

# **Native AnIML LCMS Processing and Viewing**

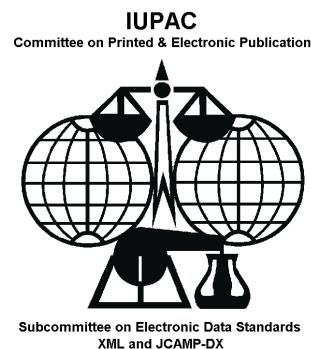
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ASTM E13.15 Subcommittee (Analytical Data)  
IUPAC Subcommittee on Electronic Data Standards (SEDS)



# AnIML – a New Home for Old Standards



JCAMP-DX

GAML





# AnIML Hybrids (LC-UV-ELS-MS)

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## Sample Alteration

**Separations**  
*chromatography*

**Chemistry**  
post-column  
addition  
reaction kinetics

→ *new samples*

4

## Sample Measurement

**Detection**  
spectroscopy  
UV, IR, MS, NMR  
point-detection  
pH, density

→ *data*

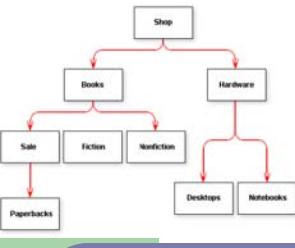
4

## Data Processing

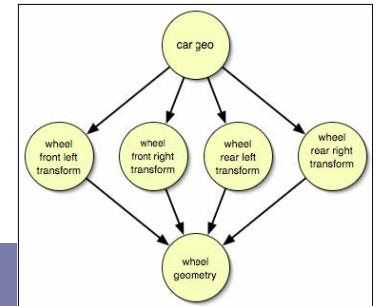
**Single Trace**  
noise filtering  
smoothing  
peak finding

**Multiple Runs**  
quantitation

→ *new data*



# Hierarchical vs. Referenced Relationships



- Hierarchical (one parent only)

xml

```

<ExperimentStep name="Spectrum">
    <Infrastructure>
        <ParentDataPointReference seriesID="ix1" startIndex="278"/>
    </Infrastructure>
    <Result name="Spectrum">
        <SeriesSet name="Spectrum" length="138">
            <Series name="Intensity" dependency="dependent" seriesID='
                <EncodedValueSet>AEExVRwDQfkcaKpRHAkanRwAqqucANsxHACTbI
            </Series>
        </SeriesSet>
    </Result>
</ExperimentStep>
  
```

- key\keyref (multiple parents possible)

xsd  
(schema)

```

<xsd:key name="sampleID">
    <xsd:selector xpath="SampleSet/Sample"/>
    <xsd:field xpath="@sampleID"/>
</xsd:key>
<xsd:keyref name="sampleIDUsage" refer="sampleID">
    <xsd:annotation>
        <xsd:documentation>An ExperimentStep may only r
    </xsd:annotation>
    <xsd:selector xpath=".//ExperimentStep/SampleRefere
    <xsd:field xpath="@sampleID"/>
</xsd:keyref>
  
