

A Protégé Ontology as The Core Component of a BioSense Message Analysis Framework

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Outline

- BioSense description
- Describe the current environment
- Describe the ontology
- Describe the ontology framework
- Describe the analysis workbench
- Future directions
- Questions



BioSense Description

What is BioSense?

- Real-time and near real-time national public health message analysis framework
- Consists of
 - Message acquisition and translation interfaces
 - Secure message transmission network
 - Message classification components
 - Data storage and query components
 - Data analysis component
 - CDC Monitors
 - Local data visualization and distribution

BioSense Functions

Confirm or refute existence of an event

- ✓ Environmental signal
- ✓ Suspect illness
- ✓ Intelligence warning
- ✓ Known outbreak/public health event

Monitor ongoing event and effectiveness of response

- ✓ Ascertain size of event
- ✓ Ascertain rate of spread
- ✓ Track efficacy of response efforts
- ✓ Monitor for adverse events
- ✓ Know when an event has passed

Data Sources

<i>Data Source 2006</i>	<i>Rationale</i>
Orders & results from 3 major commercial clinical laboratories	<i>Represent 20% of all US lab testing; 60% of independent testing; critical to many PH efforts</i>
Real-time data from VA	<i>150 hospitals and ~1000 ambulatory care clinics; share data with many state and local PH communities</i>
Real-time data from DoD	<i>45 US hospitals and ~800 ambulatory; share data</i>
Poison Control Centers call data	<i>All 62 poison control centers; display and compare with other community health data</i>
Private Hospitals	<i>500 Clinical care Hospitals provide national view and local data</i>

Target Data Types

- **Foundational***: demographics, chief complaint, discharge diagnoses, disposition, hospital utilization
- **Clinical***: vitals, triage notes, working diagnosis, discharge summary
- **Laboratory**: orders, microbiology results
- **Pharmacy**: medication orders
- **Radiology**: orders, interpretation results

All structured in HL7 2.5 BioSense messages

Current Classification

- Data mapped to 11 syndrome categories
 - Botulism-like
 - Fever
 - Gastrointestinal
 - Hemorrhagic illness
 - Localized cutaneous lesion
 - Lymphadenitis
 - Neurological
 - Rash
 - Respiratory
 - Severe illness/death
 - Specific infection
- 79 sub-syndrome categories

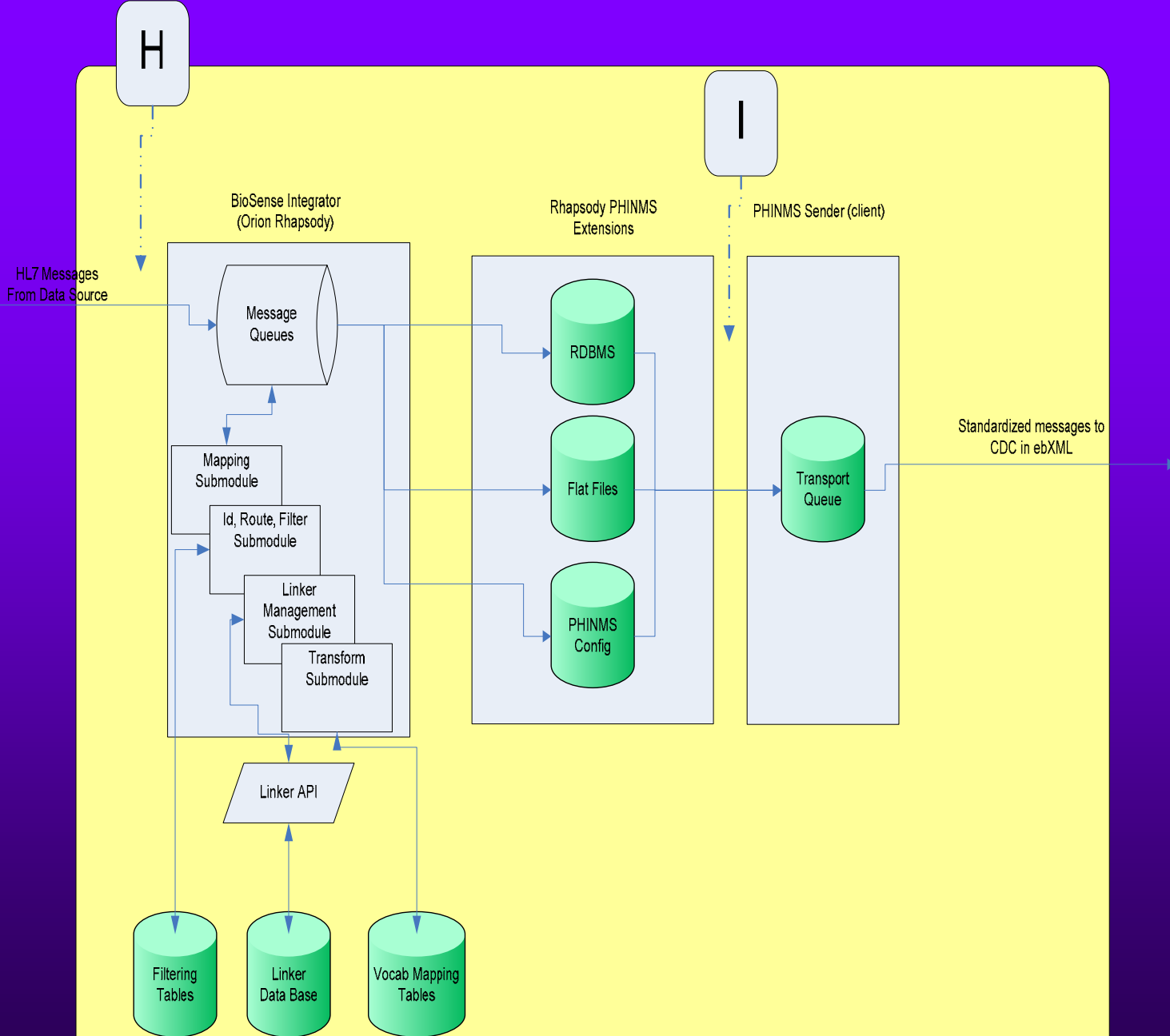
Watch what you ask for!

- BioSense message volume capacity today
 - 837 messages a second
 - >72 million messages a day
- How does an epidemiologist review that volume of data?
- How do you link messages to an individual over time to refine the diagnostic info?



