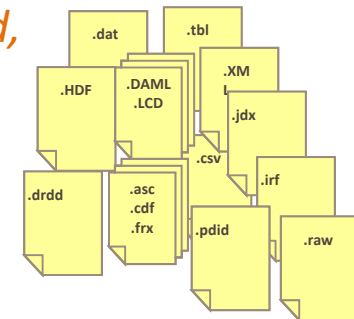
...with contextual *metadata* that are *complete, consistent & correct*

| | |
|--------|---------|
| Artist | Genre |
| Album | Date |
| Song | Artwork |

...enabling the user to *find, share* and *enjoy* it years later from any device *easily*!

Think about scientific data...

Scientific data is typically stored in a wide variety of *non-standard, proprietary* formats...



...with contextual *metadata* that are *hard to find and sometimes inconsistent*

| | |
|-----------|---------|
| Material | Process |
| Equipment | Result |

...making it *costly and sometimes difficult* to find and get value from it.

What if scientific data was as easy to access as music?

If we...

- Store scientific and process data in a *standard format* with *contextual metadata* that is...

- correct
- complete
- consistent
- compliant



We could...

- *Find data in seconds.*
- *Be confident* that the *data* that underpins our *decisions* is *accurate*, *complete*, and *compliant*.
- *Build data quality and data integrity into the system*, eliminating the need for many SOPs and quality investigations.
- *Simplify, automate* and *improve* laboratory and manufacturing processes.
- *Automatically create technical reports*, audit trails, and substantial portions of regulatory submission documents.
- *Answer complex questions*, not just those accessible via simple queries - by linking data from diverse, disparate sources.

Allotrope Foundation

Member Companies

AbbVie
Amgen
Baxter
Bayer

Biogen
Boehringer Ingelheim
Bristol-Myers Squibb
Eli Lilly

Genentech/Roche
GlaxoSmithKline
Merck & Co.
Pfizer

Secretariat

DrinkerBiddle

- Project Management
- Legal & Logistical Support

Professional Software Firm

OSTHUS
success with R&D

- Framework Development
- Technical Leadership

Partner Network



ACD/Labs
Agilent
Biovia
BSSN
IDBS

Mettler Toledo
Persistent
Riffyn
Sartorius
Shimadzu

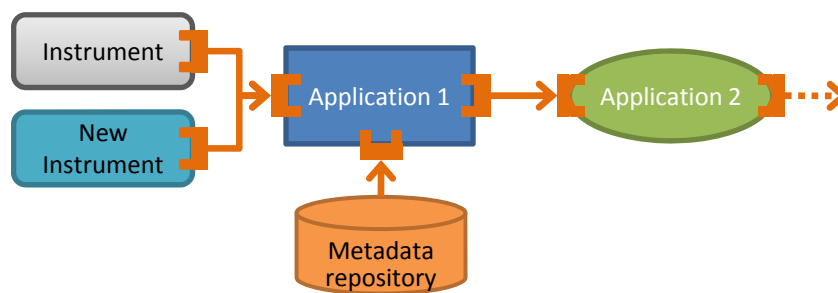
Waters

Erasmus Univ.
Med Center
University of
Southampton

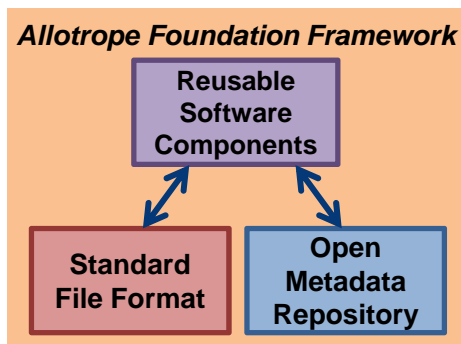
Mestrelab Research Thermo Scientific

- MOTIVATION
- **THEORY**
- REDUCING IT TO PRACTICE

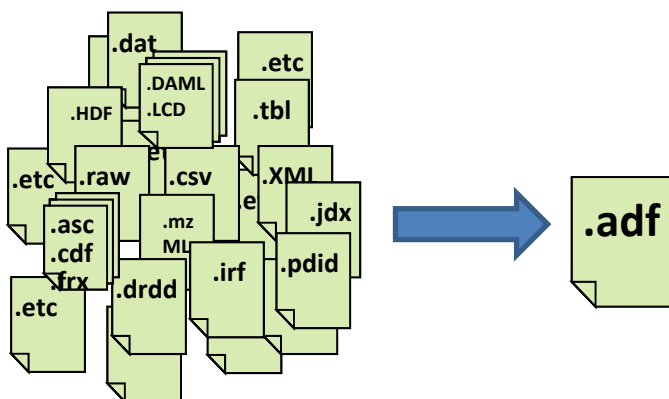
What is Allotrope Creating?



A **toolkit** that enables use of the standards & metadata in software development



File format for any technique or instrument



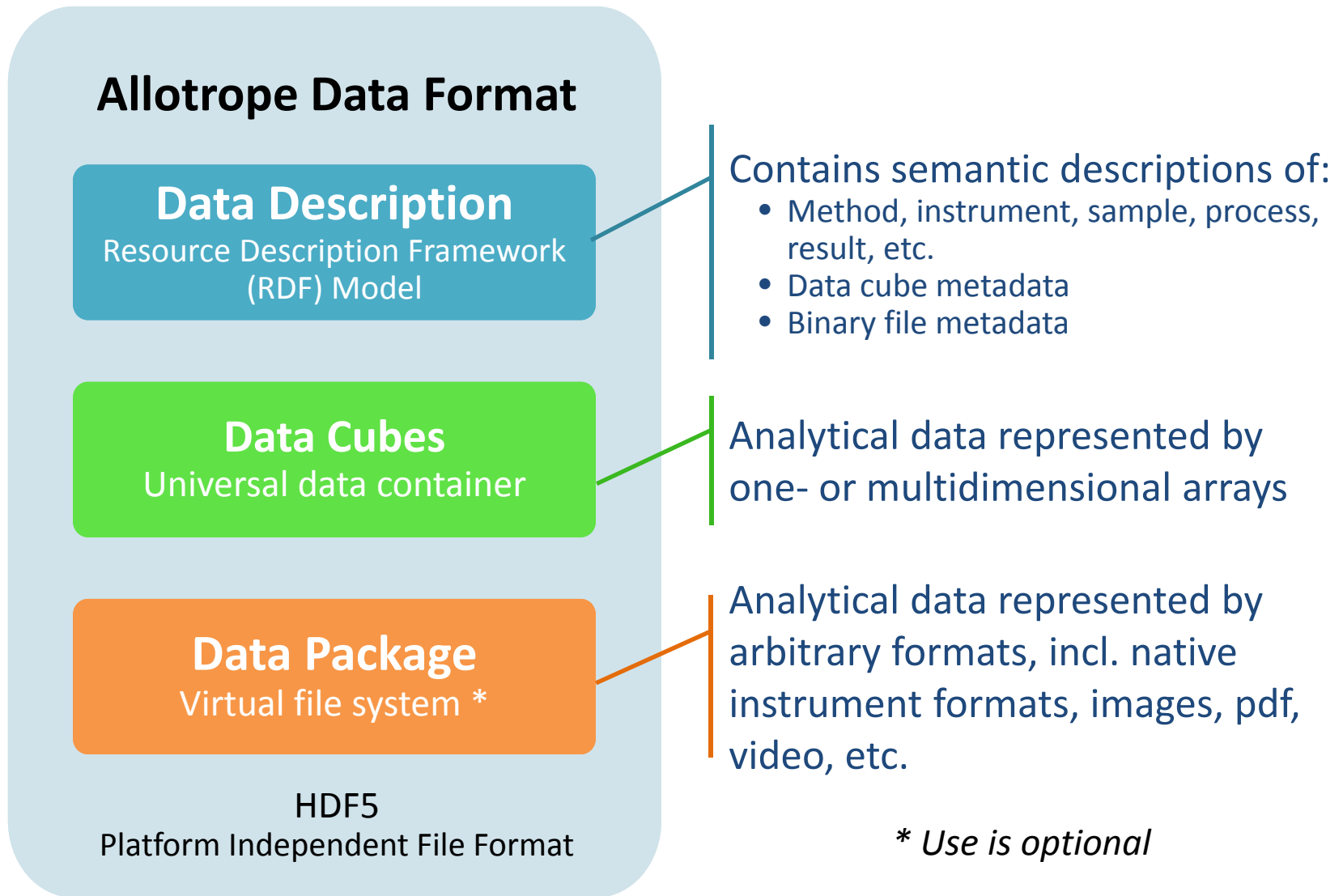
Standard vocabulary & structure for metadata

| Project | Test | Instrument |
|------------|-----------------------|------------------------|
| AF 0012354 | IR Fingerprinting | QC Lab #33B 380 FT-IR |
| AE0012764 | Bulk & Tapped Density | ASTM Standard Sieve #6 |
| AF 12989 | NMR Characterization | AM500 |
| | Tapped & Bulk Density | Sieve XXX |
| AF0045674 | Caractérisation RMN | Nouvelle DRX600 |
| AF-0034558 | IR | iS10 FT-IR |

