



**3DEXPERIENCE®**

# Simplifying Instrument Data Integration

Pathways to Digital Success in the Lab

Gene Tetreault

# Topics

## Where we are today

A Simplified Reference Data Configuration for Equipment Measurements

An Updated approach for Reading and Writing ADF

# What's Today's Starting Point?

Asset #	Name	Manufacturer	Type	Model	Status	Next Calibration Date	Maximo Level	Site	Department	Bldg	Room
1 913787	DSD ANALYZER "SIMBA"	NOVA BIOMEDICAL CORP	NOVA BIOMEDICAL C...	BIOPROFILE CDV	ACTIVE		LEVEL_3	TO	Amgen	TO-18S0148	TO-18S0148-TO18S61535A
▶ 543960	FUNCTIONAL BIOCHARACT...	BECKMAN	BECKMAN VI-CELL XR	VI-CELL XR	ACTIVE		LEVEL_3	TO	Amgen	TO-18S0235	TO-18S0235-TO18S53640C

Background | Materials | Cell Culture Splits | Seed Train Passing Performance

Material Type	Name	Product Name	Material ID	Lot Number	Container ID	Protein Alias	Protein Name	Sequence Set ID	Volume	Vessel ID	Culture Alias	Status	Parent Culture ID	VCD (cells/mL)
▶ 1 Cell Culture	Desired Culture	AMG 181	MH007939	MH007939-001	S7V	Anti-target mAb	a181-001	Seq123	69 mL	SF 3001	Active Culture	Disposed	13473	
2 Media	IMX5.0-009			1010101										
3 Cell Culture	Desired Culture	AMG 181	MH007939	MH007939-001	S7V	Anti-target mAb	a181-001	Seq123	69 mL	SF1	Thaw	Disposed	15311	
4 Cell Culture	Desired Culture	AMG 181	MH007939	MH007939-001	S7V	Anti-target mAb	a181-001	Seq123	1.5 mL	Passage 11-A1	p1	Disposed	5755	
5 Cell Culture	Desired Culture	AMG 181	MH007939	MH007939-001	S7V	Anti-target mAb	a181-001	Seq123	1.5 mL	Passage 11-A2	p1	Disposed	5755	

Material - Sample Genealogy

Instrument - Analysis Linkage

Sample Result Summary

Name	Lot #	Sample ID	Sample Name	Analysis	Result Field	Result (numeric)	Res
▶ 89 Desired Culture	MH007939-001	S1Q9	p2_Day 3	Cell Counter	Viable Cell Density	7.137492520904...	/ml (
90 Desired Culture	MH007939-001	S1Q9	p2_Day 3	Cell Counter	Viability	97.71820068359...	
91 Desired Culture	MH007939-001	S1Q9	p2_Day 3	Cell Counter	Live Diameter		
92 Desired Culture	MH007939-001	S1Q9	p2_Day 3	Cell Counter	Total Cell Diameter	16.84981045447...	
93 Desired Culture	MH007939-001	S1QA	p2_Day 3	Cell Counter	Viable Cell Density	2.9979716378...	
94 Desired Culture	MH007939-001	S1QA	p2_Day 3	Cell Counter	Viability	90.7886428833...	
95 Desired Culture	MH007939-001	S1QA	p2_Day 3	Cell Counter	Live Diameter		
96 Desired Culture	MH007939-001	S1QA	p2_Day 3	Cell Counter	Total Cell Diameter	18.25190435180...	

Sample - Result Direct Mapping

Instrument Results View

Sample ID	Sample Name	Result Field	Instrument Asset Number	Result (numeric)
▶ 2 S1Q9	Desired Culture	instrument sample id	543960	1
3 S1Q9	Desired Culture	dilution factor	543960	2
▶ 4 S1Q9	Desired Culture	sample analysis time	543960	2
5 S1Q9	Desired Culture	viability	543960	97.718200...
6 S1Q9	Desired Culture	total cell density	543960	7.3041566...
7 S1Q9	Desired Culture	viable cell density	543960	7.1374925...
▶ 8 S1Q9	Desired Culture	total cell diameter	543960	16.849810...

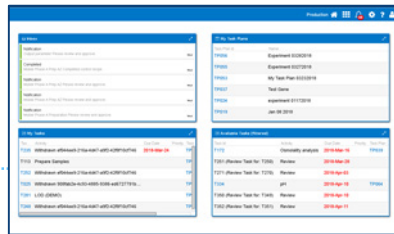
Experiment Sections

# BIOVIA ONE Lab

## Task Planner & Samples

### Lab Planning/Management

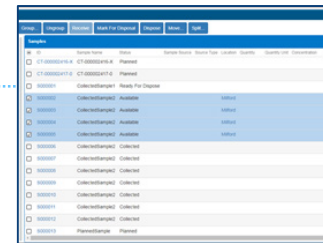
- Design and Track Studies
- Request Work
- Schedule, Review
- Confirm Requests



### CISPro& Equipment

### Resources

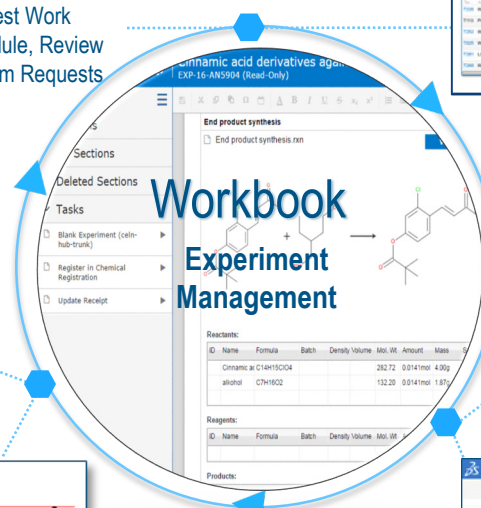
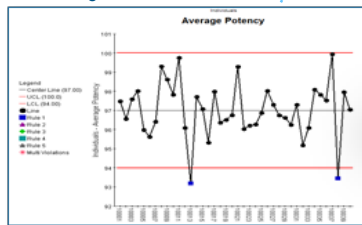
- Materials, Samples
- Personnel, Equipment



