

Knowledge Representations

- How else can we represent knowledge in addition to formal logic?

Common Knowledge Representations

- Formal Logic ✓
- Production Rules
- Semantic Nets
- Schemata and Frames

Production Rules

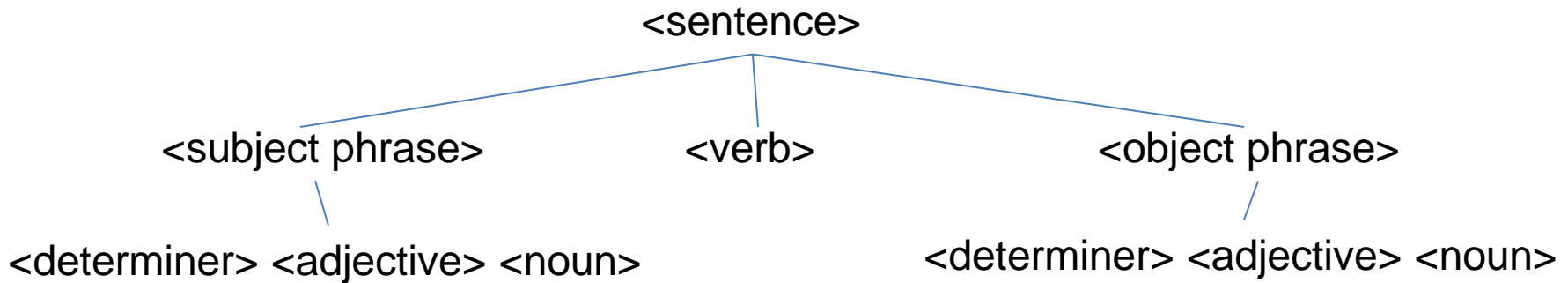
- Frequently used to formulate the knowledge in expert systems.
- BNF is commonly used in Computer Science.

◆ for a subset of the German language

```
<sentence>      -> <subject phrase> <verb>
                  <object phrase>
<subject phrase> -> <determiner> <adjective> <noun>
<object phrase>  -> <determiner> <adjective> <noun>
<determiner>    -> der | die | das | den
<noun>          -> Mann | Frau | Kind | Hund | Katze
<verb>         -> mag | schimpft | vergisst |
                  verehrt | verzehrt
<adjective>    -> schoene | starke | laute | duenne
```

◆ for a subset of the German language

<sentence>	->	<subject phrase>	<verb>	
			<object phrase>	
<subject phrase>	->	<determiner>	<adjective>	<noun>
<object phrase>	->	<determiner>	<adjective>	<noun>
<determiner>	->	der die das den		
<noun>	->	Mann Frau Kind Hund Katze		
<verb>	->	mag schimpft vergisst verehrt verzehrt		
<adjective>	->	schoene starke laute duenne		



Prolog

- Prolog was developed for AI applications.
- It specifies rules as Horn clauses, a subset of predicate logic.

- Example

```
male( albert ) .  
male( edward ) .
```

```
female( alice ) .  
female( victoria ) .
```

```
parents( edward , victoria , albert ) .  
parents( alice , victoria , albert ) .
```

```
sisterof( X , Y ) :- female( X ) ,  
                    parents( X , M , F ) ,  
                    parents( Y , M , F ) .
```

Prolog Expert System

```
% Automotive Diagnostic Expert System
```

```
defect_may_be(drained_battery) :-  
    user_says(starter_was_ok, yes),  
    user_says(starter_is_ok, no).
```

```
defect_may_be(wrong_gear) :-  
    user_says(starter_was_ok, no).
```

```
defect_may_be(fuel_system) :-  
    user_says(starter_was_ok, yes),  
    user_says(fuel_is_ok, no).
```

Picture Pattern from my 1974 Thesis

PATTERN = CIRCLE \$ C1 ¢ CIRCLE \$ C2 ¢

GT(VALU(C2,'RADIUS'), VALU(C1,'RADIUS')) ¢

AT(POINT(C2,'TOP'), POINT(C1,'BOT')) ¢

CIRCLE \$ C3 ¢

GT(VALU(C3,'RADIUS'), VALU(C2,'RADIUS')) ¢

AT(POINT(C3,'TOP'), POINT(C2,'BOT'))

What is it?