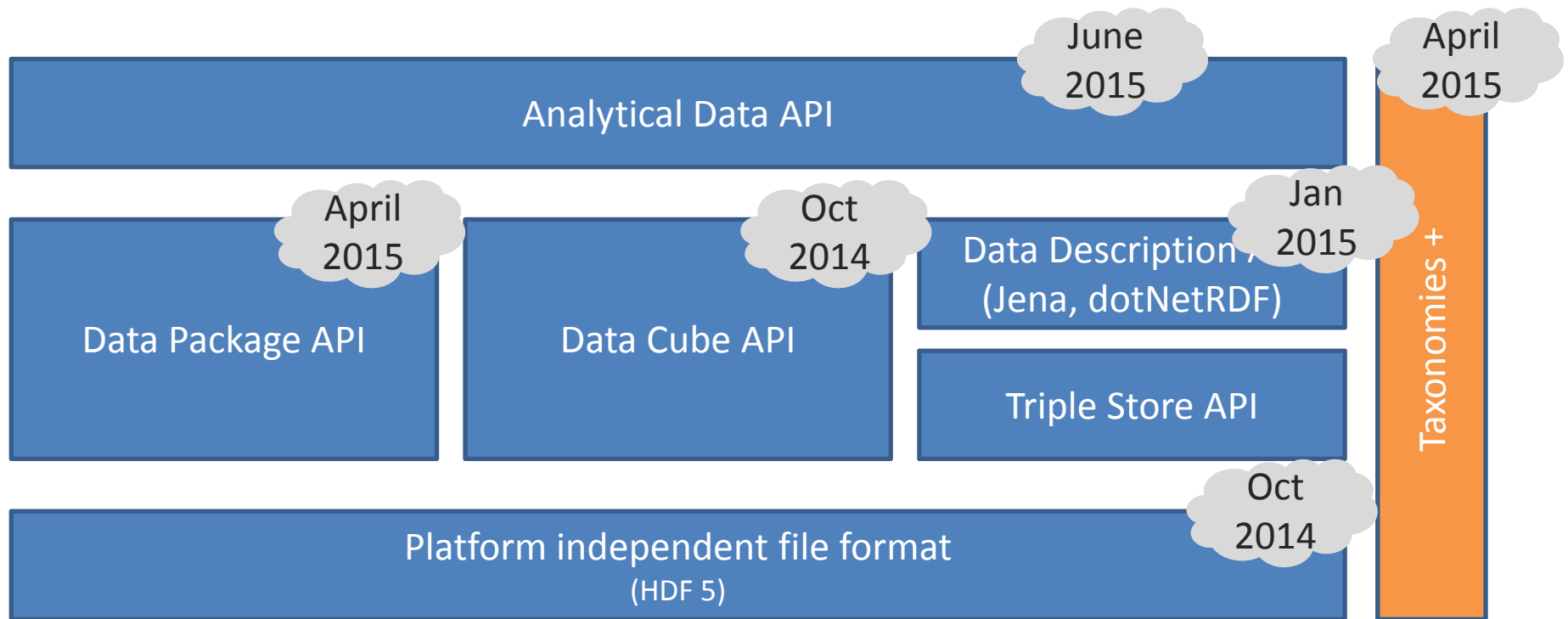
With the Metadata Repository

| Project | Test | Instrument |
|-----------|-------------------------|----------------------|
| AF0012354 | IR Fingerprinting | 380 FTIR/-SN/145453 |
| AF0012764 | Bulk and Tapped Density | ASTM Sieve-SN/3452 |
| AF0012989 | NMR Characterization | AM500-SN/0034578 |
| AF0013142 | Bulk and Tapped Density | ASTM Sieve-SN/09783 |
| AF0045674 | NMR Characterization | DRX600-SN/10234567 |
| AF0034558 | IR Fingerprinting | iS10 FTIR/-SN/341980 |