### Insight & Pipeline Pilot

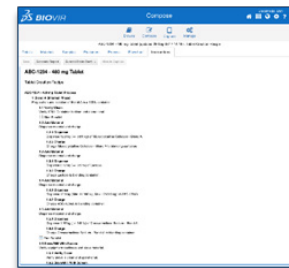
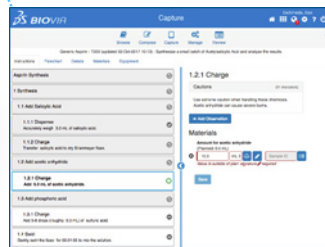
### Reporting

- Process Monitoring
- Compile, Interpret Results
- Generate Reports
- Insights



### Capture

- Prepare Tests
- Perform Tests

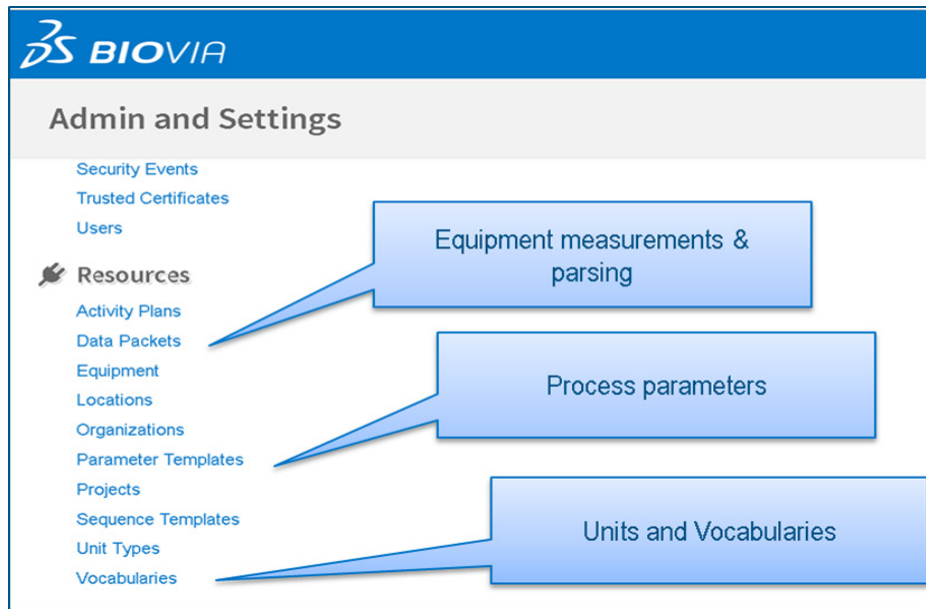
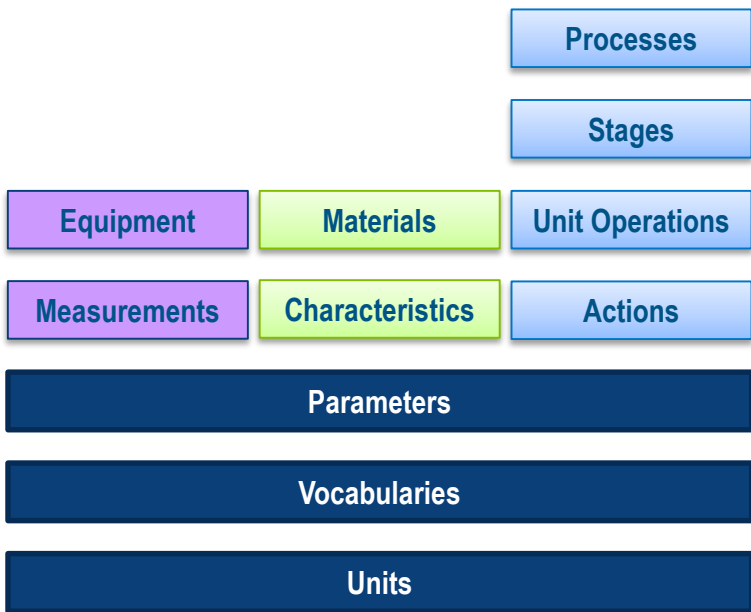


### Compose Recipes/Methods

- Develop procedures
- Adapt procedures
- Manage procedures

# Reference Data & Ontologies

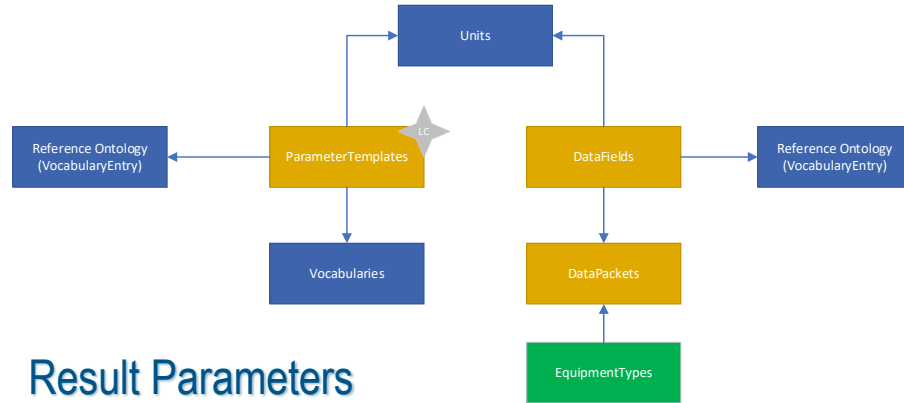
Common master data management & reference ontologies