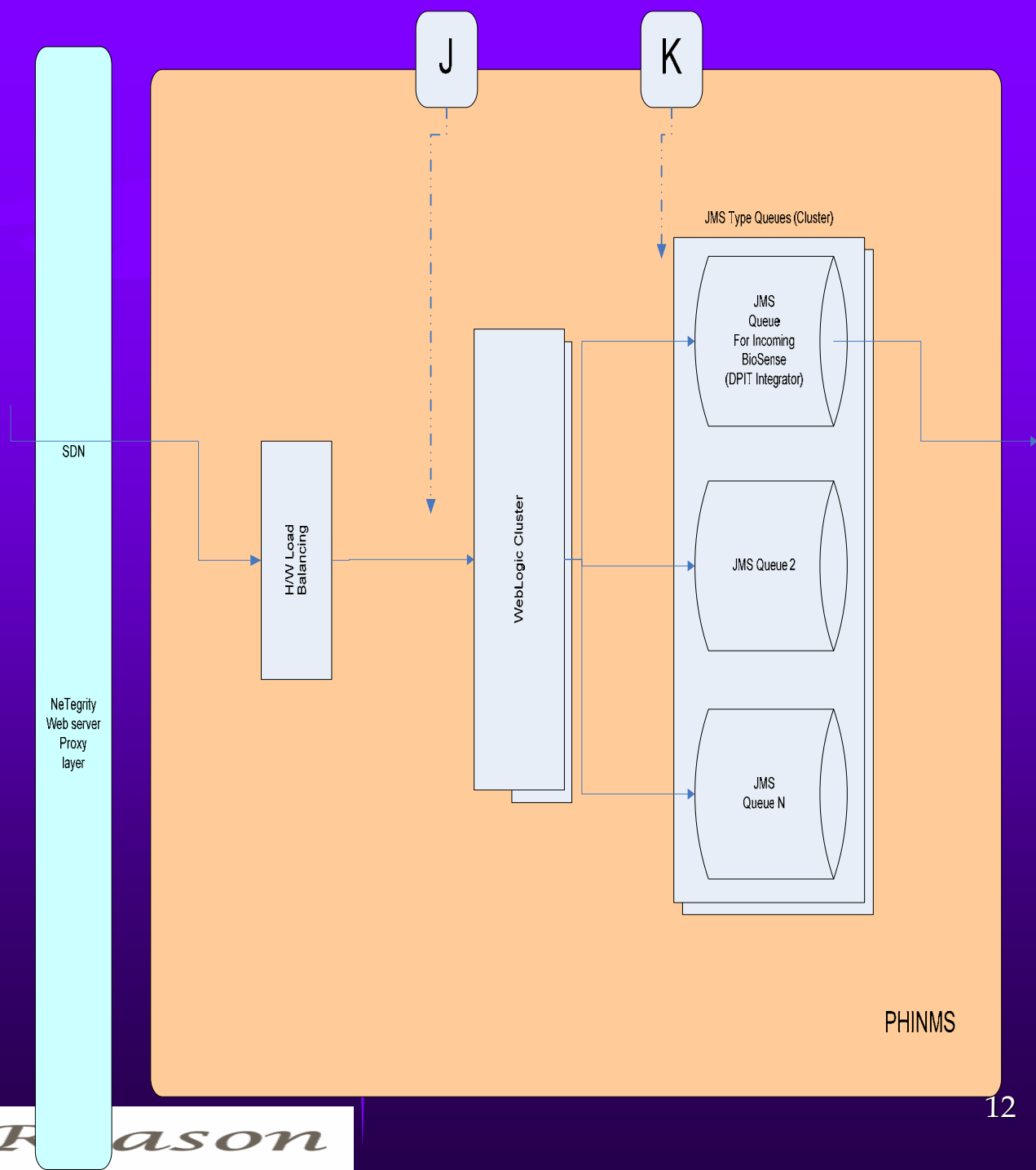
Current BioSense Framework

Message Processing

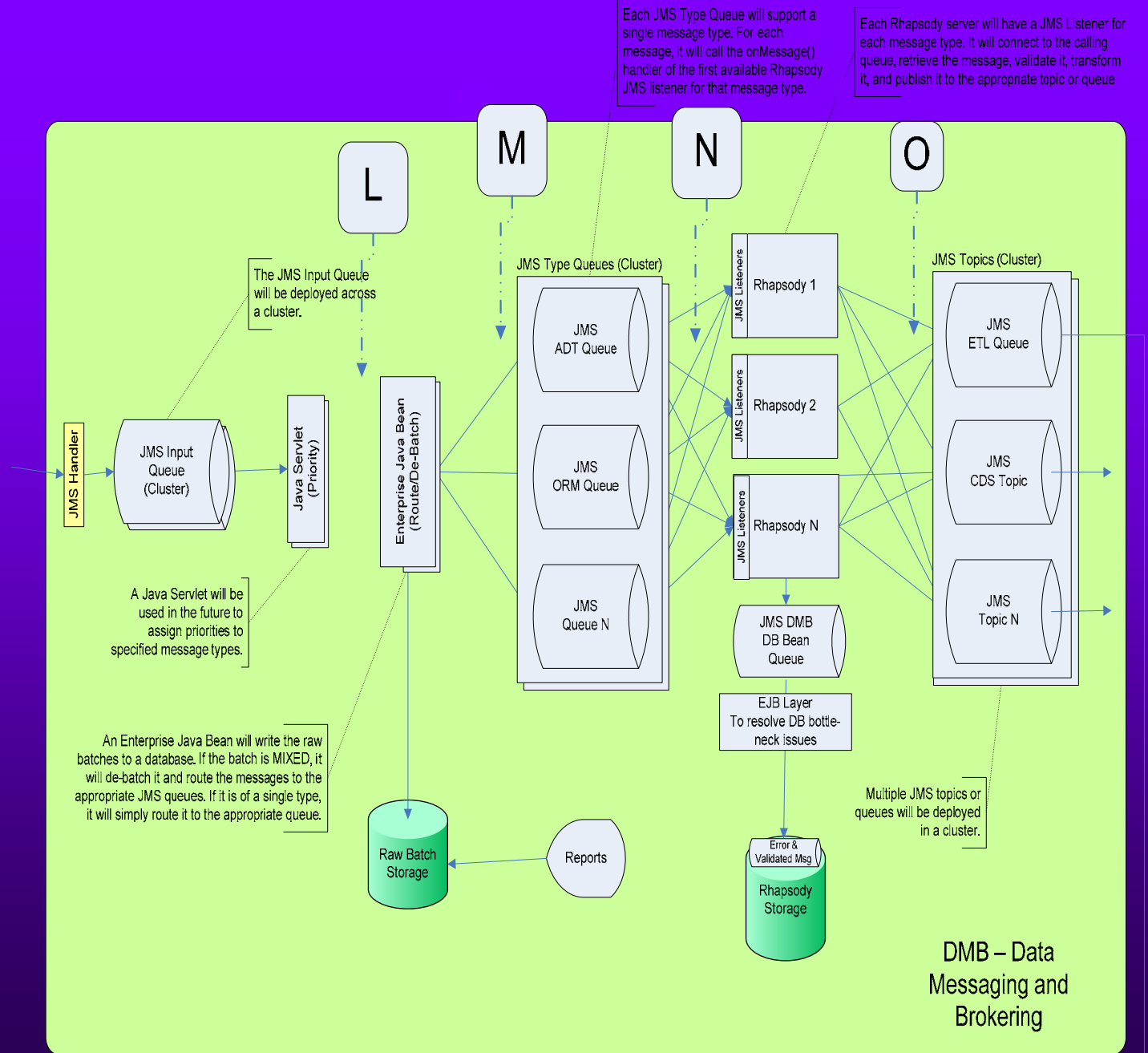


DPIT Integrator
and Linker

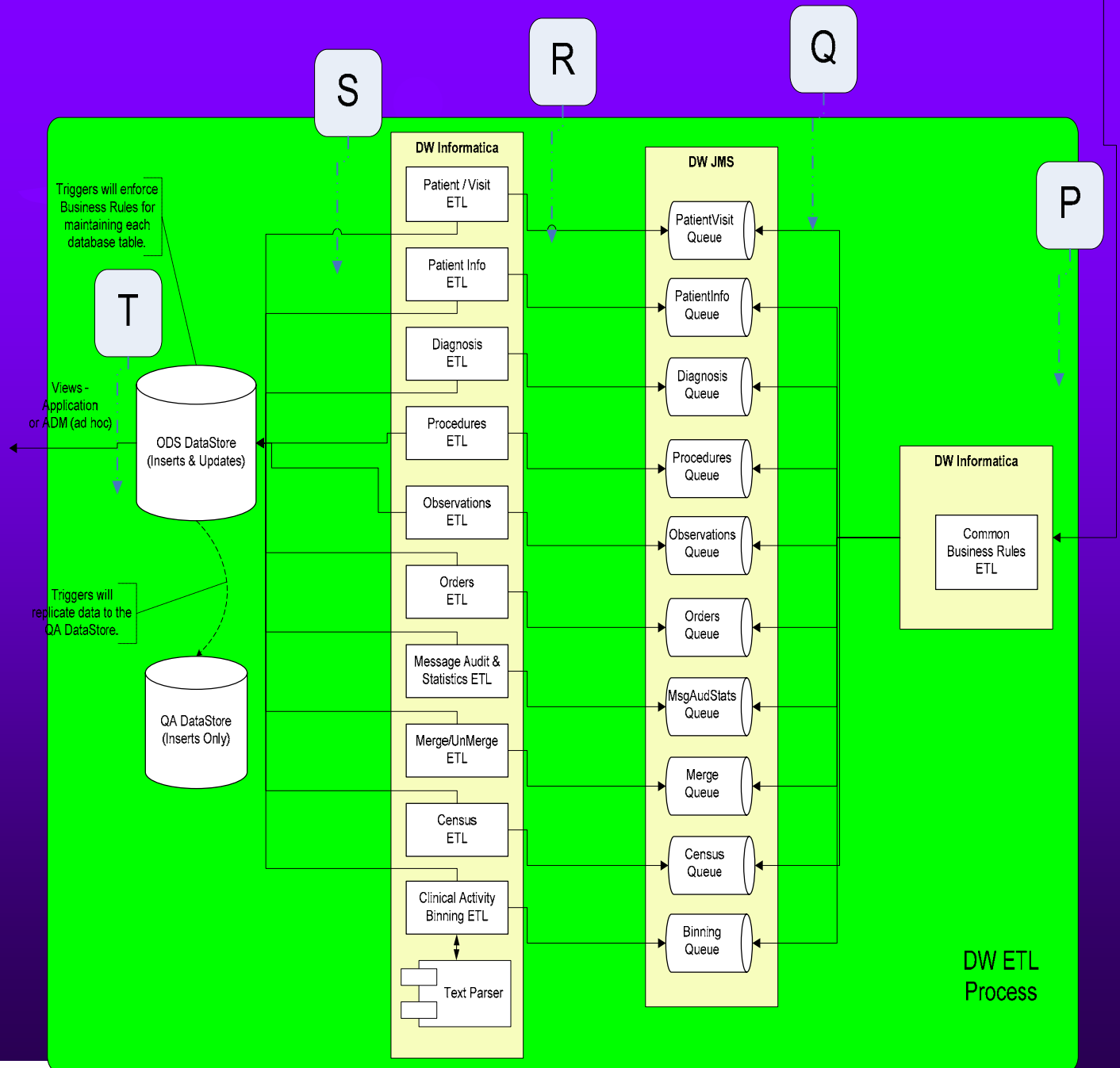
Load Balancing



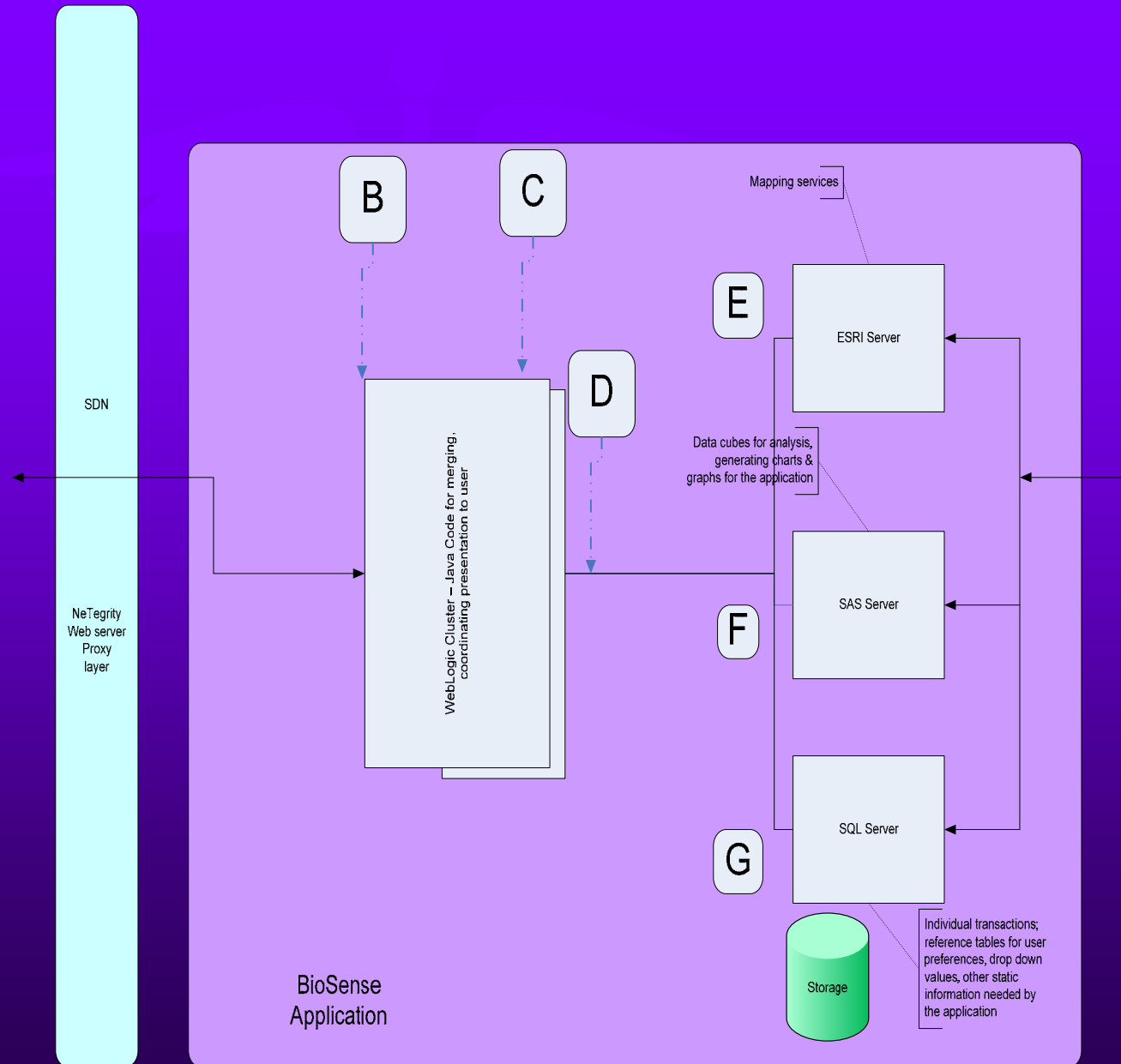
Message Type Filter



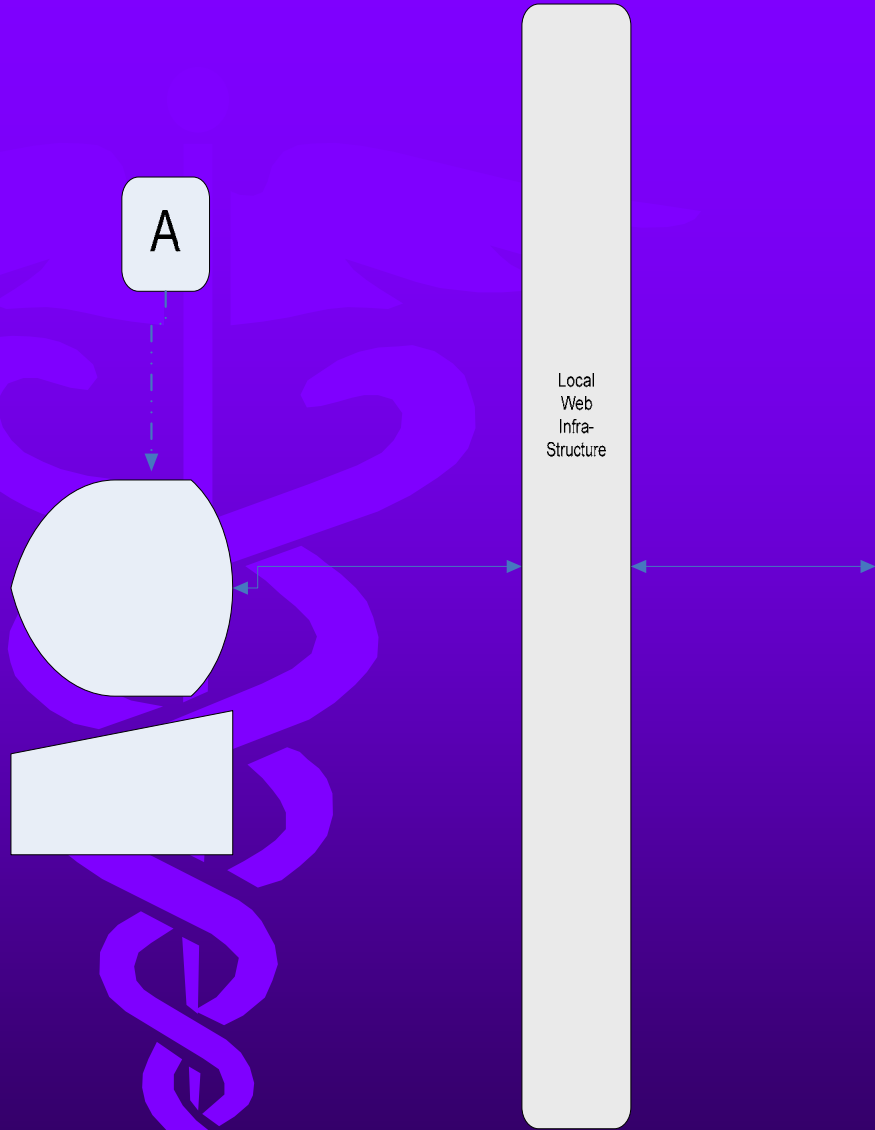
ETL Processing



AV and OTP



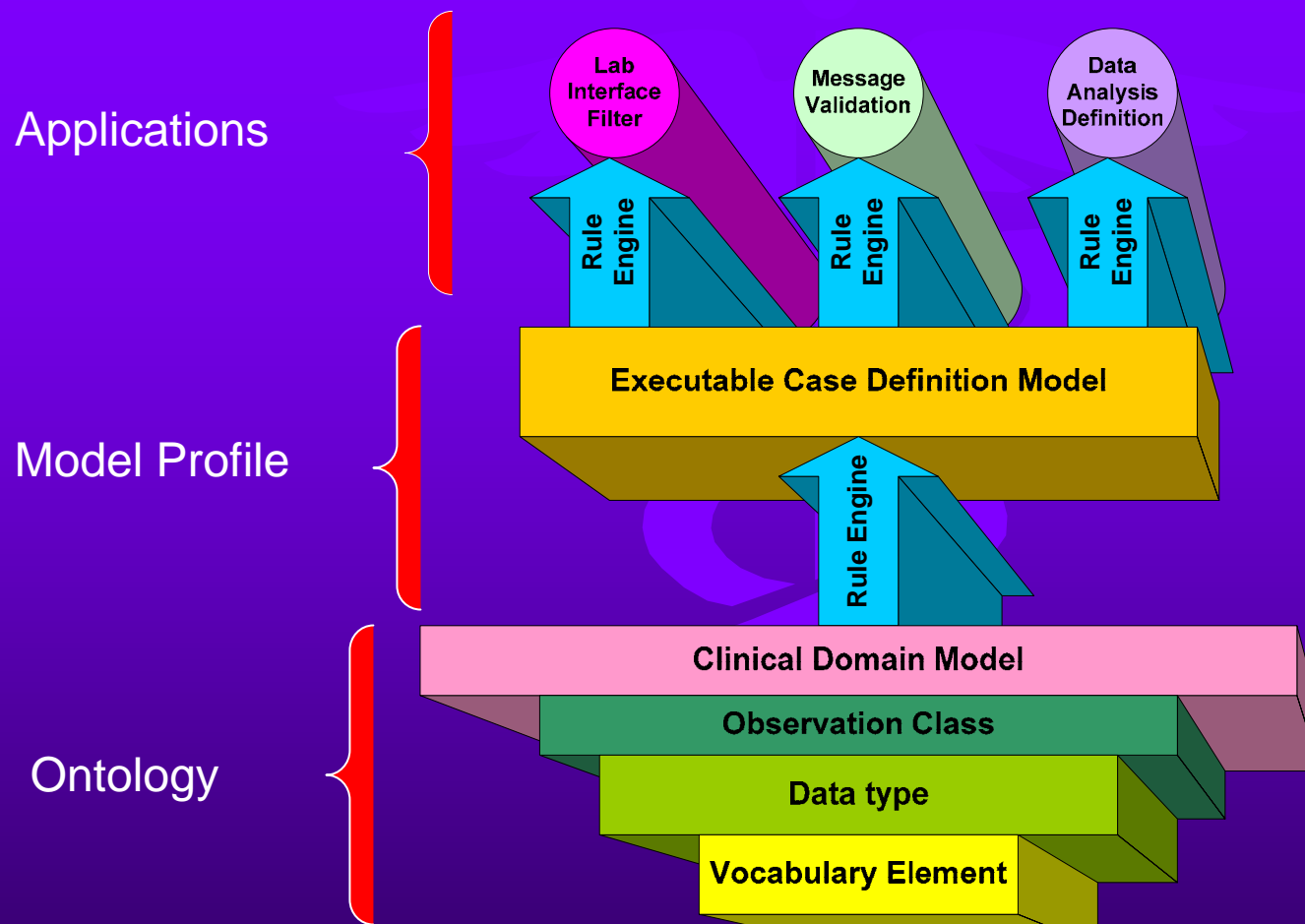
End User Views

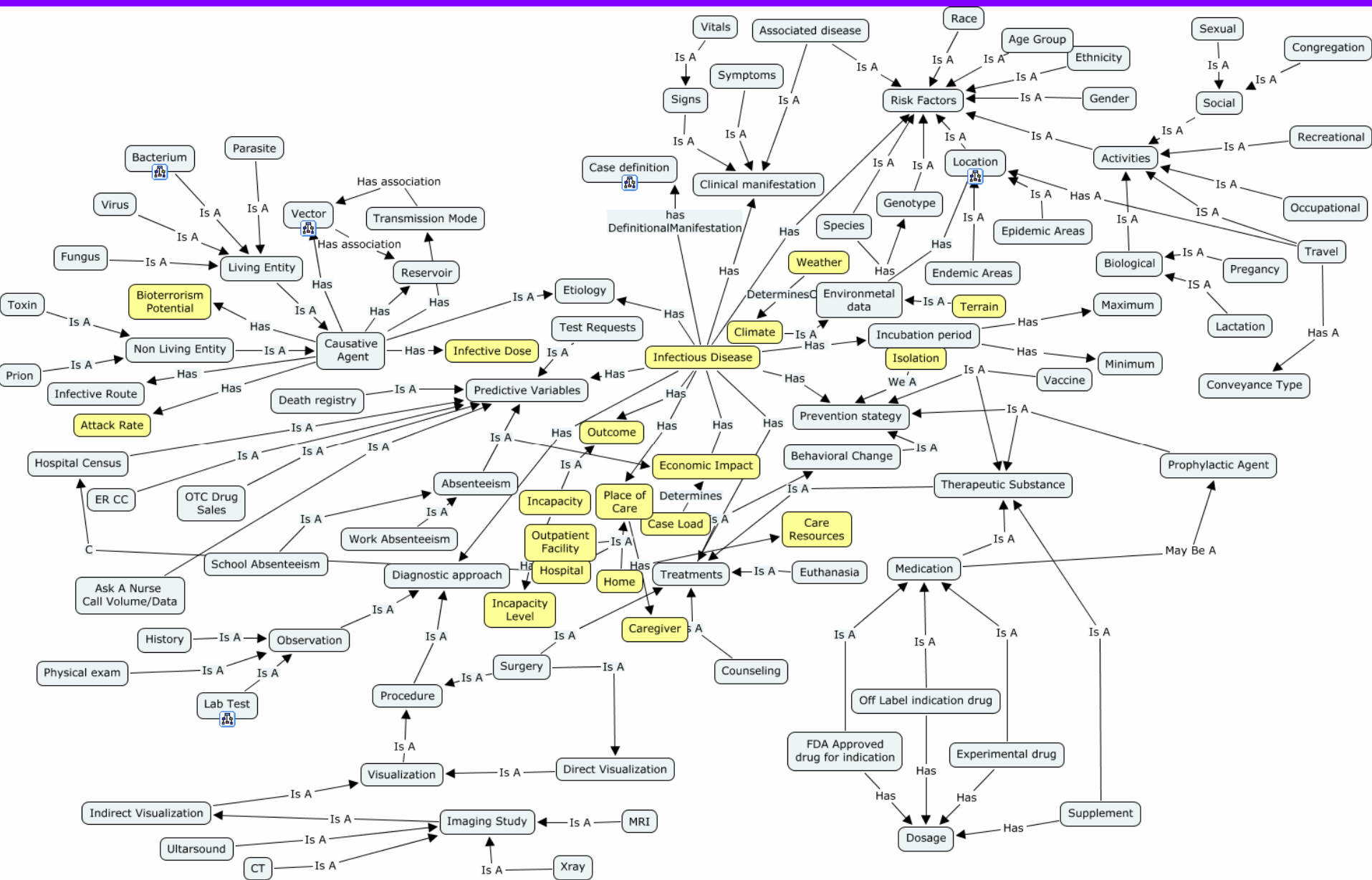


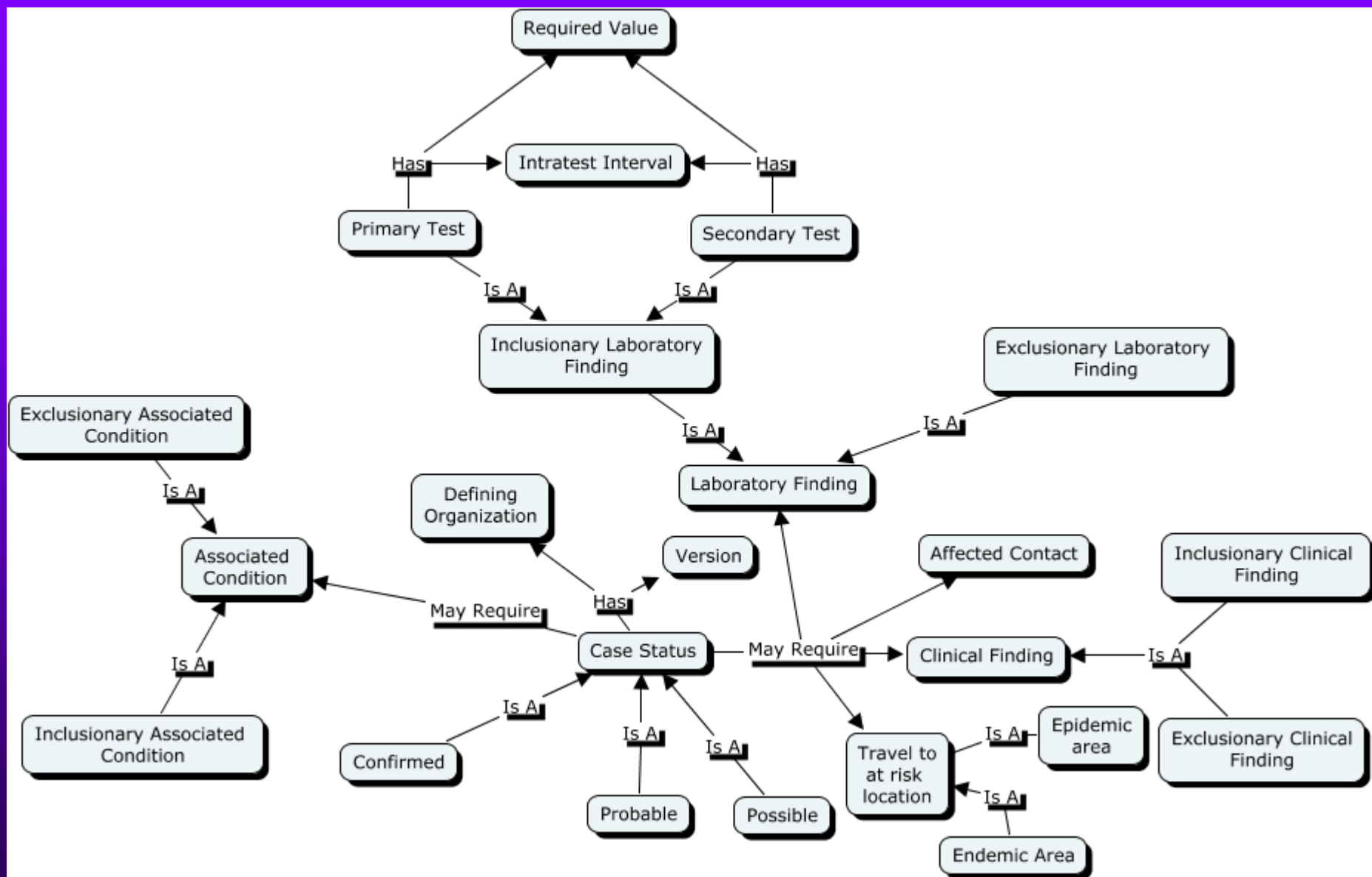
BioSense Users
On
The World Wide Web



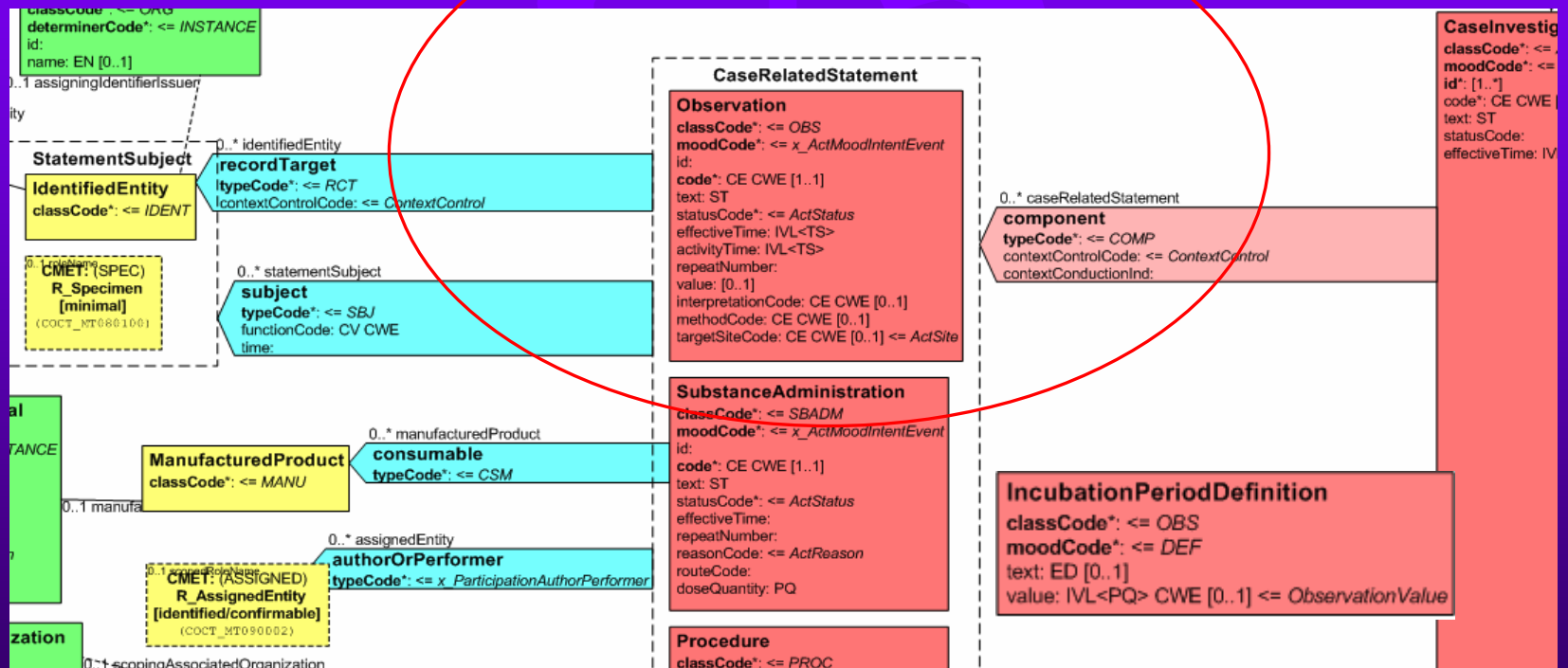
The OntoReason PH Ontology







Information Model



Concept In HL7 V3 DataType

The screenshot displays the OntoReason interface for editing an HL7 V3 Data Type concept. The main form contains the following fields:

- Code:** 49727002
- CodeSystem:** 2.16.840.1.113883.6.96
- CodeSystemName:** SNOMED CT
- CodeSystemVersion:** 0601core
- Version:** 01/31/2006
- ConceptStatus:** Active
- DisplayName:** Cough
- DateChanged:** (empty)
- OriginalText:** Cough
- Superclasses:** Functional finding of respiratory tract (highlighted)
- Subclasses:** Decreased coughing, Bovine cough, Nocturnal cough, Early morning cough, Spasmodic cough, Hemoptysis, Chronic cough, Cough at rest, Cough suppression, Postural cough, Dry cough, Cough with fever
- Translation:** Cough (finding), Observation of cough, Finding of cough, Cough (highlighted with a red box)