ADF: a universal file format for scientific data



ADF Class Libraries + Decoupled Taxonomies

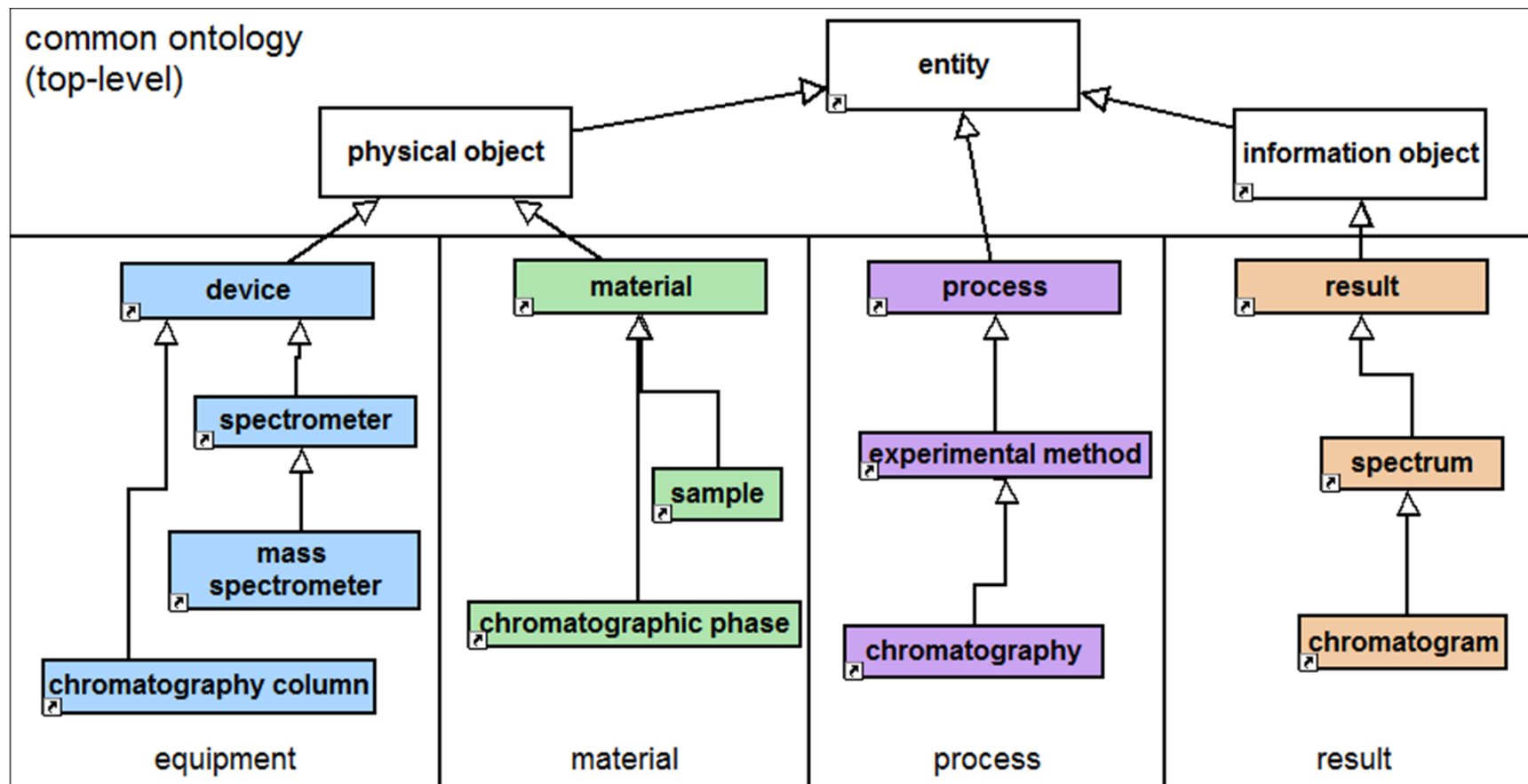


Languages: Java, C#

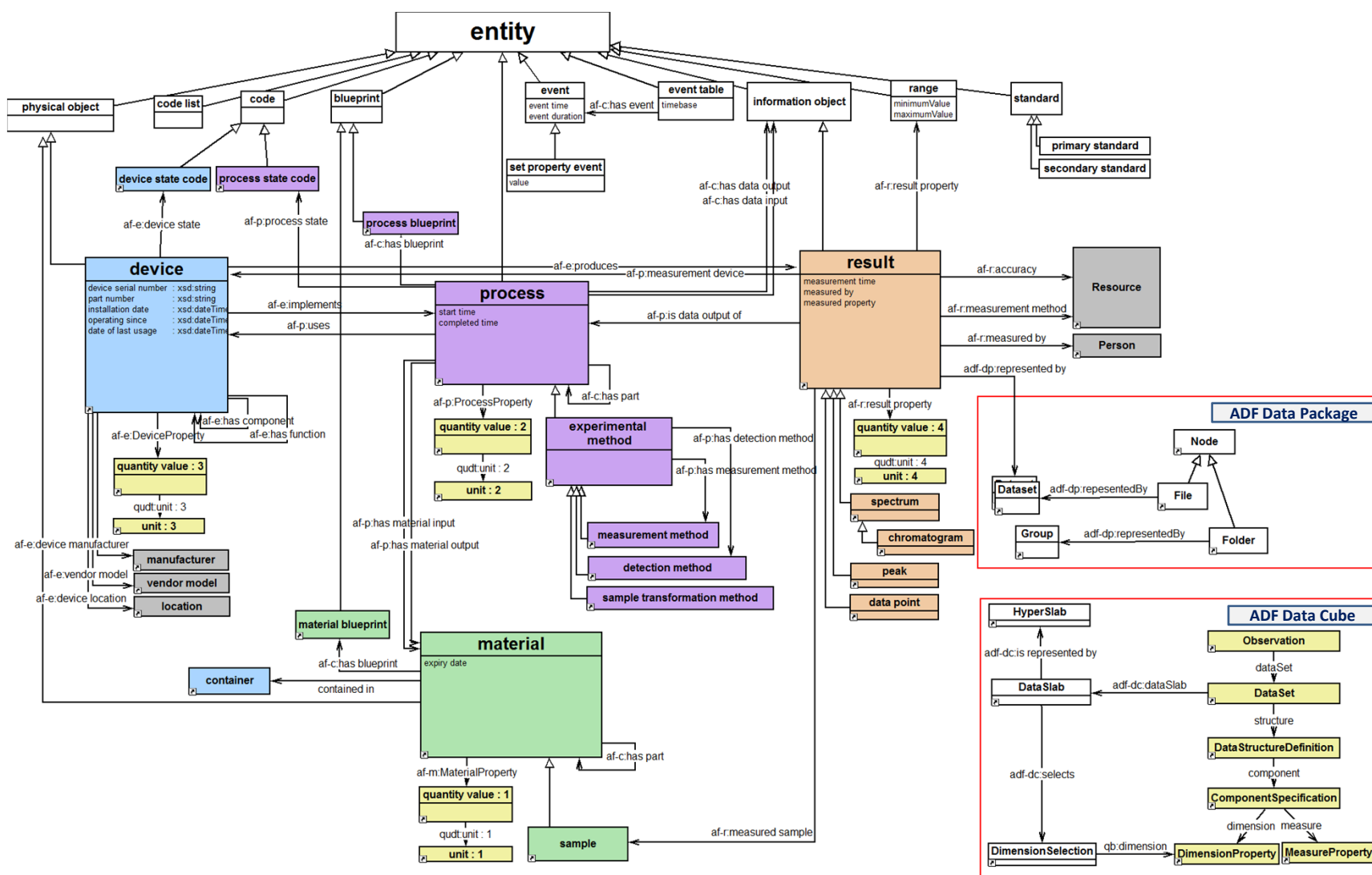
Allotrope Taxonomies: An Extensible Metadata Model

- **A library of extensible taxonomies**
 - Uses W3C standards
 - Easy to understand and maintain by SMEs and Vendors
- **Start by harvesting existing available concepts**
 - PSI-MS; IUPAC; RSC Chemical Methods Ontology; Dictionary of weighing terms; AnIML, etc
- **Reproducible & efficient collaboration model**
 - Leverages knowledge engineers & member company scientists
 - 2-3 weeks to develop initial version of a new taxonomy
- **Initial versions of 12 analytical techniques already implemented:**
 - gas chromatography
 - Karl Fischer
 - liquid chromatography
 - mass spectrometry
 - nuclear magnetic resonance spectroscopy
 - thermogravimetric analysis
 - ultra violet spectrometry
 - cell counter
 - cell culture analyzer
 - blood gas analysis
 - balance
 - pH