Advantages of Production Rules

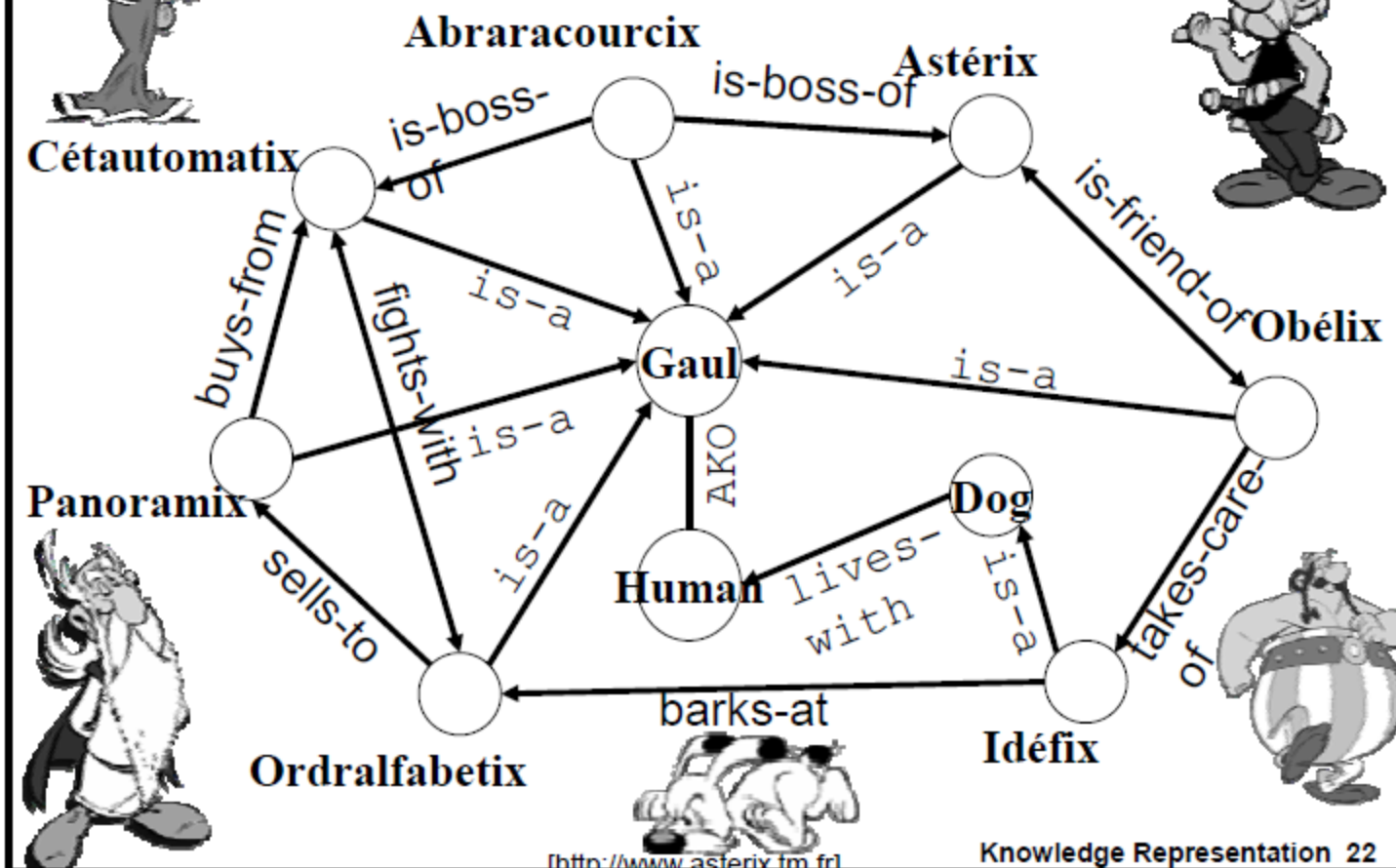
- Simpler than full predicate logic
- Still pretty expressive
- Simple backtracking search algorithms
- Easy for programmers to construct the rules
- Humans tend to understand the rules

Disadvantages?

Semantic Nets

- Graphical representation for propositional information
- Originally developed by Quillian as a model for human memory
- Nodes represent objects, concepts, situations
- Edges represent relationships

Semantix Net Example



Semantic Nets

- Relationships
 - Frequently used: IS-A, A-KIND-OF, PART-OF
 - Can be specified by the designer
- Attributes
 - Can be added to the nodes
- Advantages
 - Easy to encode and understand
- Disadvantages
 - May become large and lead to enormous searches

Related Web Developments

- **The Semantic Web**: an effort to create a web that uses the concepts from semantic nets.
- It would allow people (and programs) to better understand web content.
- Two main representations at present:
 - **RDF** (Resource Description Framework) low level, triples (**node1, relationship, node2**)
 - **OWL** (web ontology language) adds semantics to RDF

Semantic Web Languages

- **RDF (Resource Description Framework)**
 - Triples: <subject> <property> <object>
 - RDF is a data model for objects ("resources") and relations between them. These data models can be represented in an XML syntax.
- **RDFS (RDF Schema)**
 - A vocabulary for describing properties (subclass, subproperty, domain, range) and classes of RDF resources, with a semantics for generalization-hierarchies of such properties and classes.

• OWL (Web Ontology Language)

- OWL adds more vocabulary for describing properties and classes: among others, **relations between classes** (e.g. disjointness), **cardinality** (e.g. "exactly one"), **equality**, richer typing of properties, **characteristics of properties** (e.g. symmetry), and **enumerated classes**.
- There are **constraints on classes** and the types of relationships permitted between them. These provide semantics by allowing systems (**reasoners**) to infer additional information and provide classification based on the data explicitly provided.