- Name:** Lynch82824016SN
- BioSenseMatch:** toss 1221, cough 300, 7862 3902, Cough 3298

Annotations on the image:

- Code:** Points to the Code field.
- Term:** Points to the DisplayName field.
- Parent:** Points to the Superclasses list.
- Children:** Points to the Subclasses list.
- BioSense Terms:** Points to the BioSenseMatch list.
- Other code Systems and synonyms:** Points to the Translation list.

Conceptual and Syntactical

Level

Condition Observation

- Diplopia
- Blurred vision
- Blurred vision (instance)
- Sy
- Dr
- Dy
- Dy
- Mu
- Dil
- Ptc
- Me

Blurred vision NOS (instance of Disorders_MetaClass, internal name is Lynch3.)

Code	CodeSystem	CodeSystemName
267629007	2.16.840.1.113883.6.96	SNOMED CT
CodeSystemVersion	DateCreated	Qualifier
0601core	01/31/2006	
ConceptStatus	DisplayName	
Active	Blurred vision NOS	
DateChanged	OriginalText	
	Blurred vision NOS	

```

<?xml version="1.0" encoding="UTF-8"?>
< xmlns:hl7v3=" http://www.w3.org/TR/WD-xlink" xmlns:hl7="urn:hl7-org:v3"
xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"/>

<xsiType="CE" code code="267629007 "
codeSystem="2.16.840.1.113883.6.96 "
codeSystemName="SNOMED CT"
codeSystemVersion="0601Core "
displayName="blurred vision NOS "
originalText="blurry vision ">
</code>
    
```

HL V3 Class Object