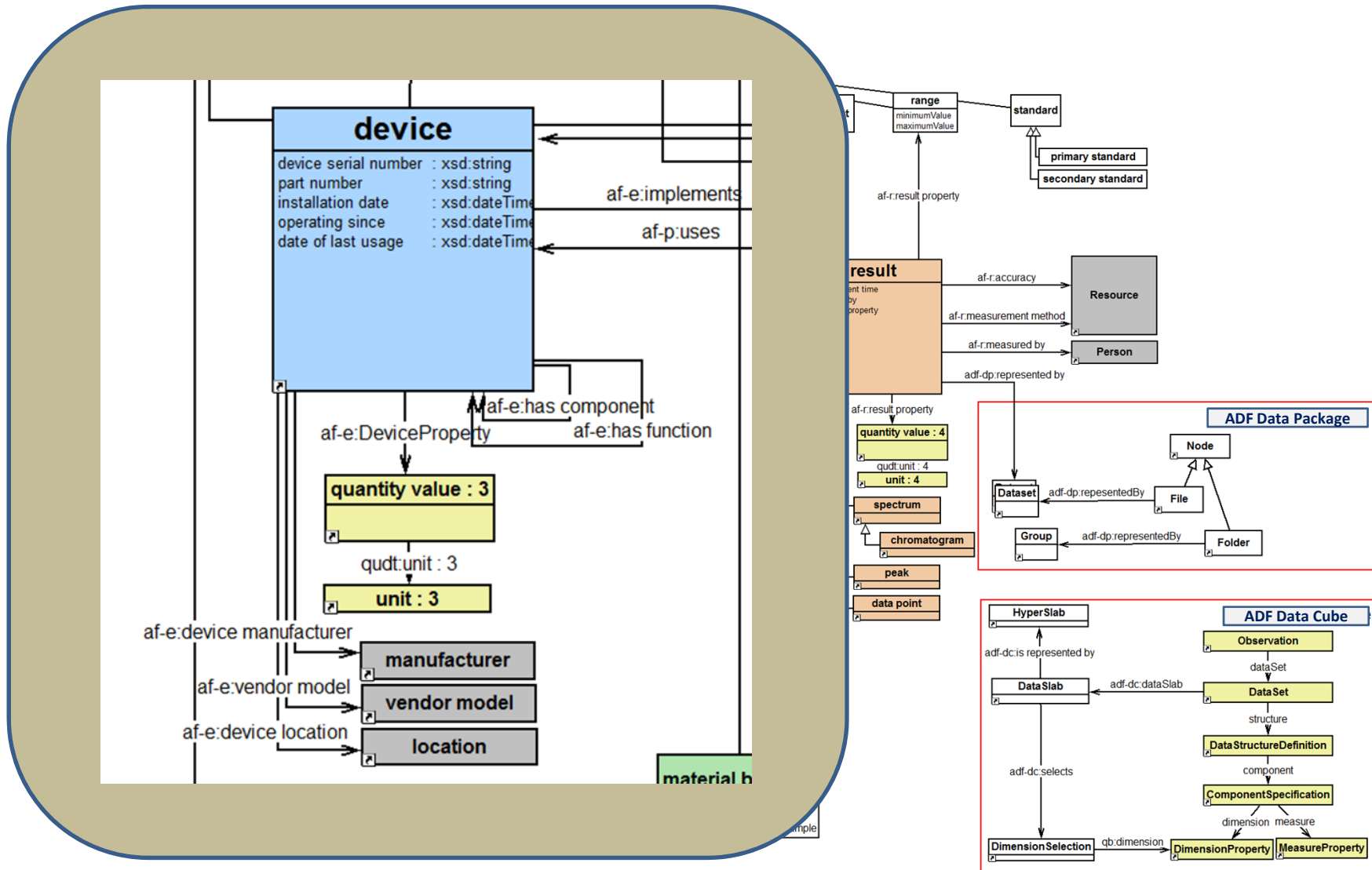
Allotrope Foundation Taxonomies



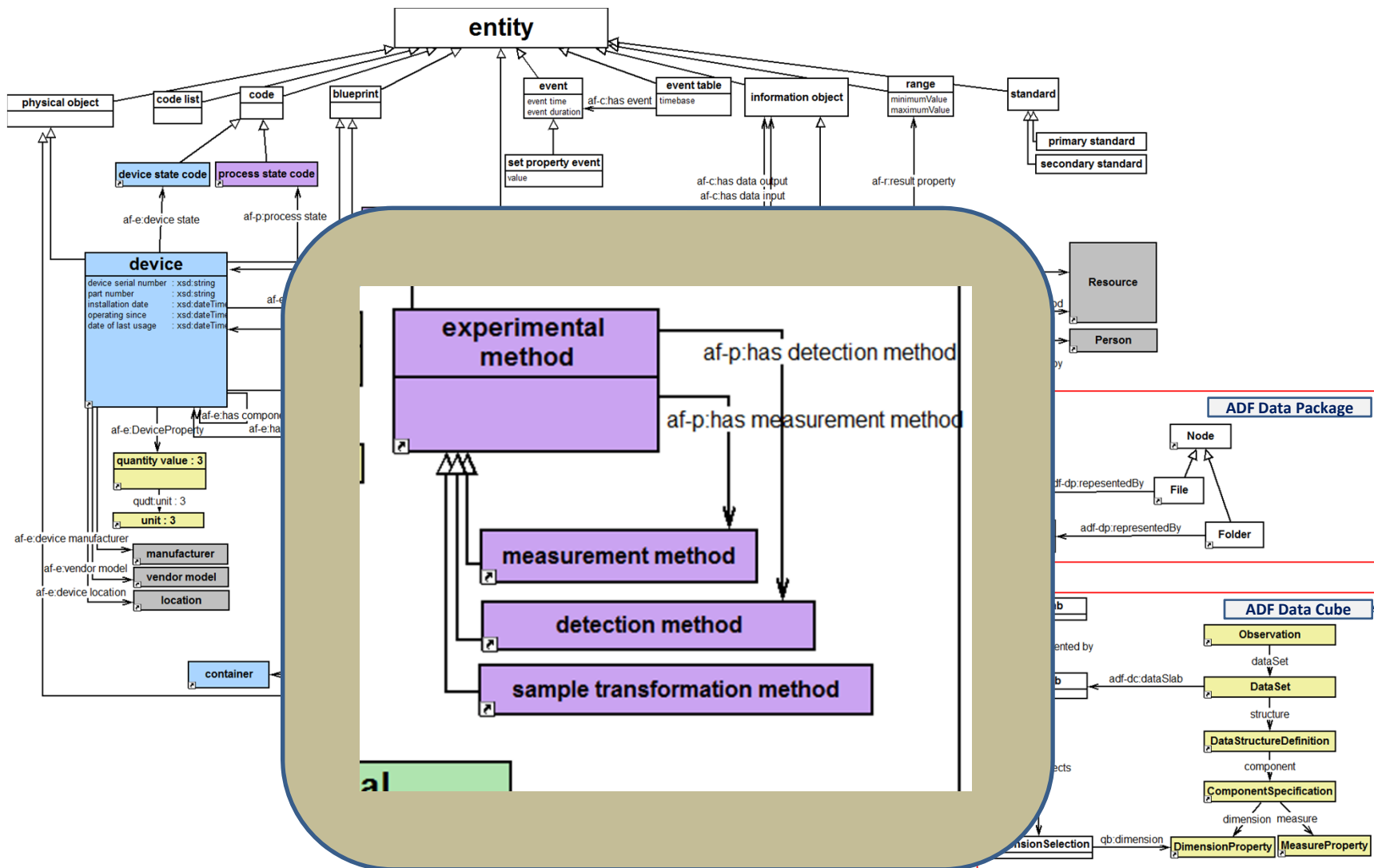
The Big Picture

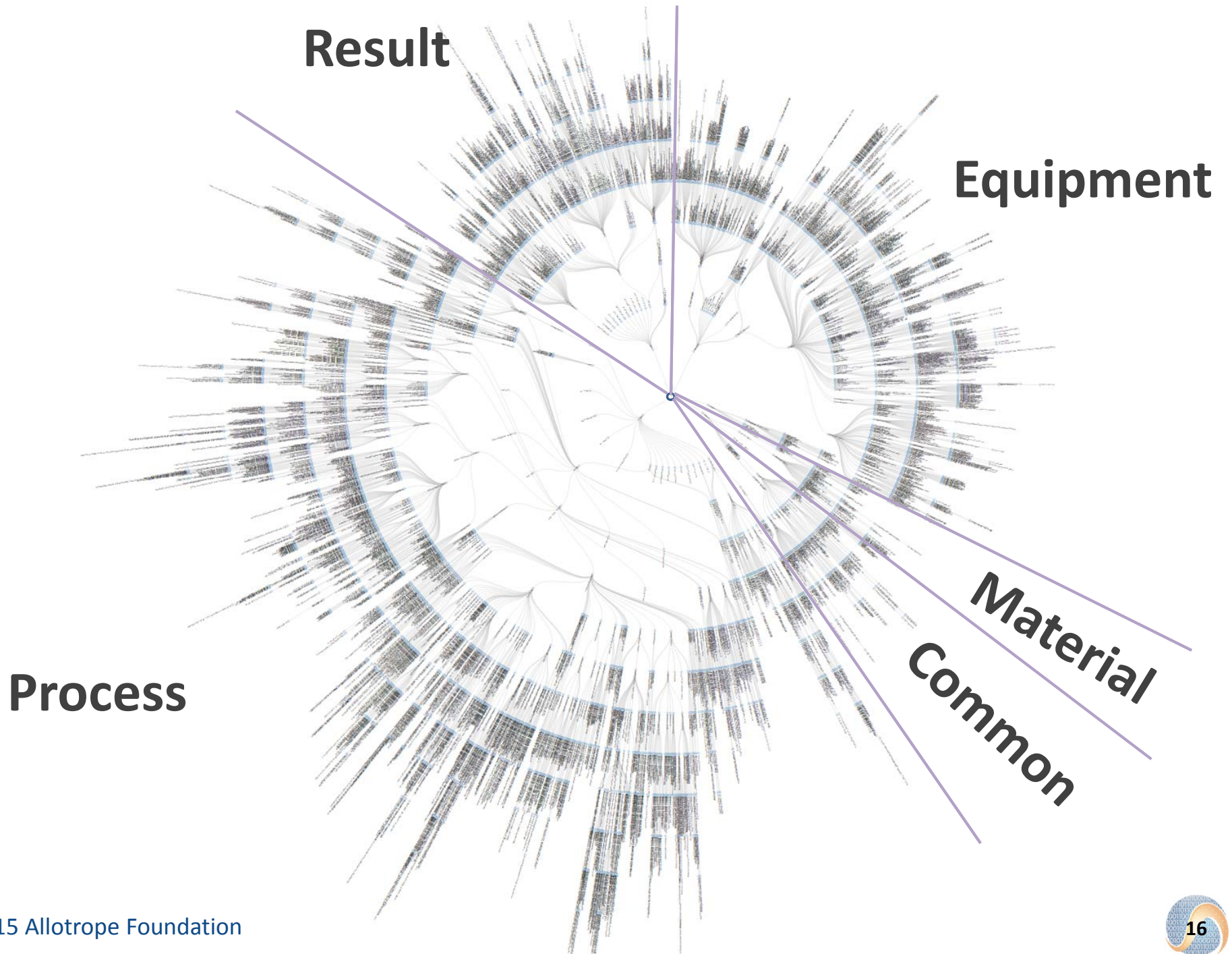


The Big Picture



The Big Picture





AF Taxonomies Documentation

Contents

- All Ontologies
- Classes (2303)
- Object Properties (245)
- Data Properties (51)
- Annotation Properties (31)
- Individuals (58)
- Datatypes (6)

skn

- 'af-p:mass spectrometer calibration'
- 'af-p:mass spectrometry data processing'
- 'af-p:mass spectrometry of recoiled ions'
- 'af-p:mass spectrometry scan'
- 'af-p:mass spectrometry'
- 'af-p:matrix solid-phase dispersion'
- 'af-p:matrix-assisted laser desorption electrospray ionization'
- 'af-p:matrix-assisted laser desorption-ionization imaging mass spectrometry'
- 'af-p:matrix-assisted laser desorption-ionization mass spectrometry'
- 'af-p:matrix-assisted laser desorption/ionization'
- 'af-p:McLafferty rearrangement'
- 'af-p:mean of spectra'

process

Class: 'af-p:mass spectrometry'

http://purl.allotrope.org/ontologies/process#AFP_0001427

Annotations (15)

- 'skos:alternative label' "MS" ()
- 'skos:alternative label' "mass spectroscopy" ()
- 'skos:change note' "0.3 ADD imported from CHMO [OSTHUS]" ()
- 'skos:change note' "0.7 DELETE hyphenations: CollisionInducedMassSpectrometry [OSTHUS]" ()
- 'skos:change note' "0.8 CHANGE definition to IUPAC [OSTHUS]" ()
- 'skos:change note' "0.8 DELETE ion mobility spectrometry - mass spectrometry hyphenated method [OSTHUS]" ()
- 'skos:change note' "0.8 DELETE ionization as classification for MS [OSTHUS]" ()
- 'skos:change note' "0.8 DELETE selective ion monitoring (duplicate) [OSTHUS]" ()
- 'skos:change note' "0.8 DELETE subclasses of molecular mass spectrometry and atomic mass spectrometry [BAYER]" ()
- 'skos:change note' "0.8 MOVE also under mass analysis [OSTHUS]" ()
- skos:definition "Study of matter through the formation of gas-phase ions that are characterized using mass spectrometers by their mass, charge, structure, and/or physico-chemical properties. [IUPAC MS REC]" ()
- skos:note "Mass spectroscopy is an obsolete synonym for mass spectrometry that should not be used to avoid confusion with spectroscopies in which the measured quantity is the absorption or emission of electromagnetic radiation. [IUPAC MS REC]" ()
- skos:note "The term is a misnomer because it is m/z rather than mass that is the independent variable in a mass spectrum." ()
- 'skos:preferred label' "mass spectrometry" ()
- 'skos:scope note' "MS" ()