```



# What Do XML Schema Do?

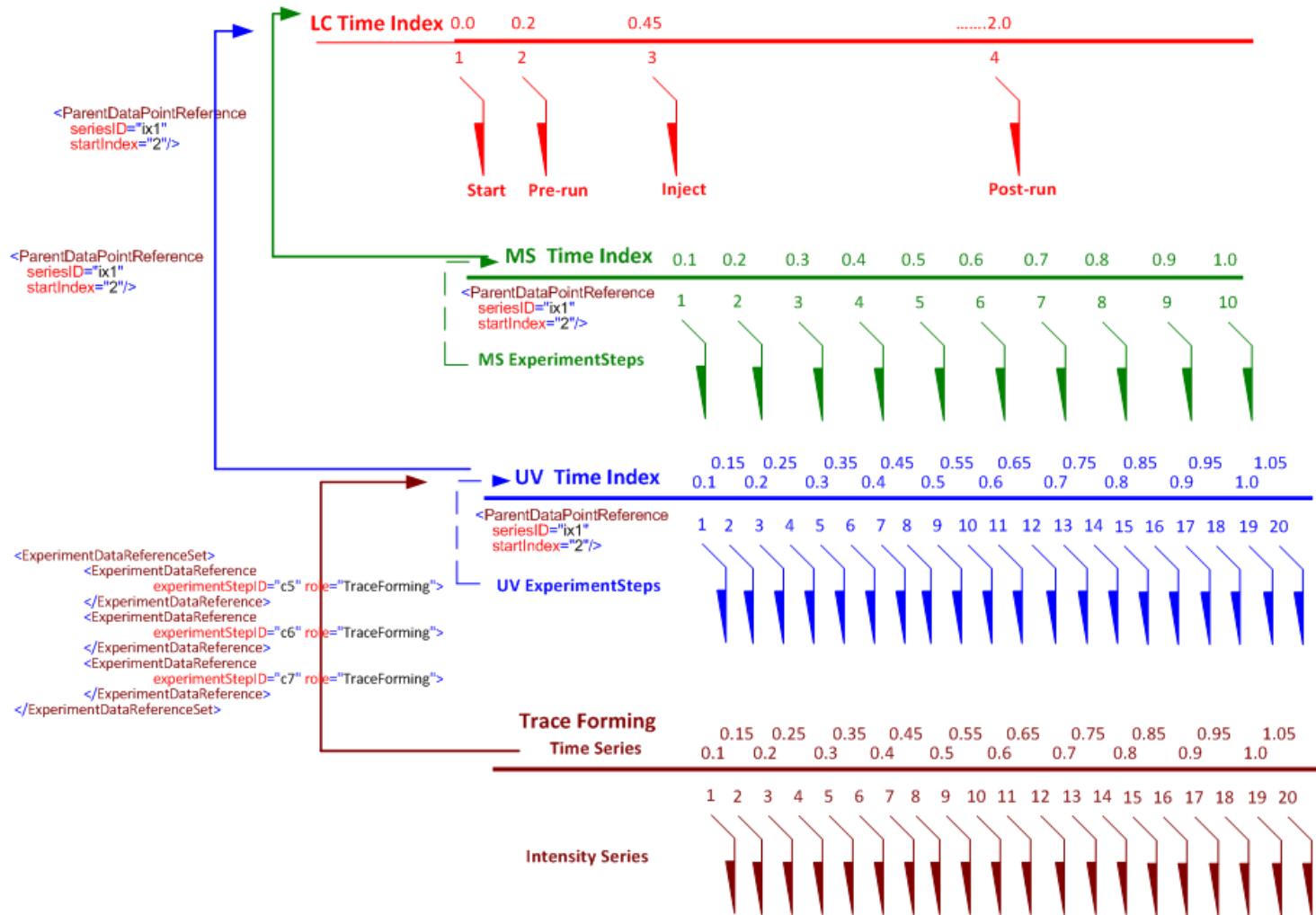
- Controls syntax -- *structure, relationships, constraints, and terminology* for XML files
- Used to validate XML documents
- The complex rules governing schema are difficult to understand.
- Hierarchical XML limits ability to model data relationships; a child can have only one parent in a hierarchical tree. This is resolved in schema by introduction of **key, keyref** (foreign key constraints).