The screenshot shows the HL V3 Class Object editor for the 'Cough' class. The interface includes a left sidebar with a tree view of the class hierarchy, a main editing area, and a right sidebar with additional information. The 'Cough' class is selected, and its properties are displayed. The 'Text' field contains 'Cough'. The 'ClinicalObservation' field is set to 'Cough'. The 'Low' field is set to '5.0'. The 'Width' field is empty. The 'High' field is set to '5.0'. The 'Unit' field is set to '% percent'. The 'ClassCode' field is set to 'observation'. The 'MoodCode' field is set to 'event (occurrence)'. The 'Citations' field contains 'Young LS, Bicknell DS, Archer BG,'. The 'NegationInd' checkbox is unchecked. The 'Subclasses' list includes 'Decrease', 'Bovine co', 'Nocturnal', and 'Early morr'. The 'Display Name' field is circled in red. The 'Frequency for each object' callout points to the 'High' and 'Unit' fields. The 'References for each object' callout points to the 'ClassCode' and 'MoodCode' fields.

CodeSystem: 2.16.840.1.1...

DateCreated: 01/31/2006

Display Name: Cough

OriginalText: Cough

Subclasses:

- Decrease
- Bovine co
- Nocturnal
- Early morr

Cough (instance of ClinicalObservation_MetaClass, inter...)

Text: Cough

Documentation:

ClinicalObservation: Cough

Low: 5.0

Width:

High: 5.0

Unit: % percent

ClassCode: observation

MoodCode: event (occurrence)

Citations: Young LS, Bicknell DS, Archer BG,

NegationInd: ☐

Frequency for each object

References for each object