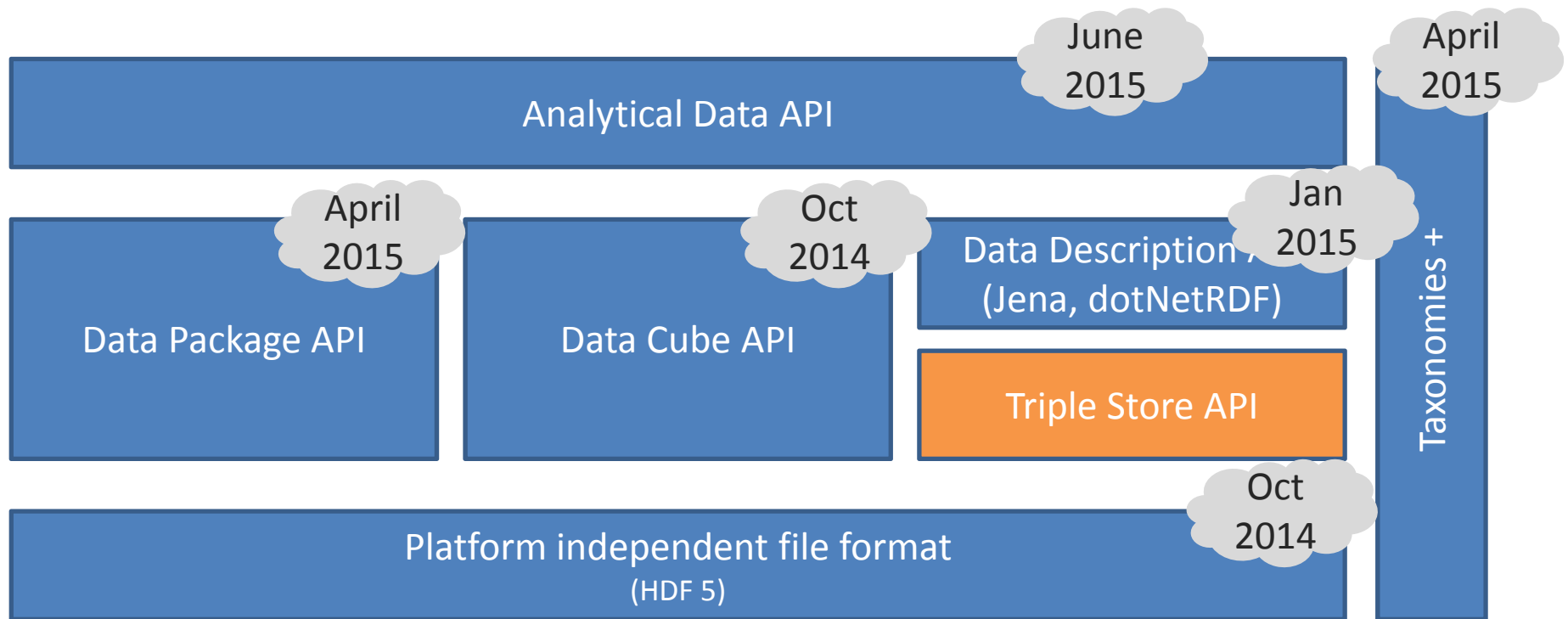
Superclasses (2)

- 'af-p:mass analysis'
- af-p:spectroscopy

Usage (6)

- Class: 'af-p:mass spectrometry'
- 'af-p:sustained off-resonance irradiation' $\hat{a}\hat{S}^+$ 'af-c:is part of' some 'af-p:mass spectrometry'
- 'af-p:stored waveform inverse fourier transform' $\hat{a}\hat{S}^+$ 'af-c:is part of' some 'af-p:mass spectrometry'
- 'af-p:time lag focusing' $\hat{a}\hat{S}^+$ 'af-c:is part of' some 'af-p:mass spectrometry'
- 'af-p:mass spectrometry property' Domain 'af-p:mass spectrometry'
- 'af-p:ionization polarity' Domain (af-p:ionization or 'af-p:mass spectrometry')

ADF Class Libraries + Decoupled Taxonomies



Languages: Java, C#

RDF Data Model

- Subject-Predicate-Object (Triple)

- Example:

<Sample 1> type <Sample>

<Sample 1> createdOn '2015-03-13'

<Sample 1> createdBy <person X>

<Sample 1> hasBarcode '1234567890'

↑
Subject

↑
Predicate

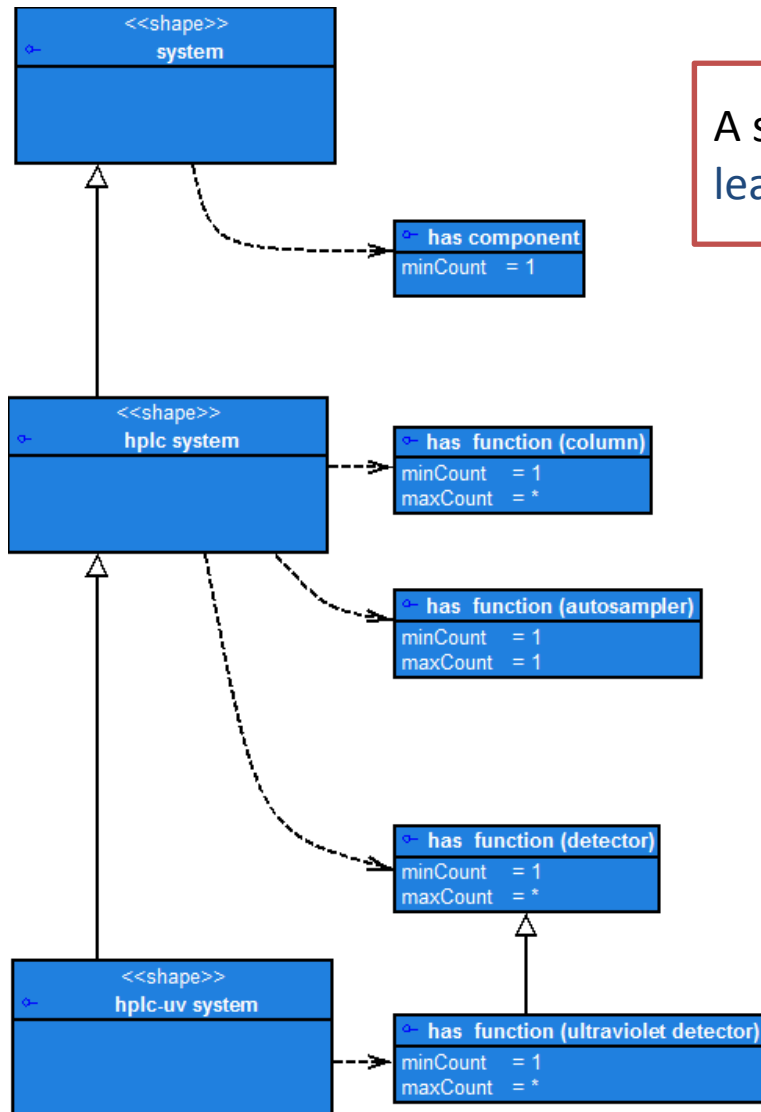
↑
Object

Data Shapes Constrain How We Use Taxonomies in the Real World

- Taxonomies provide an unconstrained vocabulary that we can use to describe things (instances) in our open world and give them a meaning (= **what** it is)
- We need a mechanism to define data structures (schemas, templates) that describe **how** to use the taxonomies for a given purpose in a standardized (= reproducible, predictable, verifiable) way
- Shapes Constraint Language (SHACL, expressed as RDF triples) is an emerging standard to do this

<http://www.w3.org/2014/data-shapes/charter>

Using Data Shapes: Equipment



A system has at least 1 component

Shape hierarchies define **additional** constraints

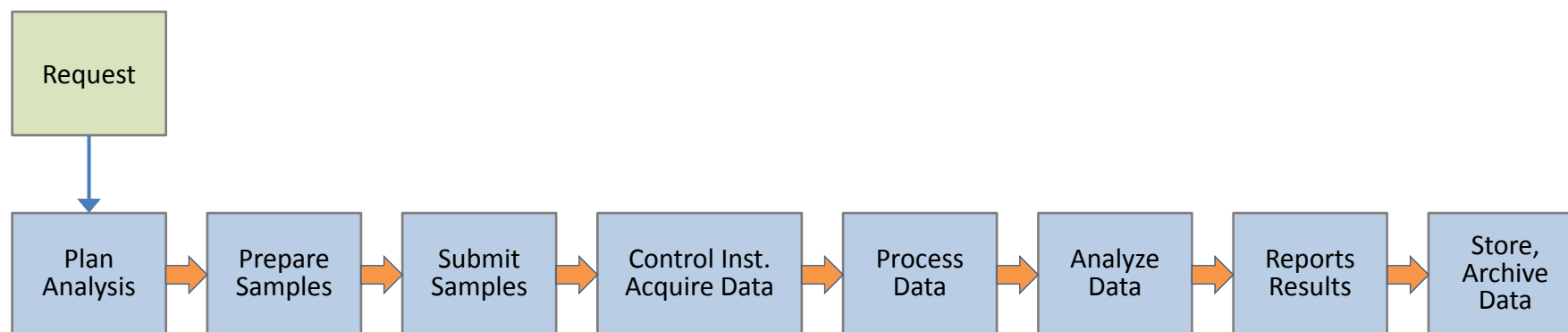
A hplc system has at least 1 component **and** has at least 1 column, exactly 1 autosampler and at least 1 detector