Three “flavors” of OWL:

OWL Full

- OWL Full includes all OWL language constructs without restrictions on how they can be used.
- Not decidable

OWL DL (Description logic)

- OWL DL includes all OWL language constructs, but they can be used only under certain restrictions.
- Decidable
- Most ontologies use OWL DL

OWL Lite (even more restricted)

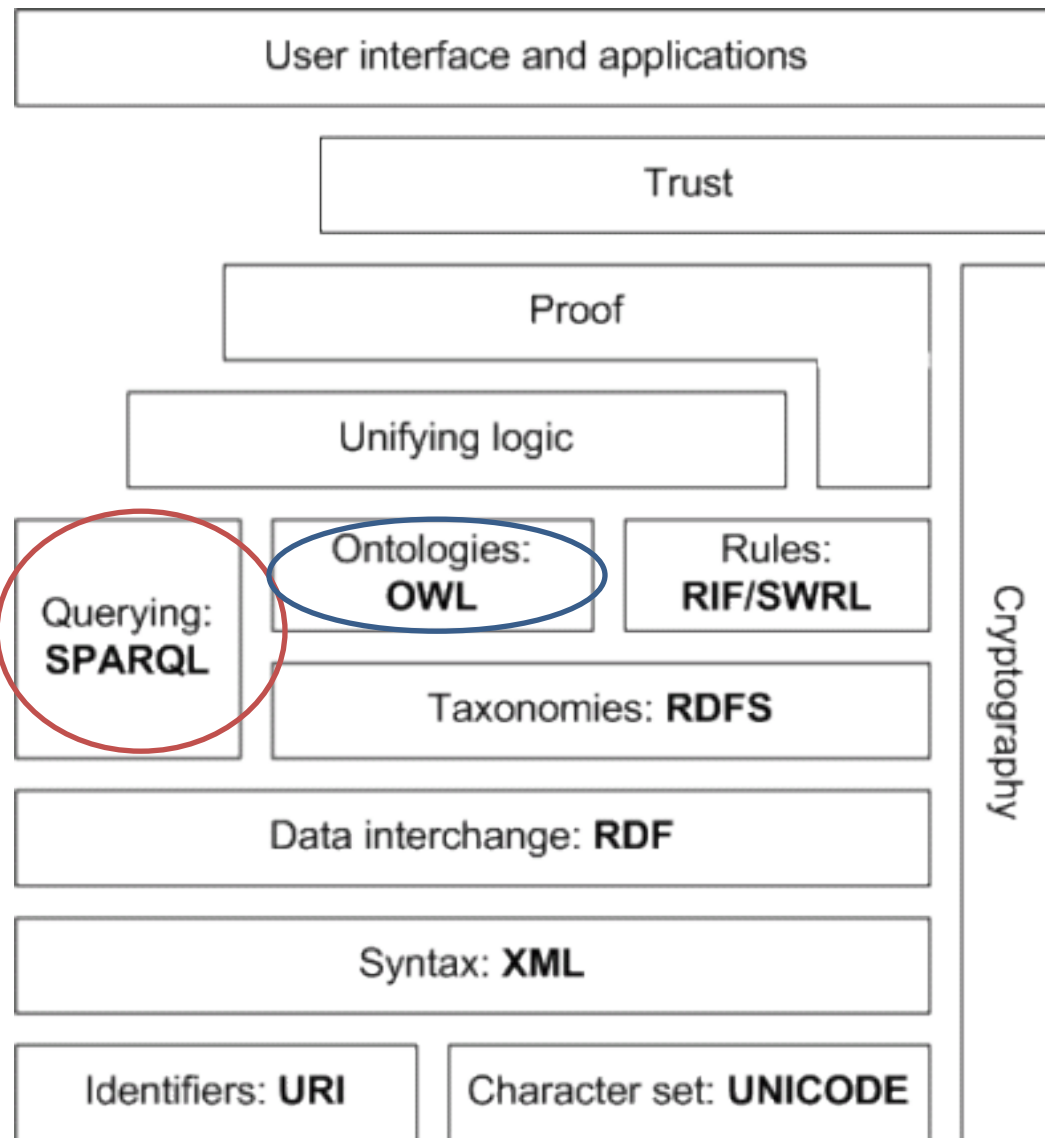
Excerpt of an OWL Ontology:

```
<rdfs:Class rdf:ID="WINE">
  <rdfs:subClassOf rdf:resource="#POTABLE-LIQUID"/>
  <rdfs:subClassOf>
    <daml:Restriction>
      <daml:onProperty rdf:resource="#MAKER"/>
      <daml:minCardinality>
        1
      </daml:minCardinality>
    </daml:Restriction>
  </rdfs:subClassOf>
  <rdfs:subClassOf>
    <daml:Restriction>
      <daml:onProperty rdf:resource="#MAKER"/>
      <daml:toClass rdf:resource="#WINERY"/>
    </daml:Restriction>
  </rdfs:subClassOf>
  <rdfs:subClassOf>
    <daml:Restriction>
      <daml:onProperty rdf:resource="#GRAPE-SLOT"/>
      <daml:minCardinality>
```


My Comments

- The international ontology community is sold on OWL, because it is powerful and portable.
- Part of the portability is that it is usually expressed in XML and then interpreted by XML readers.
- Thus, trying to read it as a human is rather difficult; it is for machines.
- But it allows people all over the world to share knowledge bases.