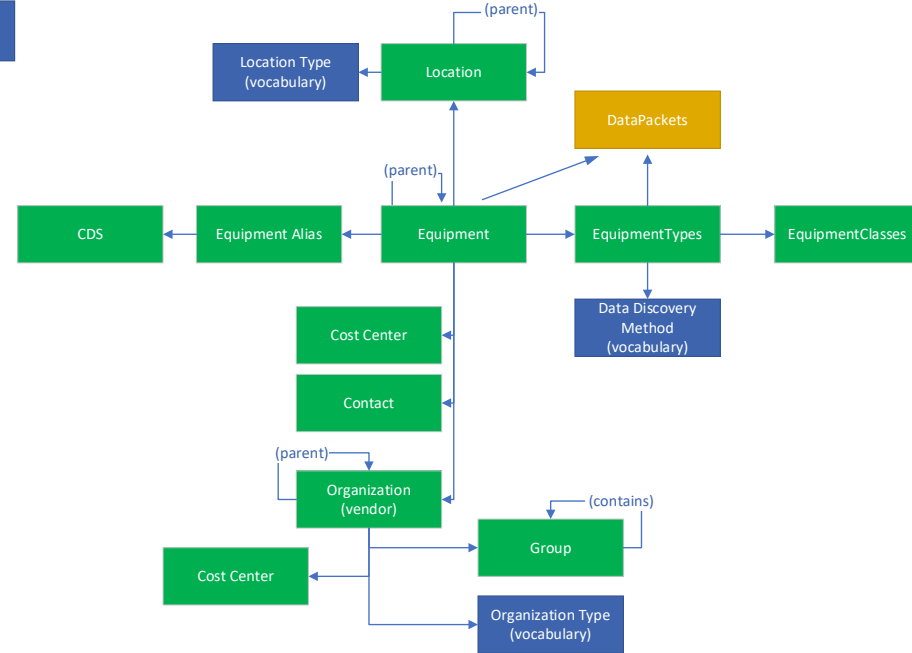
# Reference Data Schematic



## Units and Vocabularies



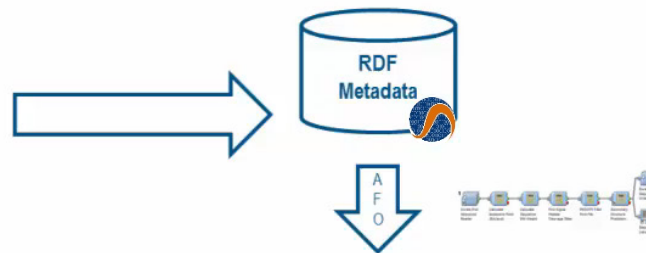
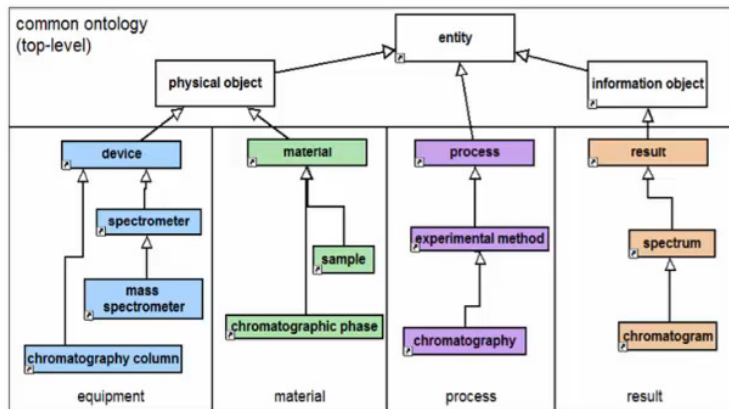
## Result Parameters



## Equipment Measurements

# Taxonomy and Ontology Synchronization

- BIOVIA ONE Lab centrally manages the ontologies
- Sync services to maintain applications usage of the vocabularies



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[http://bcn2015.europeanbioanalysisforum.eu/wp-content/uploads/2015/12/os-2015-D2A1\\_3-Gerhard-Noelken.pdf](http://bcn2015.europeanbioanalysisforum.eu/wp-content/uploads/2015/12/os-2015-D2A1_3-Gerhard-Noelken.pdf)

**BIOVIA ONE Lab**

**ASSAULT  
ISTEMES**

# Result Groups (AFR\_) as Vocabularies

## Vocabularies

Admin and Settings > Vocabularies

History

You can define lists of terms that are available to applications that use BIOVIA Foundation. These lists are maintained in the vocabularies section.

Filter Vocabularies		
	Name	Description
<input type="checkbox"/>	Allotrope Result Group: chromatography result (AFR_0000523)	Result of a chromatography run. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: data point (AFR_0000196)	A data point is the single measurement result of a series. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)	A differential scanning calorimetry peak is a peak of a differential scanning calorimetry curve. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: differential scanning calorimetry result (AFR_0000682)	The differential scanning calorimetry result is the result of a DSC measurement. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: dynamic vapor sorption measurement result (AFR_0000...	The result of a dynamic vapor sorption measurement. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: flow cytometry measurement result (AFR_0000678)	The result of a flow cytometry measurement. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: gas chromatography result (AFR_0000314)	Result of a gas chromatography run. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: isotope cluster (AFR_0000570)	Group of peaks representing ions of the same elemental composition, but different isotopic compositions. IUPAC MS REC
<input type="checkbox"/>	Allotrope Result Group: list (AFR_0000620)	A list is an ordered collection of items. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: mass spectrometry peak (AFR_0000077)	Localized region of relatively intense detector response in a mass spectrum when ions of a specified m/z are detected. If resolving power is insufficient two or more components of similar m/z may contribute to one unresolved mass peak. IUPAC...
<input type="checkbox"/>	Allotrope Result Group: mass spectrum (AFR_0000439)	Plot of the relative abundances of ions forming a beam or other collection as a function of their m/z values. IUPAC MS REC
<input type="checkbox"/>	Allotrope Result Group: Mathieu stability diagram (AFR_0000453)	Graphical representation expressed in terms of dimensionless reduced coordinates that describes the stability or instability of charged particle motion in a transmission quadrupole mass spectrometer or Paul ion trap, based on an appropriate fo...
<input type="checkbox"/>	Allotrope Result Group: nuclear magnetic resonance peak (AFR_0000451)	Peak resulting from a NMR acquisition. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: nuclear magnetic resonance spectrum (AFR_0000263)	Any spectrum that shows the response of spin-active nuclei to radio frequency radiation in an applied magnetic field. CHMO
<input type="checkbox"/>	Allotrope Result Group: numeric item (AFR_0000622)	An item in a series containing a numeric value. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: numeric series (AFR_0000623)	A numeric series is a series of numeric values. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: particle sizing result (AFR_0000745)	A particle sizing result is the outcome of a process of particle sizing. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: peak (AFR_0000413)	A peak describes a part of a spectrum/chromatogram/plot at a definite range of the experimental parameter (independent variable) of the spectrum. This definition includes a single data point of the spectrum. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: peak list (AFR_0000432)	Collection of peaks or peak groups for a specific purpose OSTHUS
<input type="checkbox"/>	Allotrope Result Group: pH measurement result (AFR_0000025)	A pH measurement result is the outcome of a the process of pH measurement. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: recorded image (AFR_0000391)	A data set consisting of a 2D representation obtained from a sample. CHMO
<input type="checkbox"/>	Allotrope Result Group: result (AFR_0000207)	The final outcome reported for a measured or computed quantity, after performing a measuring procedure including all sub procedures and evaluations. IUPAC Orange, OSTHUS
<input type="checkbox"/>	Allotrope Result Group: series (AFR_0000231)	A series is a list (ordered collection) of literal values. OSTHUS
<input type="checkbox"/>	Allotrope Result Group: spectrum (AFR_0000068)	A plot of a measured quantity against some experimental parameter. CHMO
<input type="checkbox"/>	Allotrope Result Group: temperature measurement result (AFR_0000120)	Result of a temperature measurement. Celsius temperature. Thermodynamic temperature minus 273.15 K, invariably expressed in the SI unit degree Celsius (°C) which is equal to the kelvin. Sometimes the misnomer centigrade temperature is u...