Clinical Domain Object

Name
Foodborne botulism investigation

Public Health Case Event
● Foodborne botulism

Incubation Period
● Botulism incubation period

Investigated Agent
● Botulism toxins

LabObservation

- C botulinum toxin detected in stool
- C botulinum toxin detected in gastric aspirate
- C botulinum toxin detected in food
- C botulinum toxin detected in serum
- C botulinum toxin E detected in stool
- C botulinum toxin E detected in gastric aspirate
- C botulinum toxin E detected in food
- C botulinum toxin E detected in serum
- C botulinum toxin A detected in stool
- C botulinum toxin A detected in gastric aspirate
- C botulinum toxin A detected in food
- C botulinum toxin A detected in serum
- C botulinum toxin F detected in stool

Vaccine

Radiology Observation
● CT of head normal

Condition Observation

- Diplopia
- Blurred vision
- Bulbar weakness
- Symmetric descending paralysis
- Dry mouth
- Dysphagia
- Dysarthria
- Muscle weakness

Procedures

- Tensilon test with electromyographic recording
- Diagnostic lumbar puncture

Therapeutic Medication
● {BOTULISM ANTITOXIN TRIV A,B&E 5000 UNT Injectable, E

Nested MetaClass



Investigated Agent

● Botulism toxins

● Botulism toxins (instance of EntityClassMaterial_MetaClass, internal name is 0...

Name

Botulism toxins

ClassCode

● MAT material

RiskCode

● BHZ biohazard

HandlingCode

● CREF Critical refrigerated temper...