Semantic Web Stack



Frames

- A frame represents related knowledge about a subject
- Frames contain multiple named slots
- Slots contain values of many different kinds
 - rules, facts, images, links to other frames
- Slots can have related procedures that get executed when the value is added, modified or deleted
- Frames can be arranged in a hierarchy or graph

Simple Frame Example

<i>Slot Name</i>	<i>Filler</i>
name	Astérix
height	small
weight	low
profession	warrior
armor	helmet
intelligence	very high
marital status	presumed single

Frames

- Advantages
 - Intuitive for many applications
 - Easier to understand than logic
 - Very flexible
- Problems
 - There are inheritance problems, particularly multiple inheritance in graphs

Ontologies

- An ontology is a **formal representation** of a set of **concepts** within a domain and the **relationships** among those concepts.

Does that sound familiar?

- It allows deep understanding of and reasoning about a domain.
- UW Medical School has one enormous and now famous ontology: the **Foundational Model of Anatomy** (FMA)

Ontology Tools

- Ontology-development becomes more accessible with tools.
- **Protégé**
 - Developed at Stanford Medical Informatics
 - Is an **extensible** and **customizable** toolset for
 - **constructing** knowledge bases
 - developing **applications** that use these knowledge bases

What is Protégé?

- An ontology **editor**
- A **knowledge-acquisition** tool
- A **platform** for knowledge-based applications

An Ontology Editor

The screenshot displays the Protégé ontology editor interface. At the top, there is a toolbar with various icons for file operations and editing. Below the toolbar, the main window is divided into several panes:

- Subclass Relationship:** Shows the asserted hierarchy for the project "Anatomy_Kind_04_12". The hierarchy is as follows:
 - ▼ Anatomy_Kind
 - ▼ Anatomic_Structures_and_Systems
 - ▼ Body_Cavity
 - Abdomen
 - Mouth
 - Nasal_Cavity
 - ▶ Nasal_Sinus
 - Pelvis
 - Peritoneal_Cavity
 - Pleural_Cavity
 - Thoracic_Cavity
 - ▶ Body_Fluids_and_Substances
 - ▶ Body_Part
 - ▶ Body_Region
 - ▶ Embryologic_Structure_or_System
 - ▶ Microanatomy
 - ▼ Organ
 - Biliary_Tract (selected)
 - Bladder
 - Blood
 - Bone_Marrow

- Class Editor:** Shows the editor for the class "Biliary_Tract". It includes a "Name" field containing "Biliary_Tract" and an empty "RDFS:COMMENT:" field.
- Annotations:** A table listing annotations for the class:

Property	Value	Lang
D code	C12678	
D DEFINITION	<def-source>M...	
D Display_Name	Biliary Tract	
D FULL_SYN	<term-name>Bi...	
D FULL_SYN	<term-name>T...	
D FULL_SYN	<term-name>Bi...	
D FULL_SYN	<term-name>T...	
- Properties and Restrictions:** Lists properties and restrictions for the class:
- ▶ rAnatomic_Structure_Has_Location (multiple Anatomy_Kind) (someValuesFrom Abdomen)
- ▶ rAnatomic_Structure_is_Physical_Part_of (multiple Anatomy_Kind) (someValuesFrom Gas)
- Superclasses:** Shows the class "Organ" as a superclass of "Biliary_Tract".
- Disjoints:** A section for defining disjoint relationships.

At the bottom right, there are radio buttons for "Logic View" and "Properties View", with "Properties View" selected.