# Measurements (AFX\_) as Vocabulary Entries



## Vocabularies

Admin and Settings > Vocabularies

History

You can define lists of terms that are available to applications that use BIOVIA Foundation. These lists are maintained in the vocabularies section.

+ - ↺ ↻		peak		
	Name	Description		Vocabulary
<input type="checkbox"/>	peak area	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak height	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak maximum	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak onset	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak start	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak stop	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak temperature	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak width	The peak width is the width of a peak determined at the baseline level. The peak tangents are drawn from the turning points of the leading and trailing edges. Then the points of intersect...		Allotrope Result Group: differential scanning calorimetry peak (AFR_0000647)
<input type="checkbox"/>	peak width resolution	Ratio of the maximum ion current recorded at a specified m/z value to the maximum ion current arising from the same species recorded at a neighboring m/z value. IUPAC MS REC		Allotrope Result Group: mass spectrometry peak (AFR_0000077)
<input type="checkbox"/>	base peak	Differences in overall detection sensitivities for ions of different m/z values in a mass spectrum, caused by variations in ionization efficiency, transmission efficiency through the interface ...		Allotrope Result Group: mass spectrum (AFR_0000439)
<input type="checkbox"/>	peak area	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak area corrected	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak assignment	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak asymmetry	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak asymmetry at 10%	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak height	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak position	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak position corrected	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak rank	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak width	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	relative peak area	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	relative peak area corrected	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	relative peak height	The relative retention value calculated for two adjacent peaks. By definition, the value of the separation factor is always greater than unity. IUPAC Analytic Comp		Allotrope Result Group: peak (AFR_0000413)
<input type="checkbox"/>	peak list raw scans	A list of scan numbers and or scan ranges associated with a peak list. OSTHUS		Allotrope Result Group: peak list (AFR_0000432)
<input type="checkbox"/>	peak list scans	A list of scan numbers and or scan ranges associated with a peak list. OSTHUS		Allotrope Result Group: peak list (AFR_0000432)

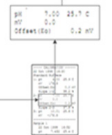
# BIOVIA Equipment

Direct Result  
Transfer

File & Database  
Transfer

Instrument  
Web Services

Data is parsed from the data stream, stored in the DB and displayed in the SOP



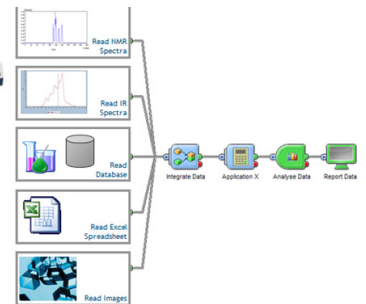
Data is collected from the instrument in RS232 and transferred to iDS using TCP/IP

Protocol Converter

DATA  
File

Database / SDMS

CDS Systems  
(Empower, Chromeleon, etc.)