DeterminerCode

● INSTANCE specific

Causative Agent|246

● Botulinum toxin type A
● Botulinum toxin type B
● Botulinum toxin type E
● Botulinum toxin type F

Citations

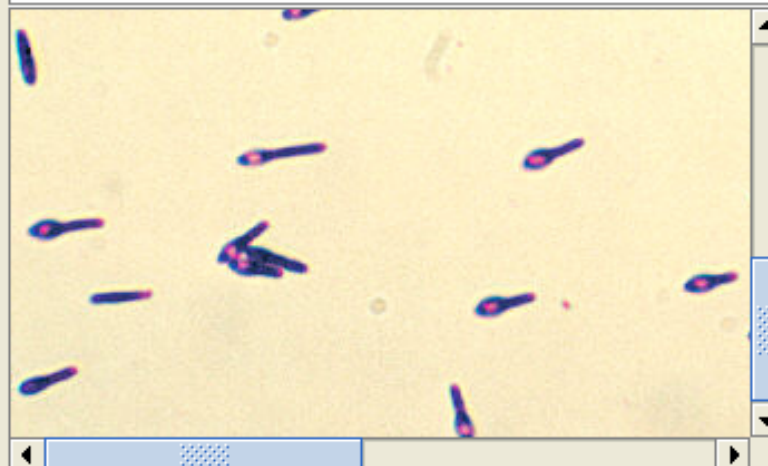
● Control of Communicable Diseases
● <http://www.dshs.state.tx.us/lab/mr>
● CDC Public Health Image Library (PHIL)

Documentation

Image from CDC/Dr. George Lombard

Image

c:\ontoreason_images\bottox.jpg



NegationValue

Laboratory Observation HL7 V3 mapped to V2

C botulinum toxin detected in stool (instance of LabObservation_MetaClass, in...

Text C botulinum toxin detected in stool	Documentation	Citations http://www.cdc.gov/epo/dphsi/cas http://www.dshs.state.tx.us/lab/mr
LabObservation.code C bot Tox XXX QI		
InterpretationCode A Abnormal	MethodCode 0025 Bioassay	ClassCode observation
TargetSiteCode Stool specimen		MoodCode event (occurrence)
Observation.Value Detected	InterpretationRange Not detected	

SPM-4

OBX-3

OBX-17

OBX-8

OBX-7

OBX-5

Map HL7 Message segments to Ontology Slots

```
<OBX.2>NM</OBX.2>  
- <OBX.3>  
  <CE.1 />  
  <CE.2>CSF protein conc</CE.2>  
  <CE.5>CSF protein mg%</CE.5>  
</OBX.3>  
<OBX.4>34</OBX.4>  
- <OBX.5>  
  <NM.1>34</NM.1>  
</OBX.5>  
- <OBX.6>  
  <CE.1>mg/dL</CE.1>  
</OBX.6>
```

LabObservation.code

This slot is populated by an instance of the LOINC coded value that represents the ordered test.

Value Type
Class

Allowed Superclasses
LOINC Code System

Cardinality
☐ required at least
☐ multiple at most

Minimum Maximum Inverse Slot

:DIRECT-TYPE
:STANDARD-SLOT
HL7V2_EQ

:SLOT-CONSTRAINTS

as HL7V2_EQ

Name
LabObservation.code

HL7V2 Field
<OBR.4> ORM
<OBX.3> ORU
<OBR.4> ORU

Documentation
This slot is populated by an instance of the LOINC coded value that represents the ordered test.

TranslationRequirement
None

Vocabulary Recommended
Laboratory Test Name



Ontology Services Platform

Technical Foundations

Platform Models

- Enterprise PHIN SOA
- Web Services
- Application Libraries
- LexPHIN Database

Application Models

- Individual Reasoners Patterns - Languages
- Intelligence & Analytics Workbench - Tools
- CTS & LexPHIN Services - Standards

Domain Models

- PH Reference Ontology
- PHIN VS
- BioSense Msg HL7 V2.x

Message Structure

Ontology Extraction

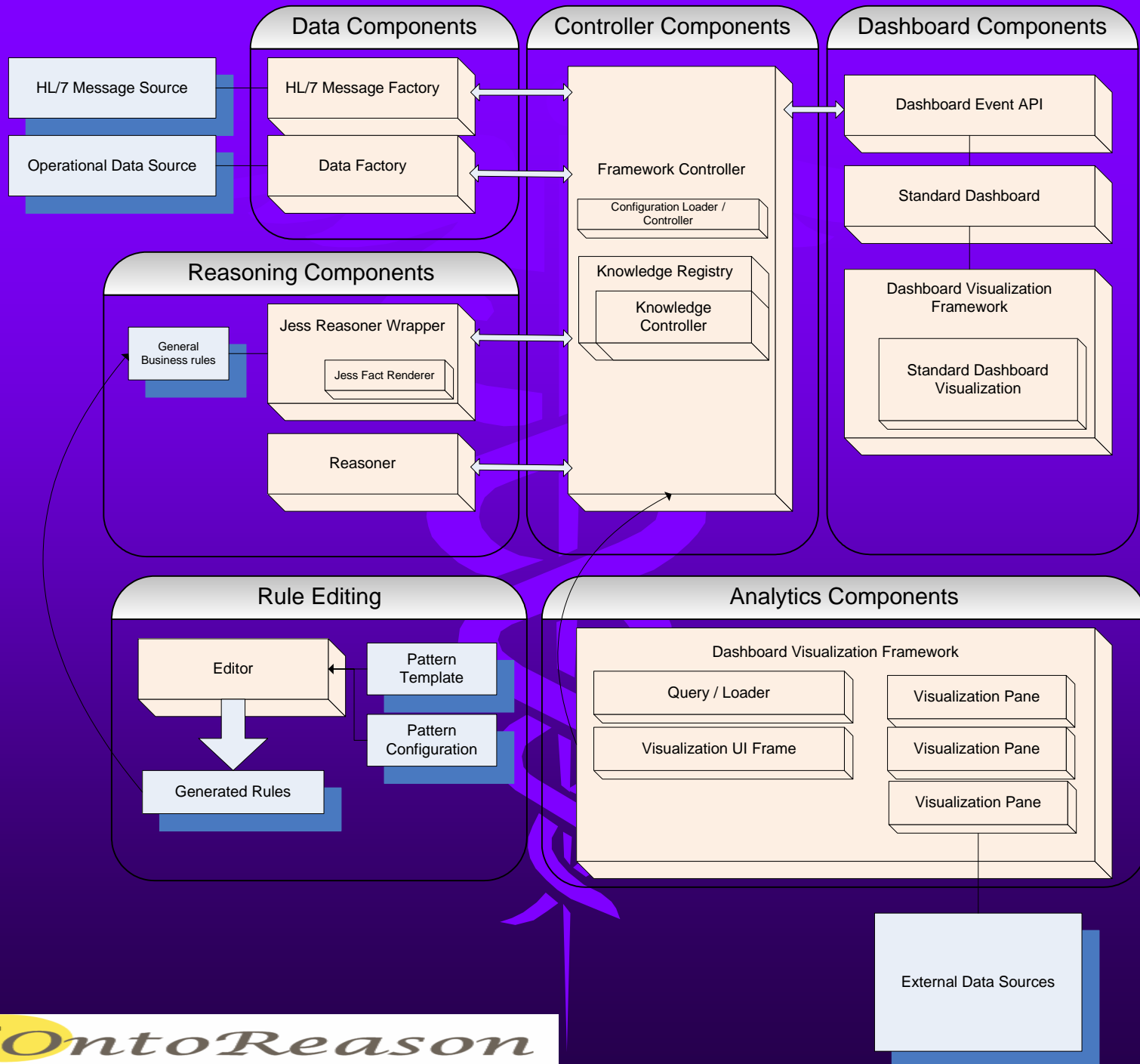
- Creating an application ontology from the reference ontology
 - Identify the core ontology classes
 - Create an object representation that maintains the ontology data
 - Generate cross reference indexes for core relationships
 - Lab tests to case investigations
 - Organism/ Agent to case investigations
 - Other significant relationships
 - Identify “Used” vocabulary
 - Create vocabulary subsets that identify specific vocabularies concepts that are used within the ontology
 - Create code to code mapping indexes
- **This produces a general purpose extraction that is suitable for various purposes**

Ontology Extraction

- **Additional activities performed for specific problem solutions**
 - Inclusion of additional vocabulary value sets
 - Generation of additional vocabulary indexes to maintain certain parent/child relationships
 - Incorporation of certain additional term mappings
 - Alternate spellings
 - Concept mappings to syndrome/sub-syndrome
 - Generation of text search algorithms
- **Loadable data married with functional API**
 - Java object serialized for easy loading
 - Java API providing lookup/query functionality