# >90% of AnIML Data Looks Like This



# Time Synchronization Issues





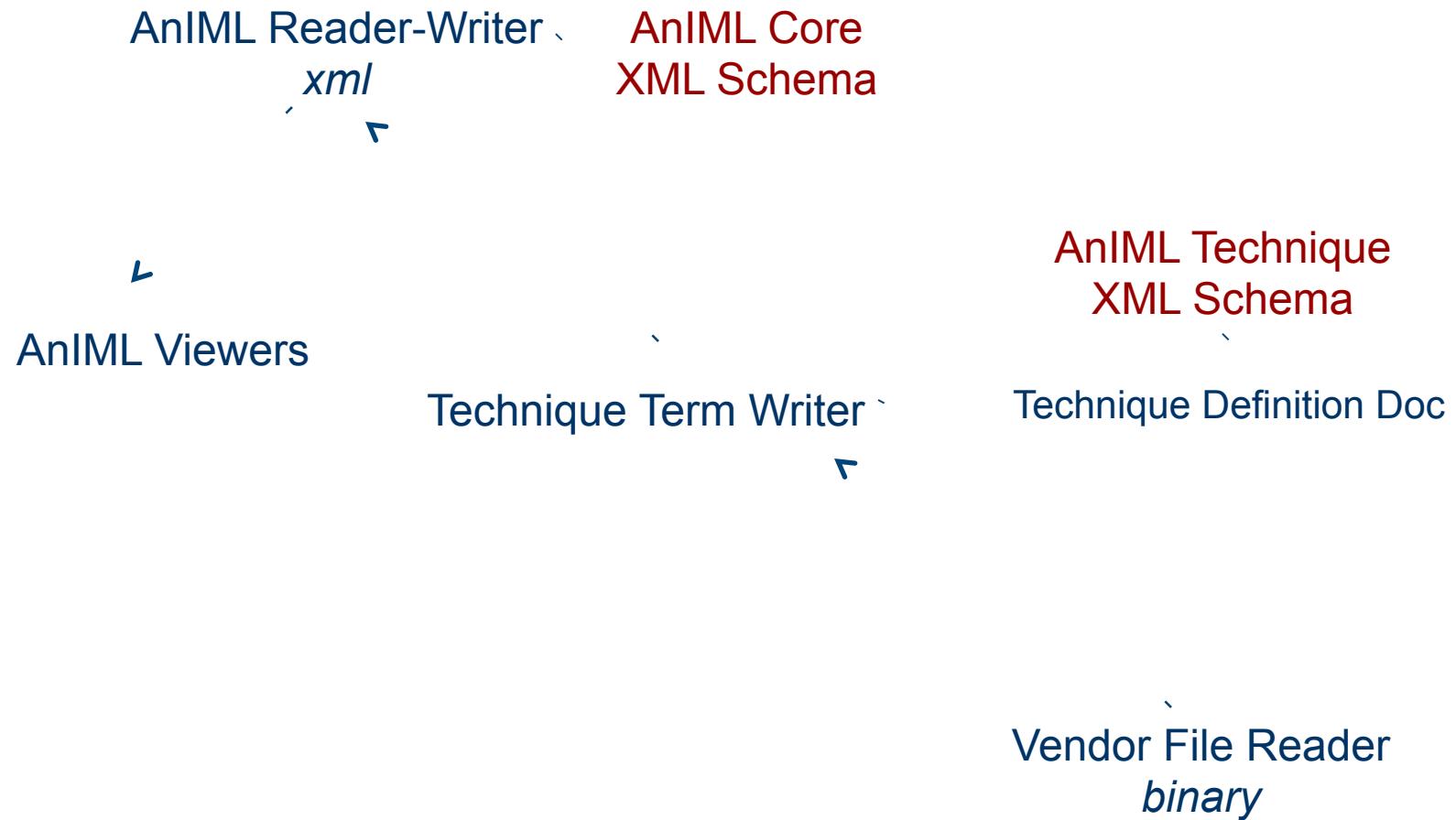
# Handling Multiple Dimensions

*One of the hardest areas to get right in AnIML*

- Are LC-UV or LC-MS truly 2D data sets?
- or are they an array of spectra against time?
- Are chromatograms the result of processing?
- 2D NMR is an array of spectra against parameter changes (T1)



# Implementing AnIML





# IRSample.AnIML

- Simple IR spectrum with metadata
- (Demo)
  - XML parsing (collaboration with Scimatic)
  - Intelligent Viewer (CANDI)



## LC-UV-GStd.AnIML

- Complete Array of UV Spectra from LC run
  - Uses Templates
  - Multiple Techniques
    - Chromatography
    - UV-Vis
    - Peak Finding
- (Demo)
  - XML parsing (collaboration with Scimatic)
  - Intelligent Viewer (CANDI)



# LC-UV-MS-Report.AnIML

- Summed & Subtracted UV and MS Spectra
  - Uses Templates
  - Multiple Techniques
    - Chromatography
    - UV-Vis
    - Peak Finding
    - Summation-Subtraction
- (Demo)
  - XML parsing (collaboration with Scimatic)
  - Intelligent Viewer (CANDI)



# AnIML = Working Together





# Acknowledgements

- **Burkhard Schaefer**, BSSN Software
- **Maren Fiege**, Waters
- **Dale O' Neill**, Agilent
- **Jamie McQuay**, Scimatic Software
- **Stuart Chalk**, UNF
- Tony Davies
- Bill Browett, GSK
- Lyle Burton, Sciex
- David Martinsen, ACS
- James Duckworth, Thermo
- Randall Julian
- Mark Mullins, Agilent
- Robert J. Lancashire, UWIMONA
- Richard Larsen

## NIST

- **Gary Kramer**
- **Dennis Backhaus**
- **Peter Linstrom**
- Alexander Roth
- Anh Dao Nguyen
- Dominik Poetz
- Frank Masur
- Kordian Placzek
- Patrick Gleichmann
- Ronny Jopp