1/12/2019 | ref.: 3DS\_Document\_2019

## Existing Direct Connections

Accumet 950	Julabo MD	Mettler MN5	Mettler XP204	Photovolt Aquatest 310	Sartorius LP1200	Thermo Orion 420A+
Accumet AR15	Kraemer Elektronik HC 87	Mettler PB1500-S	Mettler XP205	Radiometer DM220	Sartorius LP2200S	Thermo Orion 620A
Accumet AR20	Leica Auto ABBE	Mettler PB3001-S	Mettler XP205DR	Radiometer PHM210	Sartorius LP3200D	Thermo Orion 720A Plus
Advanced Instruments 3250	Metrohm 702GM T.rino	Mettler PB3002-S	Mettler XP28	Radiometer PHM220	Sartorius LP4200S	TSI 4043
Advanced Instruments 3300	Metrohm 712	Mettler PB3002-S	Mettler XP56	Radiometer PHM30	Sartorius LP6200P	Turner Quantech FMI 0635
Advanced Instruments 303	Metrohm 727 KF Coulometer	Mettler PB302	Mettler XS1003-S	Sartorius ACOO	Sartorius LP6200S	Vankel 25-1000
AND HR200	Metrohm 756 KF Coulometer	Mettler PB400-S	Mettler XS104	Sartorius ACOO-S	Sartorius M-power	Vankel W-200
Anton Paar DMA 4500	Metrohm 780	Mettler PB802-S	Mettler XS204	Sartorius AC210S	Sartorius M5P	Vankel W-7000
Anton Paar/Citzen DMA 5000	Metrohm 787 KF T.rino	Mettler PG100S-S	Mettler XS205	Sartorius AC211S	Sartorius MA40	WWR symphony SB70P
Beckman 360	Metrohm 795 T.rino	Mettler PG2002-S	Mettler XS205DU	Sartorius AC211D	Sartorius MA51	WWR symphony SB80PC
Beckman 45 pH Meter	Metrohm S31 Coulometric	Mettler PG3001-S	Mettler XS300	Sartorius BP211D	Sartorius MC210P	WWR symphony SP21
Brimmann 756 KF Coulometer	Metrohm 841 KF T.rinda	Mettler PG4002-S	Mettler XS802-S	Sartorius BP212S	Sartorius MC5	WWR symphony SP70P
Brinkmann Metrohm 713	Mettler AG104	Mettler PG403-S	Mettler XS802-S	Sartorius BP3100S	Sartorius MC6	WWR symphony SFC01
Brookfield DWH-LV	Mettler AG135	Mettler PG6002-S	Millipore Milliflex PLUS	Sartorius BP4100S	Sartorius ME215P	WWR symphony SFC01
Brookfield DWH-RV	Mettler AG204	Mettler PM100	Mitsubishi CA-100	Sartorius CP124S	Sartorius ME216S	WWR symphony SR40C
Buchi B-540	Mettler AG245	Mettler PM1300	Mitsubishi CA-200	Sartorius CP2201	Sartorius ME23S	Wescor E520
Corning 350	Mettler AT200	Mettler PM2500	Mitsubishi CAW-100	Sartorius CP2202S	Sartorius ME25P	YSI 3200
Corning 540	Mettler AT201	Mettler PM2500	Ohaus EP-413C	Sartorius CP2250	Sartorius ME23ES	Zeus MGS311
Cosa Instruments CAVA-100	Mettler AT260	Mettler PR2002	OHaus EP612C	Sartorius CP2P-F	Sartorius ME36S	
Dishak 11006	Mettler PR003	Mettler PR003	OHaus Explorer	Sartorius ED8202S	Sartorius ME5	
Dr. Schleuniger SM	Mettler AX105	Mettler PR013	Ohaus V12140	Sartorius Genius	Sartorius M346 6S	
Fisher Scientific AR25	Mettler AX105DR	Mettler PR8002	Orion 150	Sartorius L400-S	Sartorius RI60P	
Fiske 210	Mettler AX205	Mettler PR803	Orion 150A	Sartorius LC200S	Sartorius RC20-D	
Fluke 1523	Mettler AX26	Mettler PR803	Orion 162A	Sartorius LC200S	Sartorius RC210S	
GT D Bioscience M03	Mettler RE40	Mettler RE40	Orion 250A	Sartorius L4310S	Sartorius SE2	
Hach 2100AN	Mettler SB16001	Mettler SB16001	Orion 350	Sartorius LC12001S	Sartorius TEB12	
Hach Ultra Met One 3400	Mettler DE40	Mettler SevenCompact	Orion 370	Sartorius LC2021	Shimadzu ALW1200	
Hanson Research SR8 Plus	Mettler DL31	Mettler SevenEasy	Orion 420A	Sartorius LC220S	Solar HT1	
Holland 630	Mettler DL38	Mettler SevenMulti	Orion 720A	Sartorius LC200S	Stanford Research OptiMelt	
Jenway 3100	Mettler UM02	Mettler SR18001	Orion 920A	Sartorius LE2202S	Thermo Orion 150A+	
Jenway 3320	Mettler MP227	Mettler UM02	Orion 949	Sartorius LE2250	Thermo Orion 3 Star	
Julabo F26MV	Mettler MTS	Mettler MOXS20EDU	Perkin Elmer 341	Sartorius LE26P	Thermo Orion 370	
		Mettler XP120G-S	Photovolt Aquatest - 10	Sartorius LE4202S	Thermo Orion 4 Star	

## Existing Parsers

ABI 7900 HT	GUAVA EASTCYTE	Plate Reader (Envision)
ABI QuantStudio7	HeadSpace Analyzer FMS-1400	Plunger Inspection Device
Agilent 2100 Expert Bioanalyzer	HeadSpace Analyzer FMS-760	Polarimeter (Autopol V Plus)
Agilent Cary 60	ICDataCenter	PVM
AKTA Crossflow	ICP-MS	Raman Rigaku Firstguard
AKTA Explorer	Korsch XM12	Robotic Drop Tester
Analytical ultracentrifuge	Leak Detector 325	TA DSC
Barcode Labeling	Leak Detector 455	TecanMagellan
Biacore 3000	Leak Detector 655	Tecan-qPCR
BioRad ChemiDoc MP	Lyophilizer Lyostar II	Tenney Environmental Chamber
BioRad ChemiDoc ProteinGel	Lyophilizer Lyostar III	Tensiometer
Bruker B-ACS120 NMR	Malvern Particle Sizer (Mastersizer 2000 and 3000)	Thermogravimetric Analyzer GA Q500
Bruker Vertex 70 FTIR	Micro Flow Imaging (MFI)	Tiamo KF
CARY 50	Multi-Angle static Light Scattering	Transportation Lab Environmental Chamber
Cedex	Nano Drop 1000	Tristar Surface Area
Cedex HR	Nova Biomedical Flex	Varian Cary SoloVPE
Circular Dichroism	Nova CDV	Varian Spec 4000
Compression Tester	Optical Comparator	ViCell
Digital Coordinate Measuring Machine	Particle Counter (Single) HIAC	Viscosometer (Rheometer)
Dynamic Vapor Sorption	Particle Counter Auto	VMAX
Force T ester	Particle Vision System	X-Ray (Xpert Data Viewer)
Fortebio	Pendotech	XRPD

# Equipment Registry

- ▶ Equipment is registered against the list of equipment types and location
- ▶ Connection information is specified for each registered equipment
- ▶ Equipment can be deactivated if they are taken out of service
- ▶ Once registered, readings may be taken from the equipment
- ▶ The list allows for filtering and quick links to related items

The screenshot displays the BIOVIA Equipment Registry interface. The top navigation bar includes the BIOVIA logo, the user 'scitegicadmin', and various utility icons. The main content area is titled 'Equipment' and includes a 'History' button. Below the title, there is a brief description of the equipment management capabilities and a 'Related Items' section with links to 'Equipment Classes' and 'Equipment Types'. A table titled 'Filter Equipment' lists the following records:

	Display Name	Barcode	Nickname	Equipmen...	Serial Number	Location	Primary ...
<input type="checkbox"/>	BAL1 (PRP-BAL-1)	BAL1	PRP-BAL-1	<a href="#">Mettler AG245</a>	US9849878	<a href="#">Prep Lab</a>	
<input type="checkbox"/>	BAL2 (PRP-BAL-2)	BAL2	PRP-BAL-2	<a href="#">Mettler AG245</a>	US8978787	<a href="#">Prep Lab</a>	
<input type="checkbox"/>	BAL3 (CHM-BAL-1)	BAL3	CHM-BAL-1	<a href="#">Mettler AG245</a>	US8798798	<a href="#">Chemistry Lab</a>	
<input type="checkbox"/>	BAL4 (MCR-BAL-1)	BAL4	MCR-BAL-1	<a href="#">Mettler AG245</a>	US88789798	<a href="#">Micro Lab</a>	
<input type="checkbox"/>	BAL5 (MCR-BAL-2)	BAL5	MCR-BAL-2	<a href="#">Mettler AG245</a>	US8979087908	<a href="#">Micro Lab</a>	
<input type="checkbox"/>	BAL6 (MCR-BAL-2)	BAL6	MCR-BAL-2	<a href="#">Mettler AG245</a>	US87987987	<a href="#">Micro Lab</a>	

# Equipment Model Overview

## Equipment Class

Classifications make it easier for users to find equipment by grouping similar types of equipment together. Examples of classes include balances, pH meters, and HPLCs.

## Equipment Type

Equipment types define the specifications for a particular make and model of equipment. Where applicable, this definition can include the data that can be acquired from the equipment and the commands to use to obtain that data.

## Equipment

Instruments and equipment are registered with Foundation to make them available for use in a Foundation deployment. Registering a piece of equipment involves selecting its type and may include information about how to interact with it to retrieve readings.

# Equipment Example

## Equipment Class

Gas Chromatograph

## Equipment Type

Agilent 6890

## Equipment Instance

CHM-GC-1

**BIOVIA**

### Equipment Classes

[Admin and Settings](#) > [Equipment](#) > [Equipment Classes](#)

Classifications make it easier for users to find equipment by grouping similar classes include balances, pH meters, and HPLCs.

Equipment class is a required field when defining an equipment type.

Filter Equipment Classes

	Name
<input type="checkbox"/>	Analytical Balance
<input type="checkbox"/>	Analyzer
<input type="checkbox"/>	Balance
<input type="checkbox"/>	Bioreactor
<input type="checkbox"/>	Calorimeter
<input type="checkbox"/>	Conductivity Meter
<input type="checkbox"/>	Conductometer
<input type="checkbox"/>	Coulometer
<input type="checkbox"/>	Densitometer
<input type="checkbox"/>	Density Meter
<input type="checkbox"/>	Dissolution Bath
<input type="checkbox"/>	DissoPrep
<input type="checkbox"/>	Flow Meter
<input type="checkbox"/>	Fluorometer
<input type="checkbox"/>	Gas Chromatograph

**DS SYSTEMES**

# Equipment Example

Equipment Class

Gas Chromatograph

Equipment Type

Agilent 6890

Equipment Instance

CHM-GC-1

The screenshot shows the BIOVIA software interface. At the top is the BIOVIA logo. Below it, the title 'Agilent 6890 Gas Chromatograph' is displayed, followed by the breadcrumb path: 'Admin and Settings > Equipment > Equipment Types > Agilent 6890 Gas Chromatograph'. On the left side, there is a navigation menu with three items: 'General' (selected), 'Inventory', and 'Commands'. The main content area is divided into two columns. The left column contains 'Name: Agilent 6890 Gas Chromatograph' and 'Data Packet: GC Data'. The right column contains 'Manufacturer: Agilent' and 'Equipment Class: Gas Chromatograph'.

Section	Field	Value
General	Name:	Agilent 6890 Gas Chromatograph
	Data Packet:	GC Data
Inventory	Manufacturer:	Agilent
	Equipment Class:	Gas Chromatograph

# Equipment Example

Equipment Class

Gas Chromatograph

Equipment Type

Agilent 6890

Equipment Instance

CHM-GC-1

The screenshot shows the BIOVIA software interface. At the top is a blue header with the BIOVIA logo. Below the header, the title "Agilent 6890 Gas Chromatograph" is displayed in a large, bold font. Underneath the title is a breadcrumb trail: "Admin and Settings > Equipment > Equipment Types > Agilent 6890 Gas Chromatograph". The main content area is divided into two columns. The left column contains a vertical menu with three items: "General", "Inventory", and "Commands". The "Inventory" item is currently selected, indicated by a blue vertical bar to its left. The right column is titled "Equipment:" and contains the following information: "GC-1 (CHM-GC-1)", "Milford", and "Agilent 6890 Gas Chromatograph (Gas Chromatograph)".