Ontology Representation

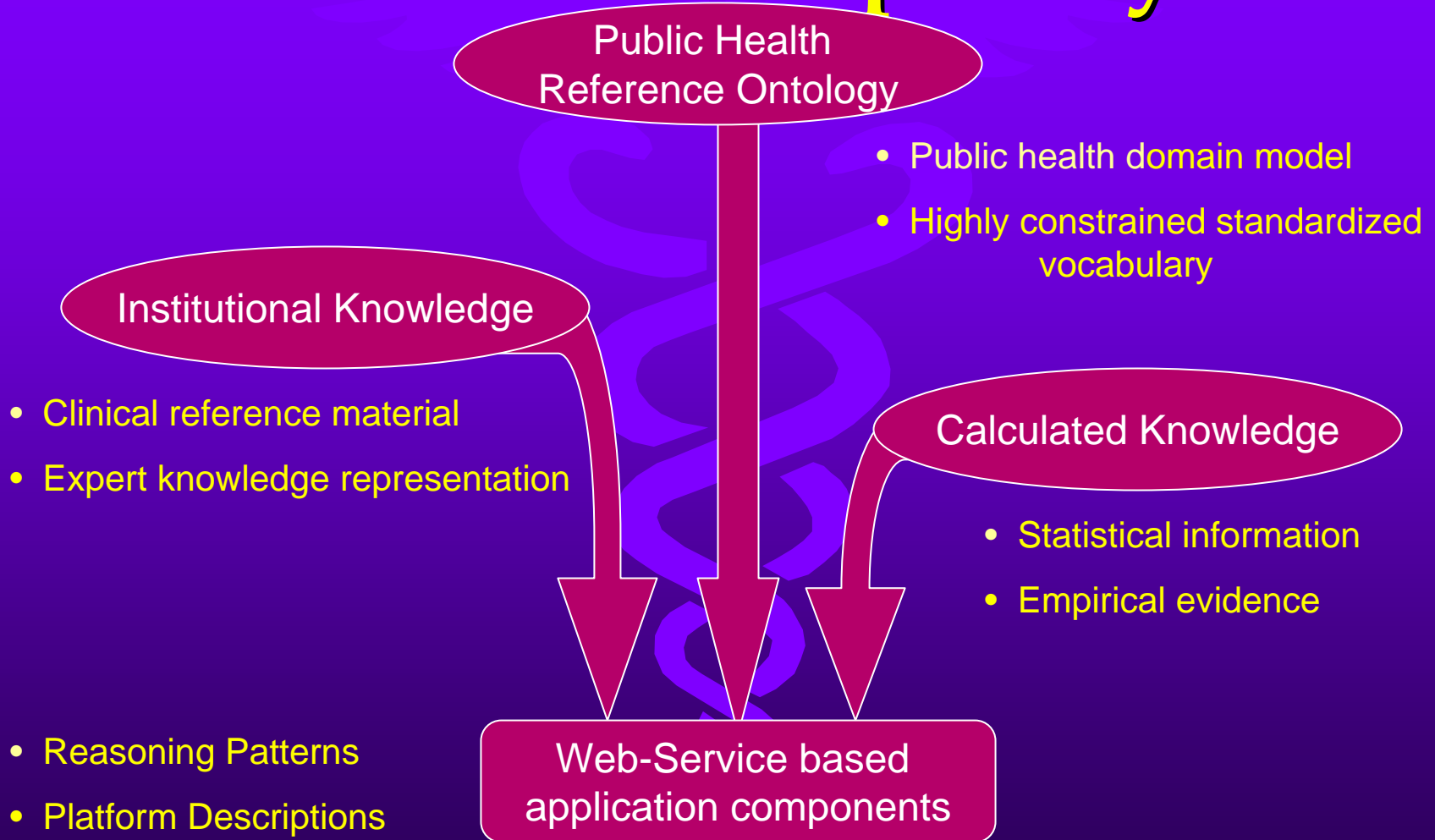
- **Jess rule engine representation**
 - The Jess rule engine utilizes an enhanced RETE algorithm to provide an execution platform for declarative rule base
 - Data in Jess is represented as a set of declared facts
 - Facts can be either structured or unstructured
 - Ontology data is represented as a set of instance data represented as structured facts
 - The ontology can either be expressed as a script or loaded directly into the rule engine at runtime
- **Rule definition**
 - Rules which describe core case definitions are constructed
 - The ontology facts are merged with the core set of rules to provide the base representation for the entire ontology





Message Analytics Workbench

Model Driven Expert System

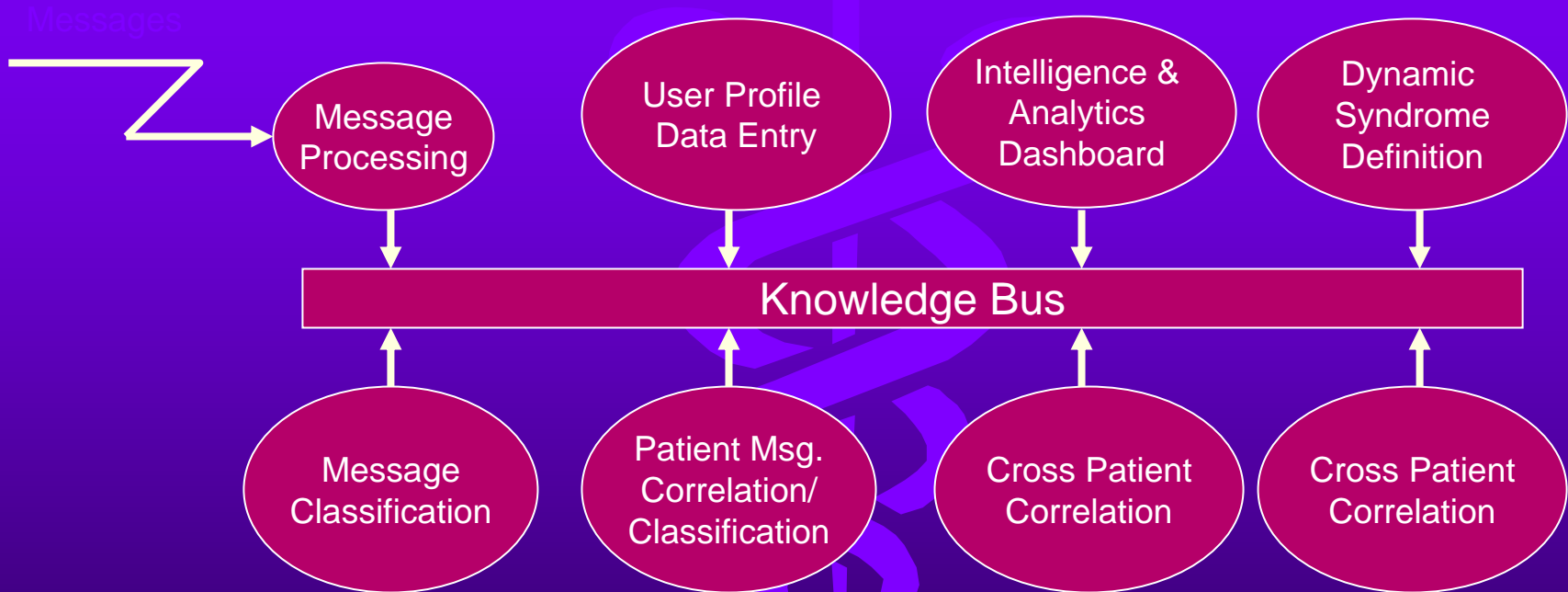


BioSense Message Data Source

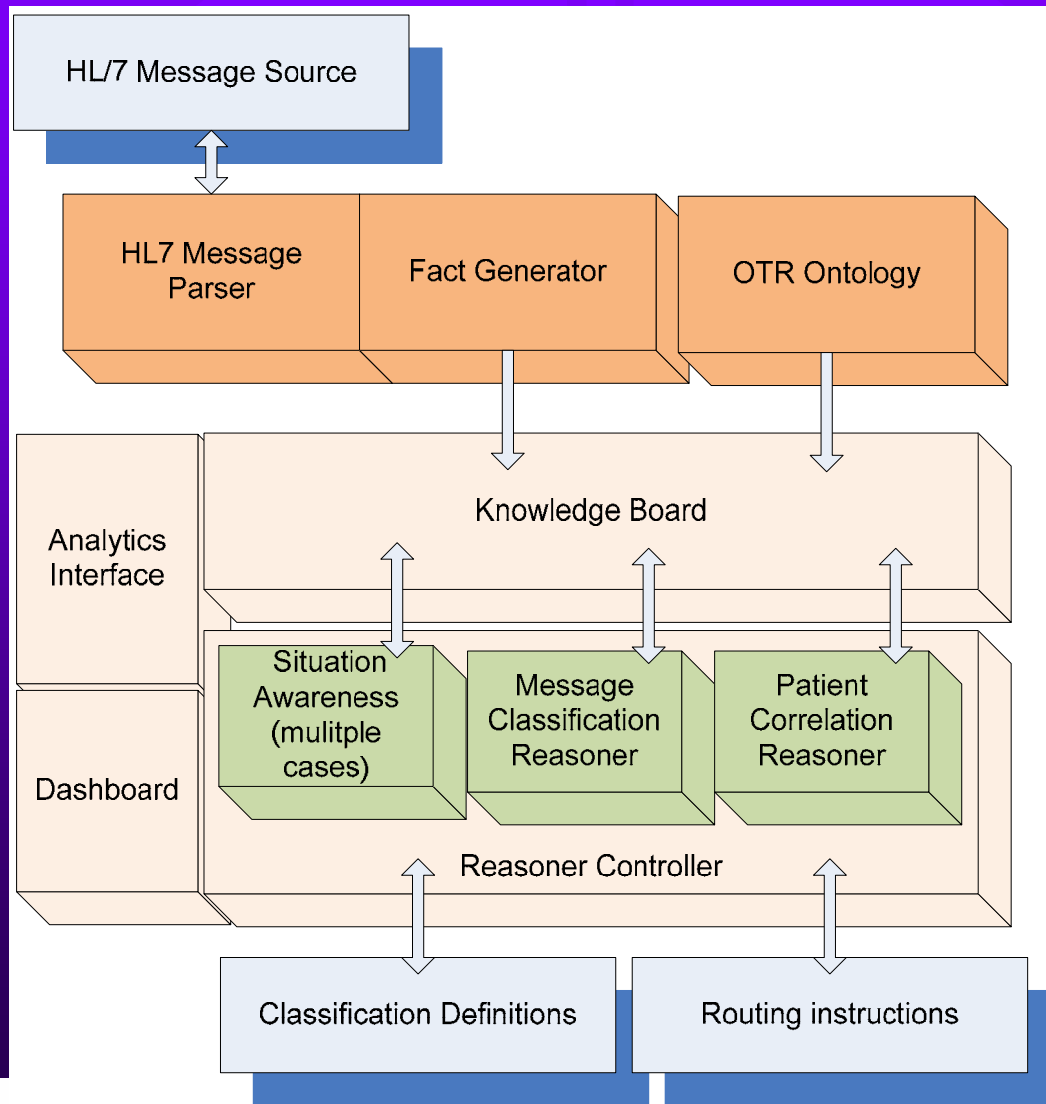
- HL7 Version 2.5
- XML representation
- Laboratory (ORU) message
- Spinal fluid protein

```
<?xml version="1.0" encoding="ISO-8859-1" ?>
<!-- edited with XMLSpy v2007 sp2 (http://www.altova.com) by Cec
- <ORU_R01>
+ <MSH>
- <ORU_R01.PATIENT_RESULT>
+ <ORU_R01.PATIENT>
- <ORU_R01.ORDER_OBSERVATION>
- <OBR>
  <OBR.1>1</OBR.1>
  + <OBR.2>
  + <OBR.3>
  - <OBR.4>
    <CE.4>CSF protein</CE.4>
    <CE.5>CSF protein</CE.5>
  </OBR.4>
  + <OBR.7>
  + <OBR.8>
  + <OBR.14>
  - <OBR.15>
    - <SPS.1>
      <CWE.4>Spinal Fluid</CWE.4>
    </SPS.1>
  </OBR.15>
  + <OBR.22>
    <OBR.24>LA</OBR.24>
    <OBR.25>F</OBR.25>
  </OBR>
- <ORU_R01.OBSERVATION>
- <OBX>
  <OBX.1>1</OBX.1>
  <OBX.2>NM</OBX.2>
  - <OBX.3>
    <CE.1 />
    <CE.2>CSF protein conc</CE.2>
    <CE.5>CSF protein mg%</CE.5>
  </OBX.3>
  <OBX.4>34</OBX.4>
  - <OBX.5>
    <NM.1>34</NM.1>
  </OBX.5>
  - <OBX.6>
    <CE.1>mg/dL</CE.1>
  </OBX.6>
```