A Knowledge-Acquisition Tool

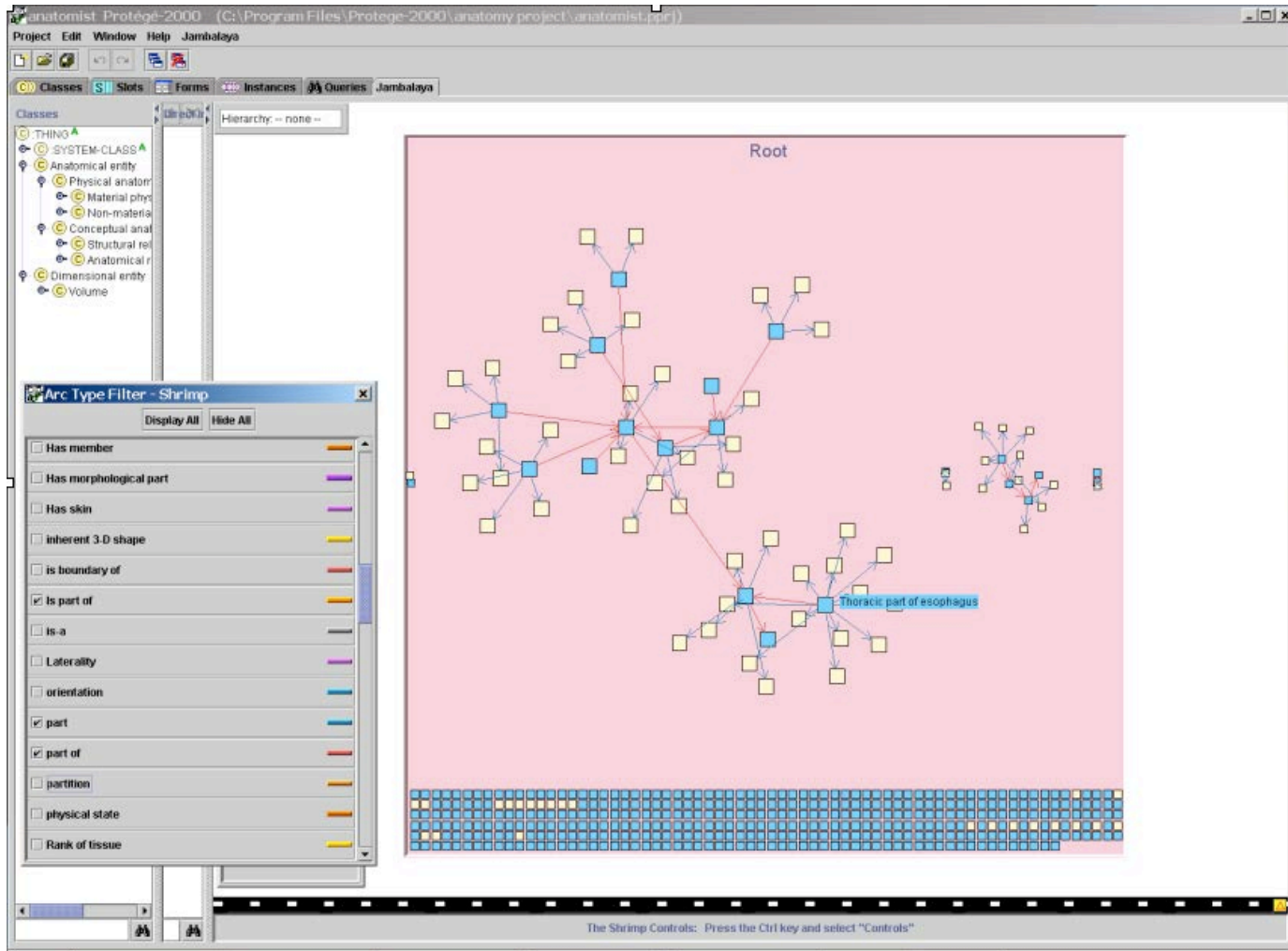
The screenshot displays a software interface for managing Gene Ontology (GO) terms. The interface is divided into several sections:

- Top Navigation:** Includes tabs for 'Classes', 'Slots', 'Forms', 'Instances', and 'Queries'.
- Left Panel (Class Hierarchy):** Shows a tree view of classes. The 'molecular_function' class is selected and highlighted in blue. Other classes include ':THING', ':SYSTEM-CLASS', 'Gene_Ontology_Entity', 'Molecular_Function_Unclassified', 'Biological_Process_Unclassified', 'Cellular_Component_Unclassified', 'biological_process', 'cellular_component', 'Annotation', 'Gene', 'Protein', and 'Transcript'.
- Center Panel (Form):** Displays the details for the 'molecular_function (Gene_Ontology_Metaclass)' class. Fields include:
 - Name:** 0003674
 - Term:** molecular_function
 - Synonyms:** (empty)
 - InterPRO ID:** (empty)
 - Documentation:** (empty)
 - Secondary GOIds:** (empty)
 - EC Numbers:** (empty)
 - TC Numbers:** (empty)
 - SwissProt Keyword:** (empty)
- Right Panel (Definition and Annotations):** Contains text boxes and controls for:
 - Definition:** The action characteristic of a gene product.
 - Definition Reference:** GO:curators
 - Associated Annotations:** (empty)
 - Part-Of:** (empty)
- Bottom Panel (Superclasses):** Shows the 'Gene_Ontology_Entity' class as a superclass of the selected term.