A hplc-uv system has at least 1 component **and** has at least 1 column, exactly 1 autosampler and at least 1 detector **and** at least 1 uv-detector

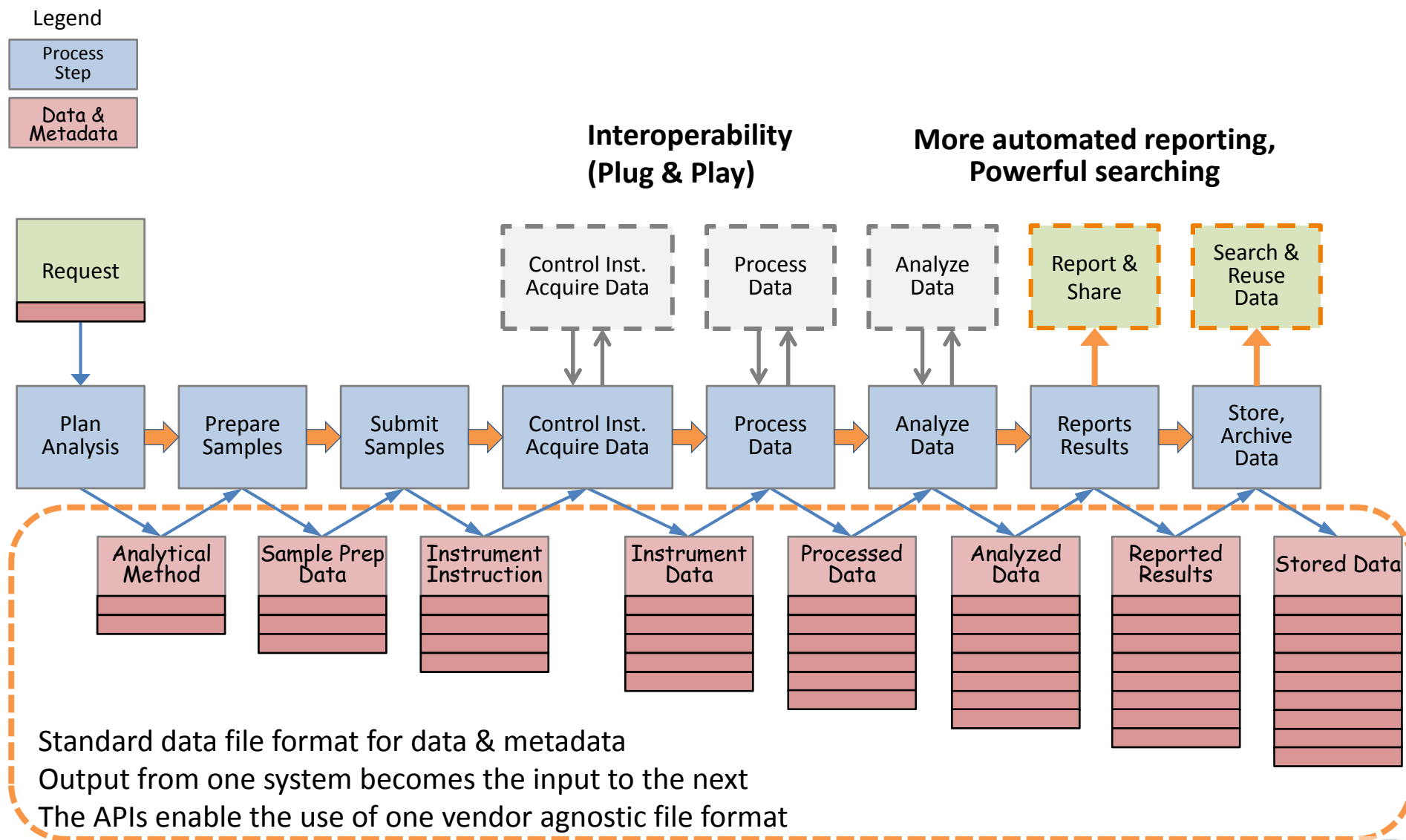
The ADF enables a self-contained documentation of the data & metadata

Legend

Process
Step



The ADF enables a self-contained documentation of the data & metadata



Data Description

Data acquisition

```

<myHPLC/2> a nan;
--
adf-p.Result <myHPLC/1>/chromatogram; <myHPLC/2>/chromatogram;
--
<myHPLC/1>/chromatogram a qb.Dataset; adf-r.Chromatogram;
adf-measurementDevice <HPLC/U/Detector>;
adf-measurability <Iand>;
adf-measurementTime 2015-08-24T14:05:55+1;
adf-qb.representedby <chdf.databases/2D HDF>;
qb.structure <Chromatogram2DCube>;

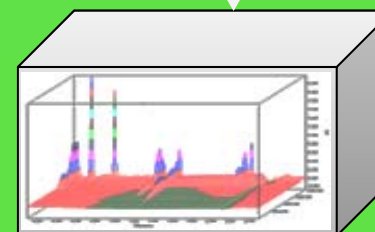
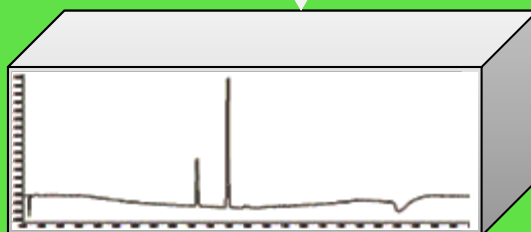
<myHPLC/2>/chromatogram a qb.Dataset; adf-r.Chromatogram;
adf-measurementDevice <HPLC/P/Da>;
adf-measurability <Iand>;
adf-measurementTime 2015-08-24T14:05:55+1;
adf-qb.representedby <chdf.databases/3D HDF>;
qb.structure <Chromatogram3DCube>;

<Chromatogram2DCube> a qb.DataStructureDefinition;
qb.component [ qb.dimension adf-r.Time ]
[ qb.measure adf-r.Absorbance];

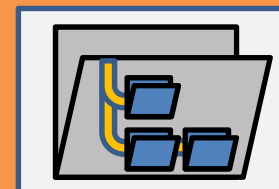
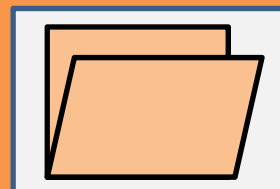
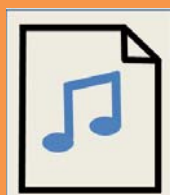
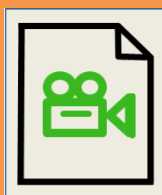
<Chromatogram3DCube> a qb.DataStructureDefinition;
qb.component [ qb.dimension adf-r.Time; qb.order 1 ]
[ qb.dimension adf-r.Wavelength; qb.order 2 ]
[ qb.measure adf-r.Absorbance];

```

Chromatogram: 3D HDF



Data Package(s) (optional)



- MOTIVATION
- THEORY
- **REDUCING IT TO PRACTICE**

Alpha & Beta Release available to AF & APN members now

- **Taxonomies (work in progress)**
 - Mature versions for:
 - MS, LC, pH, weighing, UV
 - Initial versions of:
 - NMR, cell counter, blood gas, capillary electrophoresis, cell culture analyzer, thermogravimetric analysis, Karl Fischer, GC,
 - OWL, OWL + SKOS, Excel and OWLDoc formats
- **APIs**
 - Data Package, Data Description, Data Cube, Analytical Data
- **ADF & API documentation**
- **New Example Applications for the ADF APIs**



Allotrope Data Format

Allotrope First Public Working Draft 30 April 2015

This version:

<http://purl.allotrope.org/TR/2015/WD-adf-20150430/>

Latest published version:

<http://purl.allotrope.org/TR/adf/>

Editor:

Wolfgang Colman, [OSTHUS](#)

Author:

Christoph Weidmann, [OSTHUS](#)

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Abstract

The Allotrope Data Format (ADF) is a family of specifications designed to standardize the acquisition, exchange, storage and access of analytical data captured in laboratory workflows. This document is an overview of ADF. It provides an entry point to its specifications.

Status of This Document

This section describes the status of this document at the time of its publication. Other documents may supersede this document. A list of current Allotrope publications and the latest revision of this technical report can be found in the [Allotrope technical reports index](#) at <http://purl.allotrope.org/TR/>.

This document was published by the [Allotrope Standards Working Group](#) as a First Public Working Draft. This document is intended to become a Allotrope Recommendation. If you wish to make comments regarding this document, please send them to james.vergis@dbi.com. All comments are welcome.