Demonstration of Basic Platform

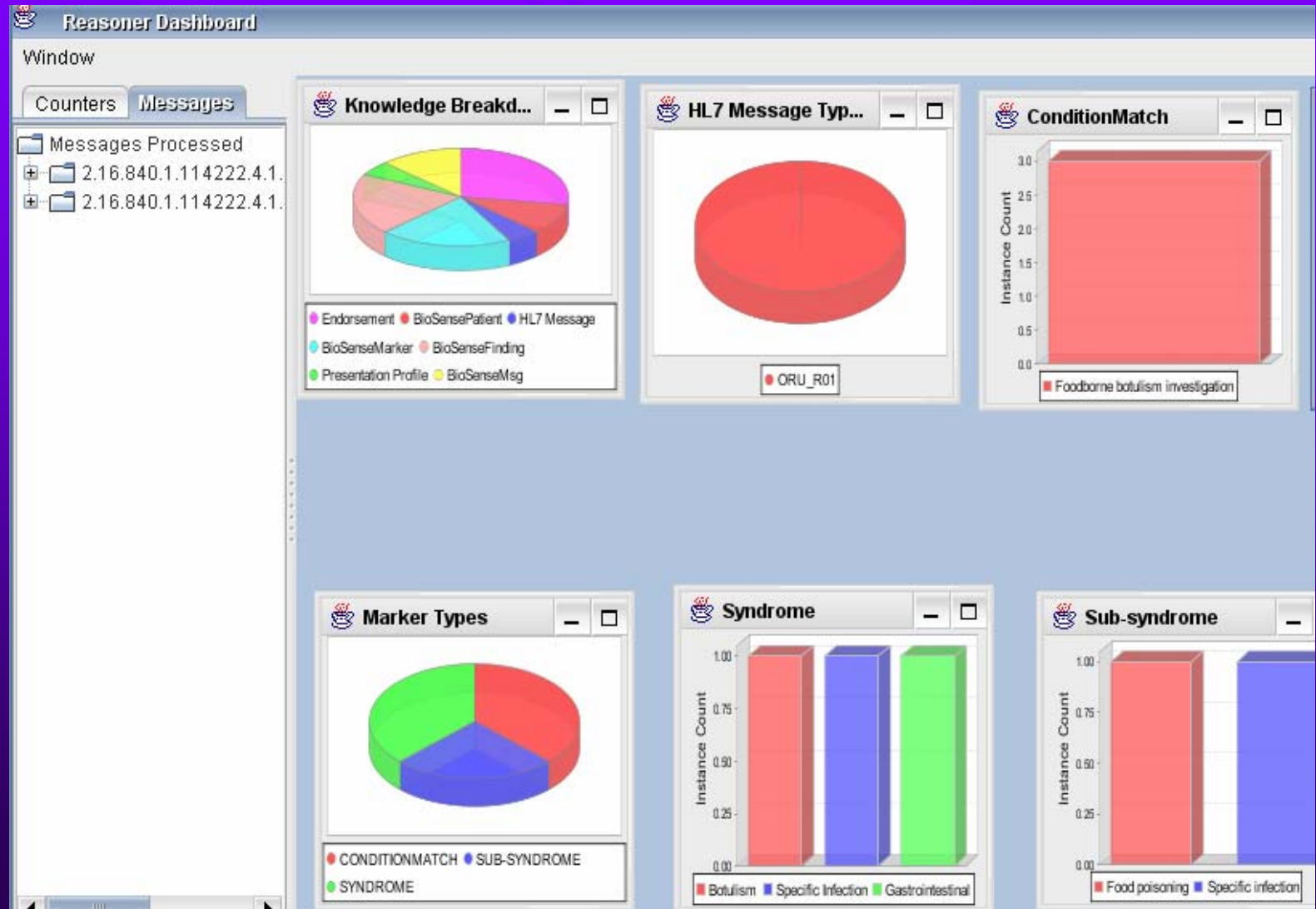


Message Analytical Workbench





Reasoner results





OntoReason

How the Rules Work

- JESS template is like a class in Java
- Template can but does not have to declare attribute type, default values, and if an attribute is a single value or a list
- Facts asserted into the expert system directly from the ontology - based upon the template

Interaction of Ontology And Rules

- Information from the ontology and generated template facts imported into the expert system and operated on by a variety of reasoners
- This way the reasoner knowledgebase can have a relatively small footprint vs the reference ontology

JESS Rule From Template

```
(defrule ClinicalFindingConditionMatchesInitial
  (Classifier-State DevelopFindingsAndEndorsements)
  (observation (obsId ?obsId)(msgId ?mId)(patientId ?patientId)(chiefComplaint ?chief)
    (code ?observation) (obsType ClinicalFinding) (dataQualityFactor ?quality)
    (dqfReason ?qualityReason) (originationDate ?oDate) (originationDateType ?oType))
  (nnd-finding (cond-code ?condCode) (finding-code ?observation)(high ?prob) (ratio
?ratio))
  (nnd-condition (cond-code ?condCode) (description ?desc))
  (not (BSFinding (msgId ?mId)(finding ?condCode)))
=>
  (assert (BSFinding (msgId ?mId)(patientId ?patientId) (findingId ?*ClassId*)
    (finding ?condCode)(findingType Condition)(findingDesc ?desc)
    (originationDate ?oDate)(originationDateType ?oType)))

  (assert (Endorsement (msgId ?mId)(findingId ?*ClassId*)(findingCorrelation ?ratio)
    (findingType Condition)(finding ?condCode)(findingProb ?prob)
    (endorsementId (+ ?*ClassId* 1))(endorsement
?*Supportive*)(endSymbol *Support*)
    (endorsementType *ClinicalFinding*)(rule
*ConditionMatch*)(endorsementContext ?context )
    (obsId ?obsId)(obsCode ?observation)(obsQuality
?quality)(explanation ?qualityReason )))
  (bind ?*ClassId* (+ 2 ?*ClassId*)))
```

A couple of things to remember

- This is an Intelligence and Analytics toolkit
 - Used to exploit the expert knowledge of the organization to provide simple to configure application components
 - Real-time processing
 - Historical data for analysis, knowledge discovery and re-classification
 - Findings can be reused to tune and validate real-time processing
- Classification tools are based upon a very quick assessment generalized across all conditions
- The classification weights can be greatly improved based upon empirical data analysis
- Algorithms are simple to tune and extend (including geo-spatial and temporal services)
- The use-cases were made from some limited set of assumptions
- We used a condition centric analysis

Next Steps

- Add additional domain centric rules for better agent classification
- Overcome limitations of ontology size and maintenance issues by subdividing into smaller ontologies
- Apply a novel technique to use the best aspects of Frames and OWL structures
 - (see the demo)
- Develop simple domain expert editing tools for rules and knowledge

Questions and Answers