A Platform for Other Applications

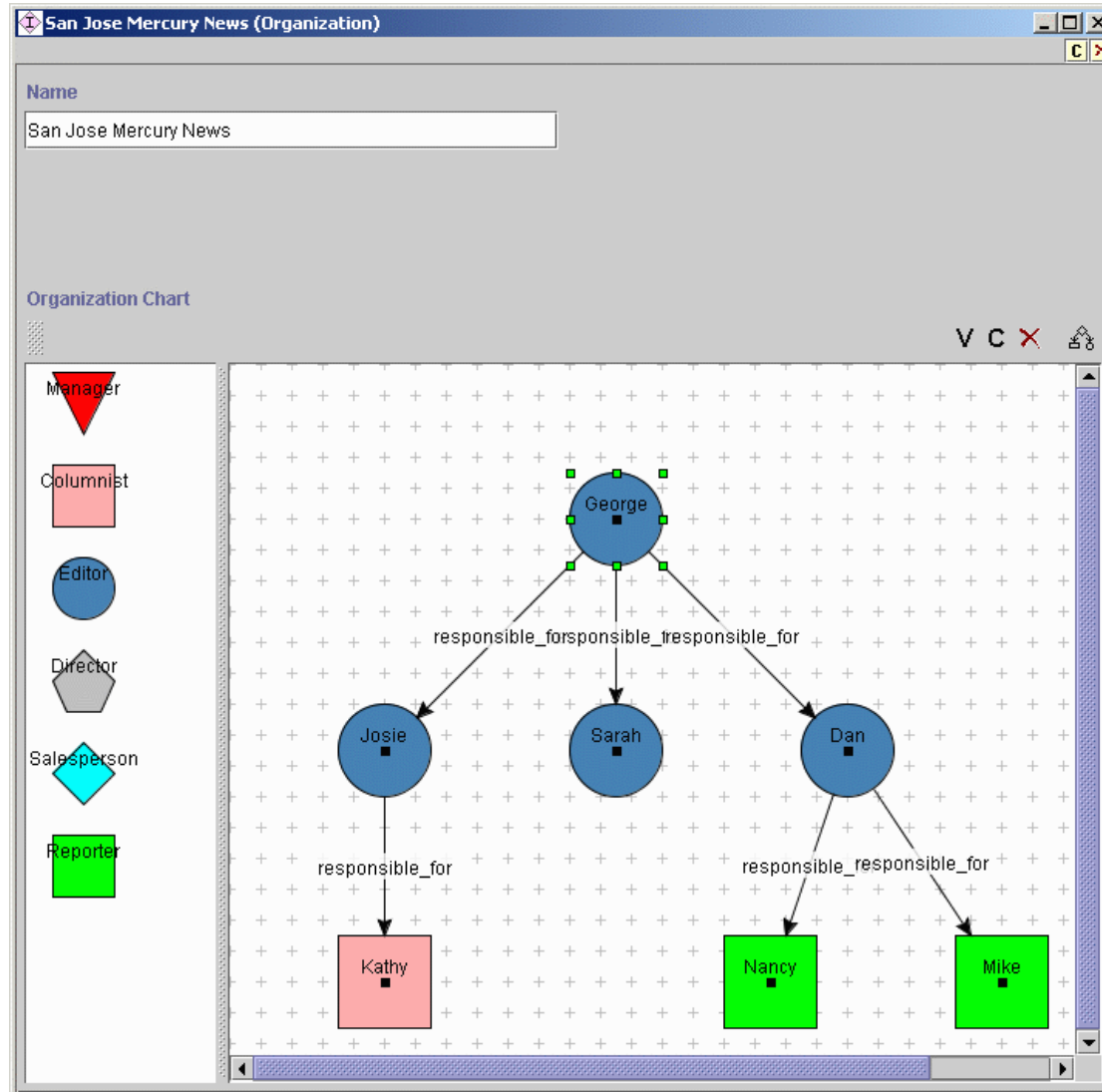
- A **Java API** that enables developers to write plugins for
 - Visualization systems
 - Inferencing systems
 - Scripting facilities
 - Import and export formats
 - User-interface features
 - Means of accessing external data sources
- About **60** plugins currently in the library (developed at Stanford and elsewhere)

Visualization: Jambalaya



Developed at University of Victoria

Graph Widget



Some Applications Supported by Protégé:

- Surveillance of data sources for evidence of potential bioterrorism
- Concept-based information retrieval
- Modeling of metabolic pathways
- Automation of guideline-based therapy

What Makes Protégé Different

- **Easy-to-use** graphical interface
- **Scalability**
 - currently can handle up to 5 million concepts
- **Plugin** architecture
 - active international community of plugin developers
- It's a **platform** for other applications
 - Integration with Eclipse (Mayo Clinic)
 - A server and a client for (Semantic) Web Services
- **Open source**

Foundational Model of Anatomy

- Developed here at UW in the **Department of Biological Structure** by Prof. Cornelius Rosse and Dr. Onard Mejino and other team members (including me)
- Purpose was to provide a comprehensive **framework for the study of the human body**
- Can be used by students, doctors, medical researchers, even the general public

Foundational Model of Anatomy (FMA) Ontology

Motivating Hypothesis

“A sound ontological framework of biological structure (anatomy) provides a logical, comprehensive and efficient framework for organizing all types of information about biological organisms”

Why Anatomy?