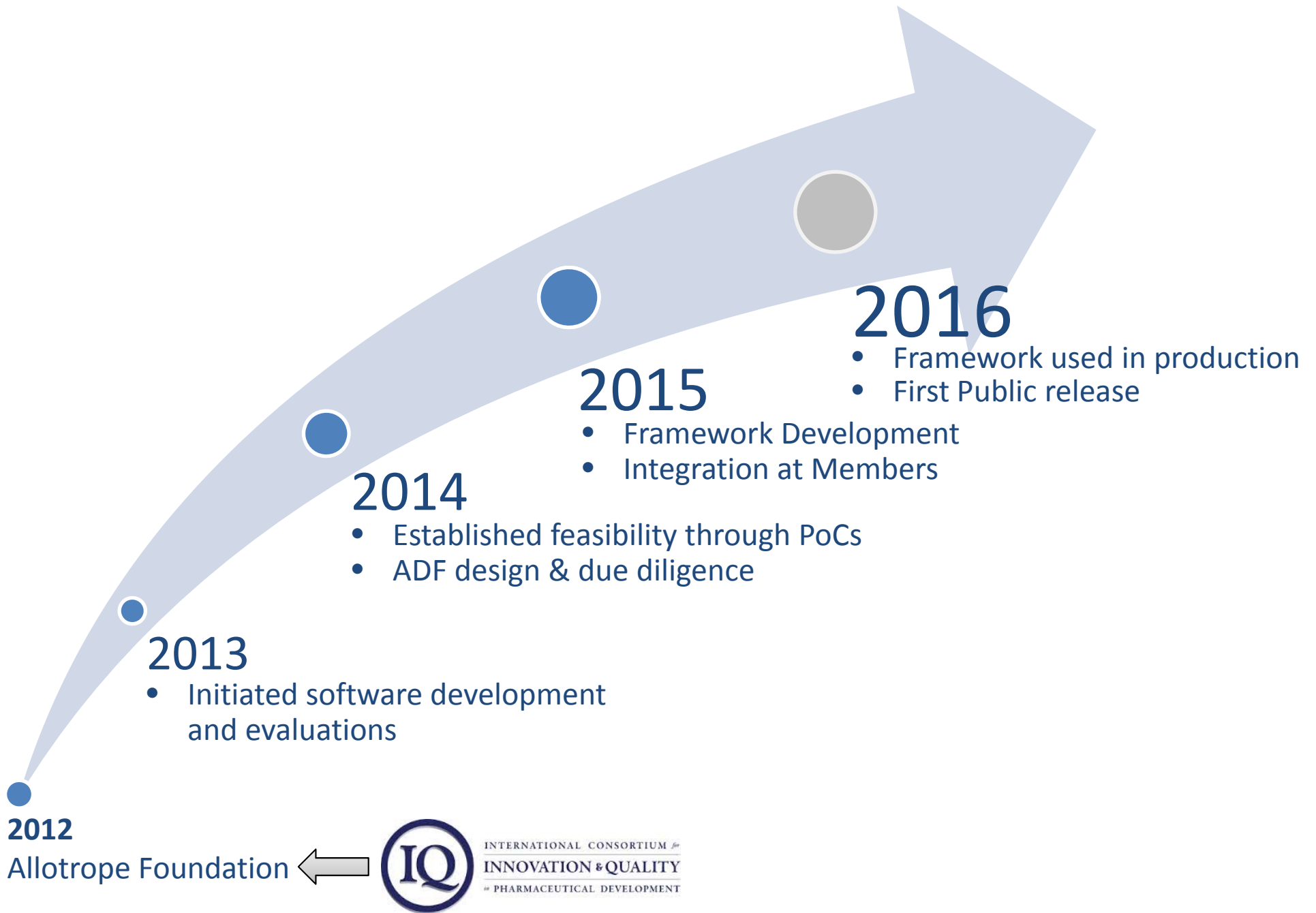
Publication as a First Public Working Draft does not imply endorsement by the Allotrope Membership. This is a draft document and may be updated, replaced or obsoleted by other documents at any time. It is inappropriate to cite this document as other than work in progress.

Table of Contents

- 1. Introduction
- 2. Key Requirements
- 3. ADF high level structure
- 4. ADF API stack
 - 4.1 Analytical Data API
 - 4.2 Data Package API
 - 4.3 Data Cube API
 - 4.4 Data Description API
 - 4.5 Triple Store API
 - 4.6 Platform Independent File format
 - 4.7 Ontologies
- A. References
 - A.1 Normative references

Current Allotrope Integration Project Dashboard

| Project Type | Description | Company | Status | | | | | |
|-----------------------------------|---|---------|--------|---------------|---------|---------------|-----------------|----------------|
| | | | Idea | Business Case | Scoping | Specification | Agile Execution | Implementation |
| Data Converter | <ul style="list-style-type: none"> •A temporary, expedient solution to transform data into ADF to mitigate obsolescence; enables first step to full adoption | A | | | | | | |
| | | B | | | | | | |
| | | C | | | | | | |
| Lab & Plant Automation | <ul style="list-style-type: none"> •Platform for the planning, execution, analysis & reporting of analytical chemistry leveraging the Framework •Includes IoT instrument integration; metadata repository/method management; workflow execution; ADF I/O integration with COTS and in-house software •Enables significant opportunities for automating data flow | D | | | | | | |
| | | E | | | | | | |
| | | F | | | | | | |
| | | G | | | | | | |
| | | H | | | | | | |
| | | I | | | | | | |
| Taxonomies | <ul style="list-style-type: none"> •Leverage Allotrope taxonomies to provide metadata for enriched index •Lightweight universal viewer for any technique | H | | | | | | |
| | | J | | | | | | |
| Data Lake | <ul style="list-style-type: none"> •Repository based on ADF/AT/APIs | H | | | | | | |
| CRO Integration | <ul style="list-style-type: none"> •Convert at CRO to return raw & processed data back to company in ADF | K | | | | | | |



Thank you!

Networking with Peers: upcoming workshops and meetings

- Sep 15, 2015 (Chicago, IL): Allotrope Partner Network F2F Workshop
- Sep 16, 2015 (Chicago, IL): Cross Industry Workshop

To join or get additional information, contact:

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Science Advisor | Drinker Biddle & Reath LLP

1-202-230-5439

James.Vergis@dbr.com

more.info@allotrope.org

www.allotrope.org



What brought me to Uppsala ??

- Meeting Thomas Balzer at an IDMP conference in Berlin
- Allotrope Framework presented as a long-term option to establish a smooth dataflow
 - from analytical data creation in the lab
 - to analytical Data reporting e.g. for Group 4 Specified Substances
- Allotrope closing a gap in terms of missing analytical data standards in the CMC area
- Discussion:
 - Can we identify synergies between Allotrope Framework for analytical data management in the Lab and analytical information in a substance database

Specified Substance Group 4 Analytical Data

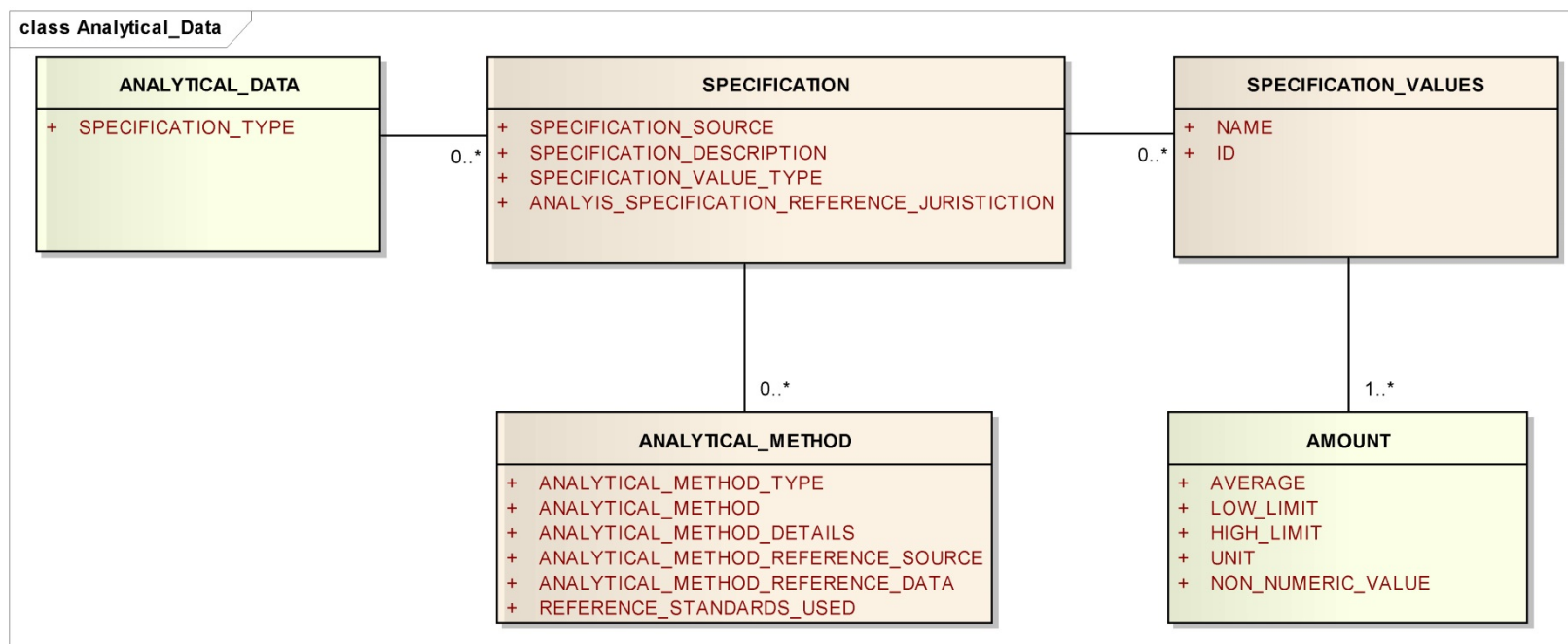


Figure 19 Information model for the analytical data class

These are Data that are in our:

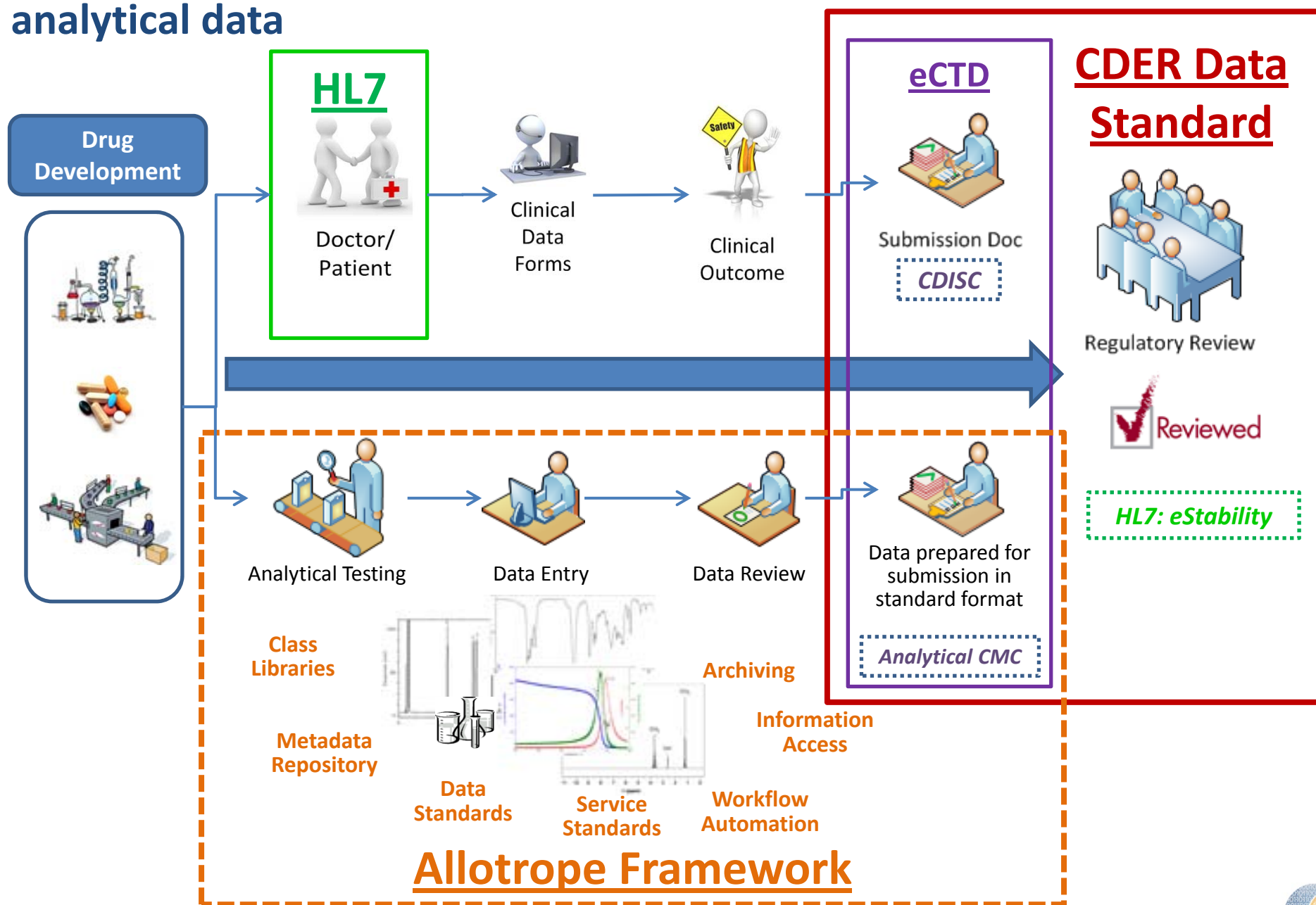
LIMS (Laboratory Information Management System) and

ELN (electronic Lab Notebook) and CDS (Chromatographic Data systems) today

Raw Data from Instruments in many different mainly proprietary Data formats

Processed Data (results) with non-harmonized meta-data

Allotrope Framework addresses the gap in standards for CMC analytical data

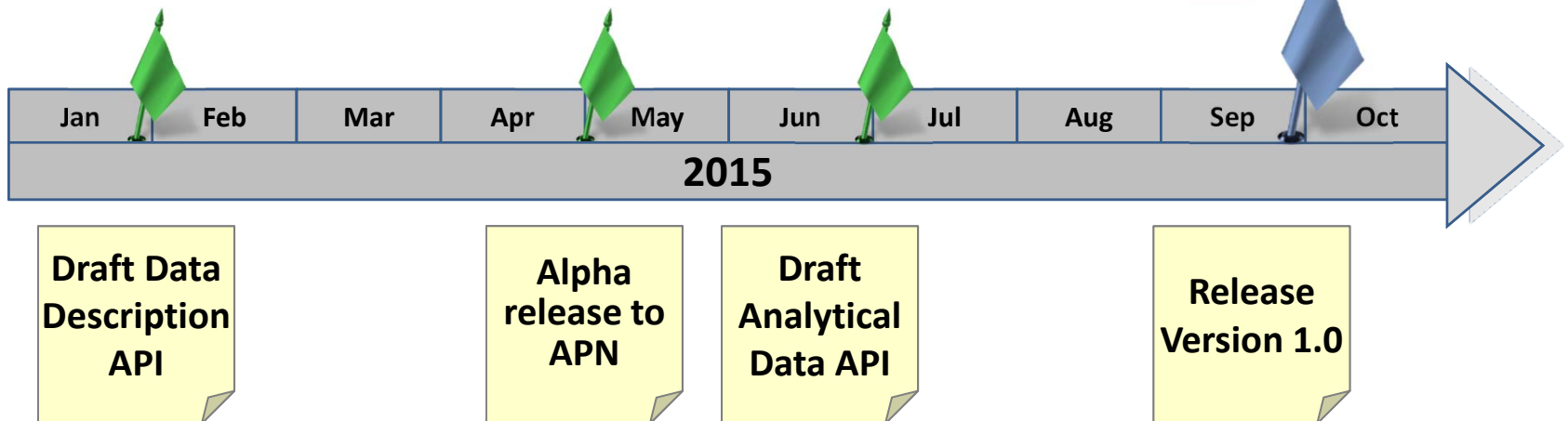


September, 2015 Release: Version 1.0

| Draft ADF Format & API | | ADF Format & API Version 1.0 |
|------------------------|------------------|----------------------------------|
| Data Description API | | Data Description API Version 1.0 |
| | Data Cube API | Data Cube API Version 1.0 |
| | Data Package API | Data Package API Version 1.0 |

| | |
|---------------------------|---------------------------------|
| Draft Analytical Data API | Analytical Data API Version 1.0 |
|---------------------------|---------------------------------|

| Draft Taxonomies | | Taxonomies Version 1.0 |
|-----------------------|---------|------------------------|
| Equipment | Process | Taxonomies Version 1.0 |
| Material | Result | |
| Analytical Techniques | | |



| | | | | | | | | | | |
|-------------------------|--------------|--|---|--|--|--|--|--|--|--|
| ISO IDMP Implementation | CDER CBER | Implement International Organization for Standardization (ISO) Identification of Medicinal Products (IDMP) standards with reliable and robust repositories and processes to support efficient, consistent, and timely decision making in the regulation of medicinal product throughout the product development lifecycle. | <p>December 2015:</p> <p>ISO 11238 pilot was completed based on the Global Substance Registration System (GSRS) in April 2015 with positive outcome. Project is moving to establish GSRS in production environment and Phase 1 to be completed FY2016 Q1.</p> | | | | | | | |
|-------------------------|--------------|--|---|--|--|--|--|--|--|--|