# Equipment Data Acquisition

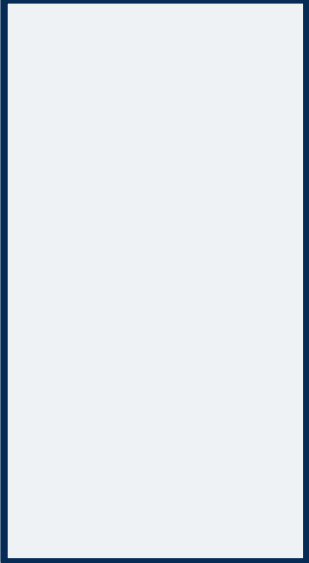
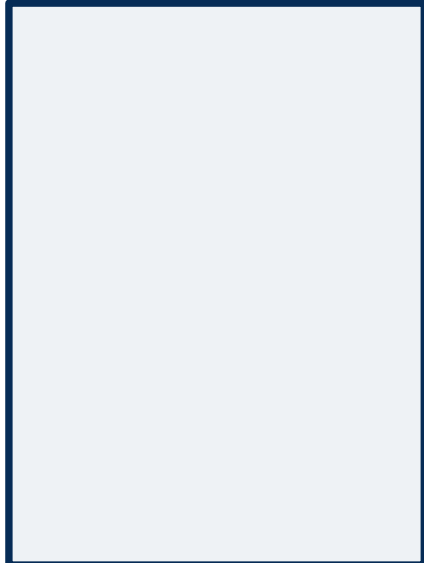
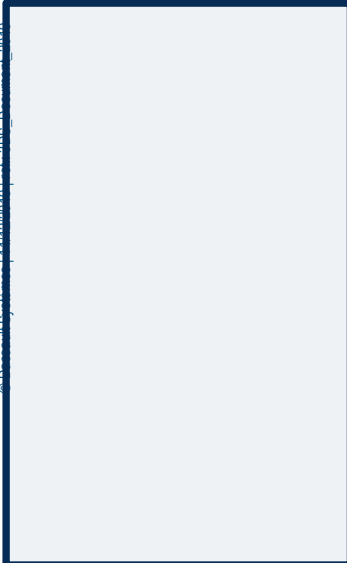
Data Packet  
(File)

Mappings

Parsed Name  
(Parameter)

Measurement  
Ontology

Triple Store



Equipment  
Vendor

Equipment Data Format

Measurement  
Ontology

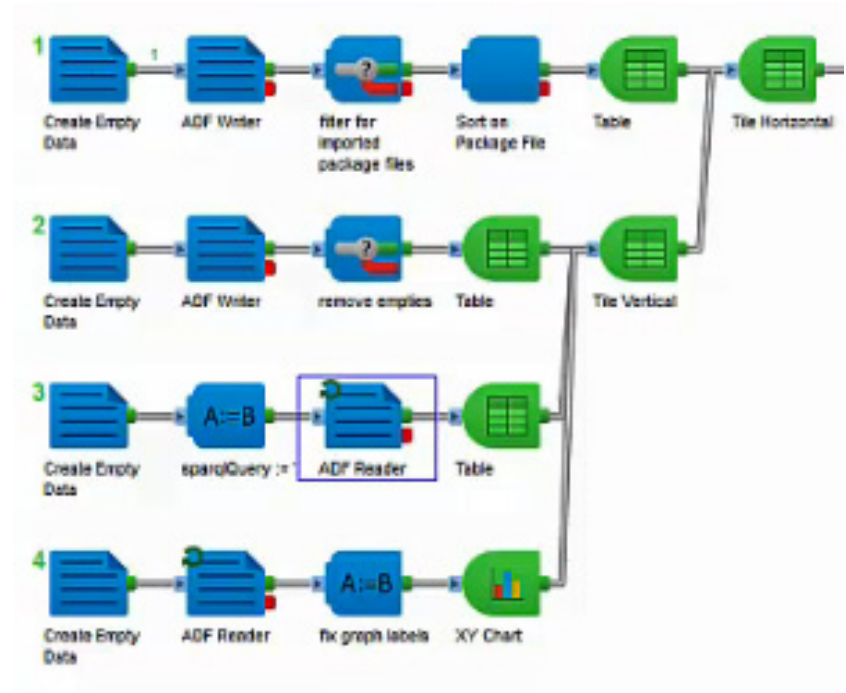
ONE Lab





# BIOVIA –Allotrope ADF Support in ONELab

- ▶ Pipeline Pilot Components for Allotrope ADF manipulation
  - ▷ Read and write ADF version 1.3 since Fall 2017
- ▶ Designed to support custom ADF read or write capabilities



# ADF Parsing (Click to play video)

ADF File Reader and Writer Example.

Protocol illustrates reader and writer components.

In line 1 the writer creates an ADF file and writes metadata and files into it. The files are specified as contained in a directory and the metadata as a TTL file.

In line 2 data cubes are imported and appended to the ADF file. The data cubes are defined by a metadata file and a directory location containing the data cube data files.

In line 3 a SPARQL query is passed to the ADF Reader to cube retrieve information about the imported data cube data.

In line 4 the data for the first data cube is read using the ADF Reader and graphed.

# Going Forward: Simplified Equipment Configuration

- ▶ Why is this needed?
  - ▷ Configuring readings for equipment can be complex
  - ▷ Requires creation and configuration of Parameter Templates, Data Packets, Parsers, Data Fields for every Equipment Type
  - ▷ Need simpler way to add equipment from any known class

The screenshot displays two configuration panels. The top panel is for 'Weight', with a breadcrumb trail 'Admin and Settings > Data Packets > Data Fields > Weight'. It includes a 'Data Packet:' field with the value 'Balance', and checkboxes for 'Is Sample Id:' and 'Is Group By:'. The 'Display Order:' field is a dropdown menu. The bottom panel is for 'BGA Data', with a breadcrumb trail 'Admin and Settings > Data Packets > BGA Data'. It features a 'Description:' field containing 'This is a data packet for BGA', and a 'Data Fields:' section with a list of fields: mPO2, mpH, mPCO2, mBP, and iFIO2. Below the list is a pagination control showing 'Page 1 of 2' and navigation arrows. At the bottom, it shows 'Last Updated: 2019-Aug-14 14:14:57' and a 'Data:' label.