Hypothesis 1: Manifestations of health and disease are attributes of **anatomical structures**.

Hypothesis 2: Representation of anatomy should facilitate representation of other domains and interoperability between biomedical domains.

Foundational Model of Anatomy (FMA) Ontology

Theory:

The FMA is a *spatio-structural ontology* of the entities and relations which together form the *phenotypic structure* of a biological organism at all salient *levels of granularity*.

High level Objectives of the FMA theory

Foundational Model of Anatomy

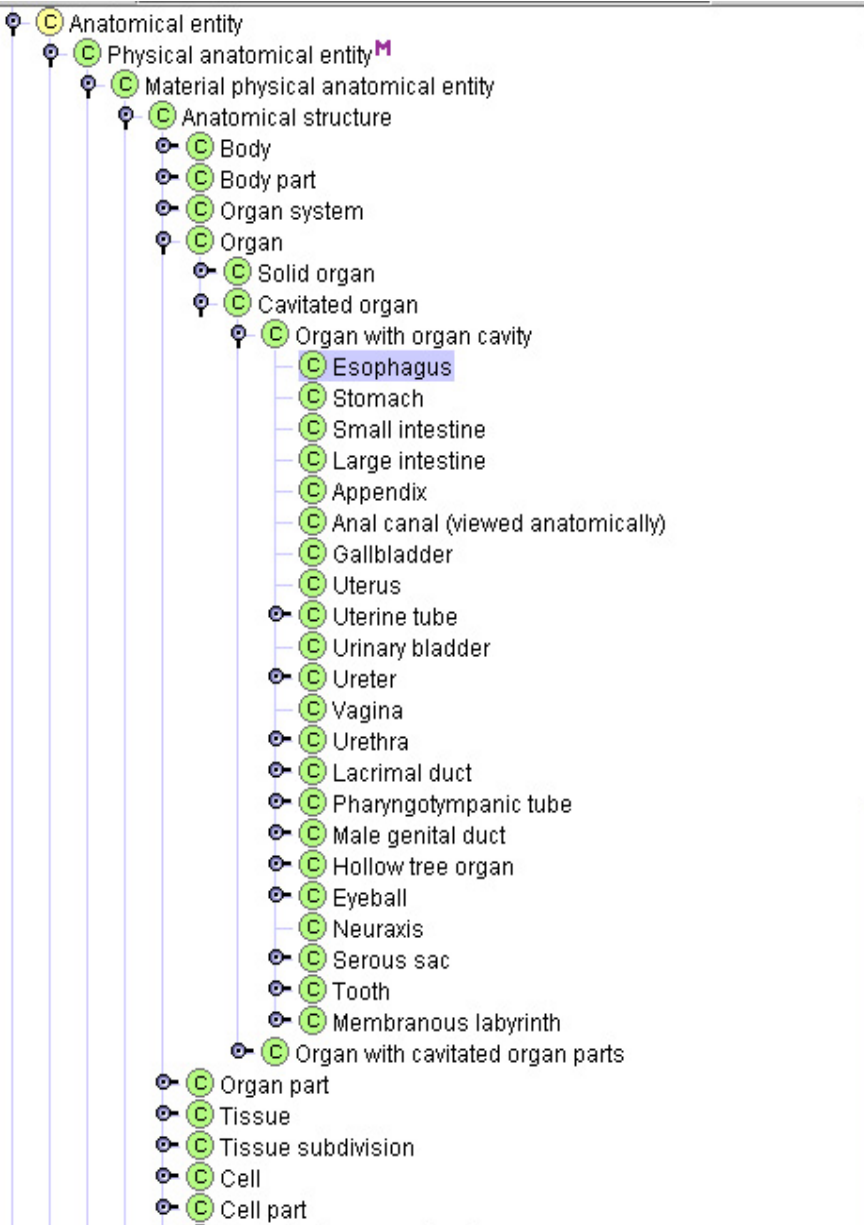
declare the principles

for including entities and relations that are implicitly assumed when knowledge of anatomy is applied in different contexts;

explicitly define

entities and relations necessary and sufficient for consistently representing the structure of a biological organism.

Foundational Model of Anatomy Ontology



Preferred Name **LWDAID**

Synonyms **Non-English Equivalents**

Definition
 Organ with organ cavity which is continuous proximally with the pharynx and distally with the stomach.
 Examples: There is only one esophagus.