# Going Forward: Simplified Equipment Configuration

- ▶ Re-align the Equipment Class-Equipment/Data Field relationships
- ▶ Support association of the Allotrope Result Group ontology (AFR) with the Equipment Class (AFX- AFO)
  - ▷ Provides higher level of re-use of configuration defined by Allotrope and simplification of the definition of Measurement results
  - ▷ Allows pre-definition of readings for Equipment Classes, requiring an Admin only to map vendor readings to AFO readings
  - ▷ When parsing vendor-supplied ADF files, readings would not require user definition other than mapping to parameter and setting context

# BIOVIA – Allotrope Ontology Support

- ▶ Currently ULM supports the association of results with the Allotrope Measurement ontology (AFX) through Data Field reference vocabularies and Parameter Template external Ids
- ▶ Roadmap for ONE Lab ULM Equipment
  - ▷ Re-align the Equipment Class-Equipment/Data Field relationships to support association of the Allotrope Result Group ontology (AFR) with the Equipment Class
  - ▷ Provides higher level of re-use of configuration defined by Allotrope and simplification of the definition of Measurement results

# Positive Allotrope Direction – Simplification

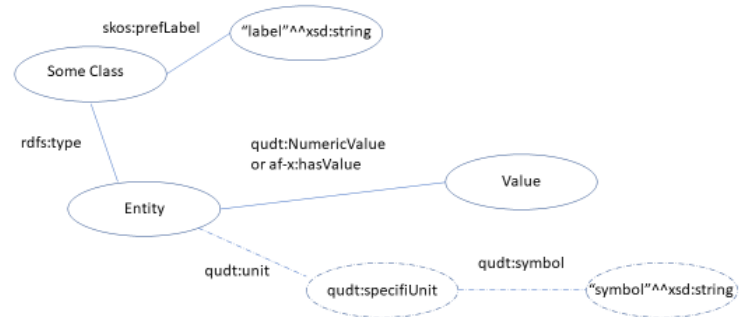
- ▶ Leaf Node Model
- ▶ Fundamentally more practical for broader array of implementations
- ▶ More facile engineering with broader array of tools
- ▶ Much more generalizable to the Equipment Class Level
- ▶ More comprehensible to a broader array of users

Example: Cell Counter Data

This is essentially name:value pairs

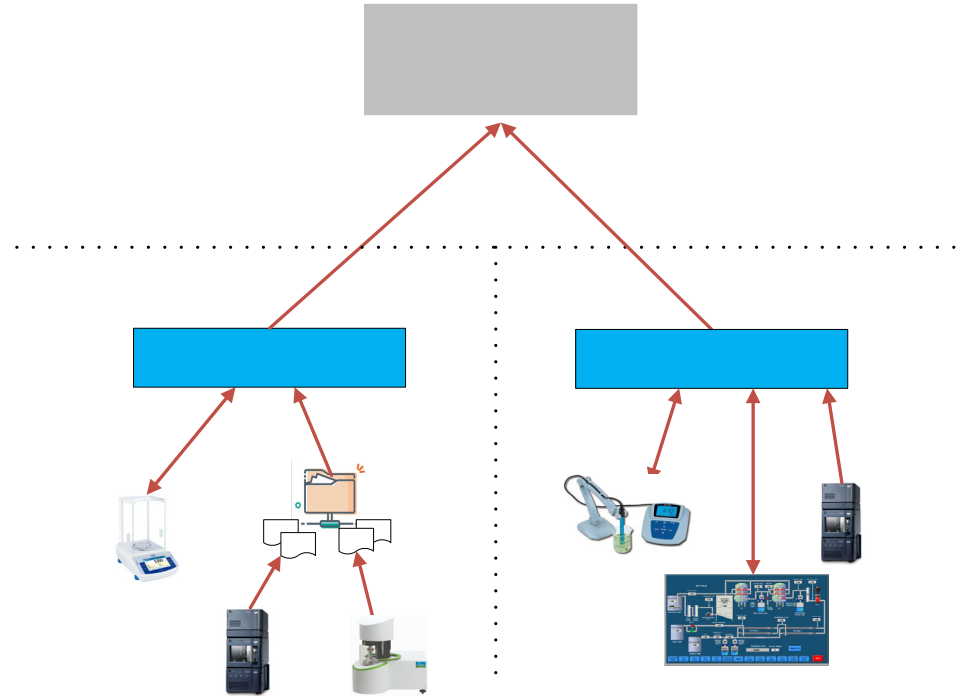
Vi-CELL XR 2.04							
Beckman Coulter, Inc.							
Sample ID	Dilution factor	Sample date	Viability (%)	Total cells /ml (x10 <sup>6</sup> )	Viable cells /ml (x10 <sup>6</sup> )	Avg. diam. (microns)	
water check 03 Aug 2017	1.0	3 Aug 2017 10:55:55 AM	30.8	0.013	0.004	9.39	
water check 080317 2	1.0	3 Aug 2017 11:09:19 AM	33.3	0.003	0.001	6.05	
wil2s 080317 pre ff	1.0	3 Aug 2017 11:47:19 AM	97.4	1.61	1.57	13.59	
wil2s 080317 post ff	1.0	3 Aug 2017 11:56:43 AM	97.0	0.71	0.68	13.68	
U20S IL23R P2 CF 03Aug17	1.0	3 Aug 2017 1:19:02 PM	94.4	0.16	0.15	15.37	
U20S IL12R P3 CF 03Aug17	1.0	3 Aug 2017 1:31:15 PM	85.3	0.15	0.13	16.21	
NR92 P30 CF 04Aug17	1.0	4 Aug 2017 1:11:06 PM	94.2	2.85	2.68	15.57	

And the associated triple pattern



# ONE Lab Integration Appliance

- ▶ The Integration Appliance provides a bridge between any number of site laboratories and the central installation of ONE Lab
- ▶ The Integration Appliance will support
  - ▷ Discovering data files from laboratory instruments for results parsing (e.g. TGA)
  - ▷ Reading data from network-connected port devices (e.g. balances)
  - ▷ Local label printing
  - ▷ Reading data from OPC-enabled equipment (e.g. protein purification equipment)
  - ▷ Providing data to OPC-enabled software (e.g. MES)
  - ▷ **Writing Allotrope ADF files from ONE Lab-connected equipment**
  - ▷ **Reading Allotrope ADF files from Allotrope-enabled instruments**



# Conclusion

- ▶ The new Equipment Class mode in ONE Lab will permit the **alignment of Equipment Class and Allotrope Result (AFR) ontology**.
- ▶ Configure an Equipment Class once, align it with the AFR and for each new manufacturer, you only need to alias the field names to the Equipment Class/AFR. Much **easier to maintain and scale Equipment** definitions.
- ▶ The **leaf-node approach** lets us create a more general solution to support **ADF** read/write enhancing **scalability**.
- ▶ The **Pipeline Pilot** components available for use in any application that require more **specific/deeper use of the ADF** file (e.g. data cubes writing or reading).