Member Of **Part**

- Wall of esophagus
- Lumen of esophagus
- Cervical part of esophagus
- Thoracic part of esophagus
- Abdominal part of esophagus
- Broncho-esophageus

Part Of

Attributed Part

related part	anatomical/arbitrary	shared/unshared	partition
Wall of esophagus	Anatomical	Unshared	Partition 1
Lumen of esophagus	Anatomical	Unshared	Partition 1
Cervical part of esophagus	Arbitrary	Unshared	Partition 2
Thoracic part of esophagus	Arbitrary	Unshared	Partition 2

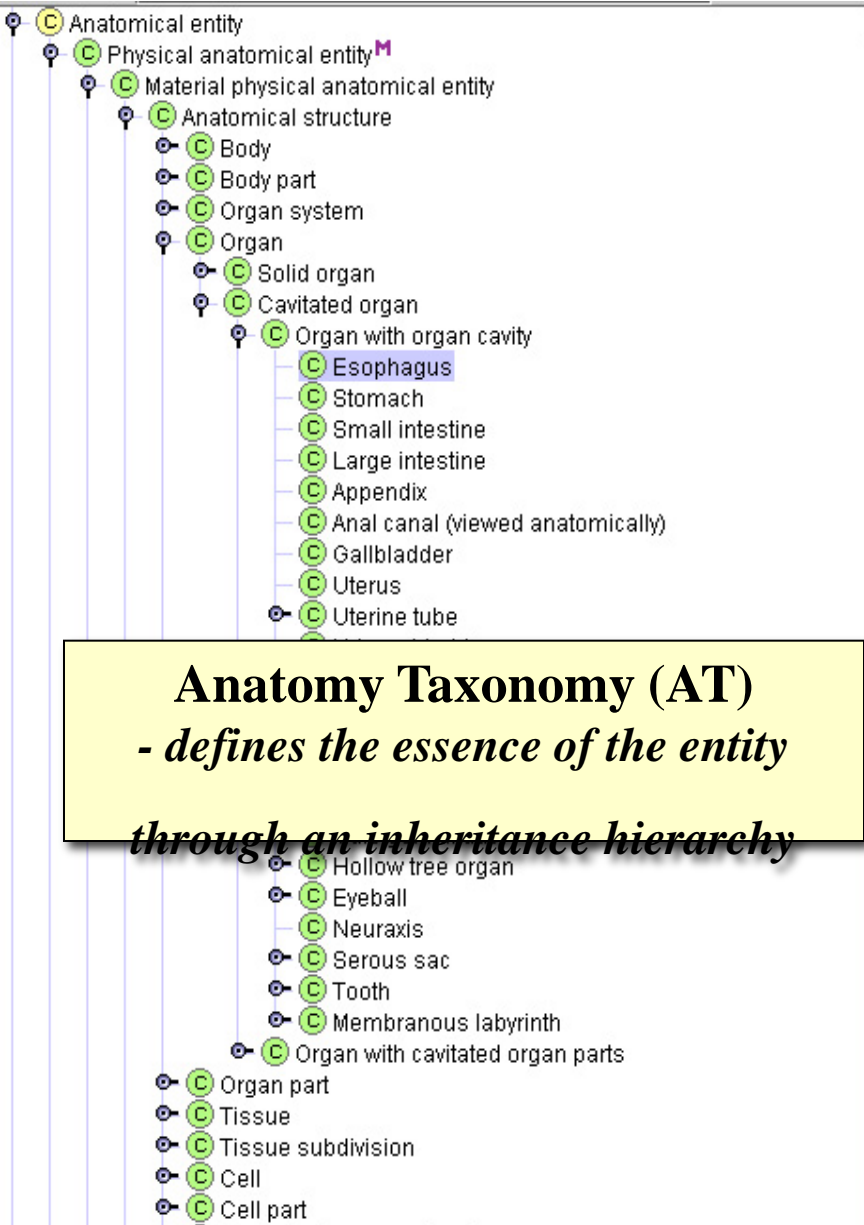
Attributed Continuous With

related object	coordinate	laterality
Pharynx	Superior	
Stomach	Inferior	

Orientation

related object	coordinate	laterality
Plane of pharyngoesophageal junction	Superior	
Plane of esophagogastric junction	Inferior	

Foundational Model of Anatomy Ontology



Anatomy Taxonomy (AT)
- defines the essence of the entity
through an inheritance hierarchy

Preferred Name **UWDAID**

Synonyms **Non-English Equivalents**

Definition
 Organ with organ cavity which is continuous proximally with the pharynx and distally with the stomach.
 Examples: There is only one esophagus.

Member Of **Part**

- Wall of esophagus
- Lumen of esophagus
- Cervical part of esophagus
- Thoracic part of esophagus
- Abdominal part of esophagus
- Broncho-esophageus

Part Of

Attributed Part

related part	anatomical/arbitrary	shared/unshared	partition
Wall of esophagus	Anatomical	Unshared	Partition 1
Lumen of esophagus	Anatomical	Unshared	Partition 1
Cervical part of esophagus	Arbitrary	Unshared	Partition 2
Thoracic part of esophagus	Arbitrary	Unshared	Partition 2

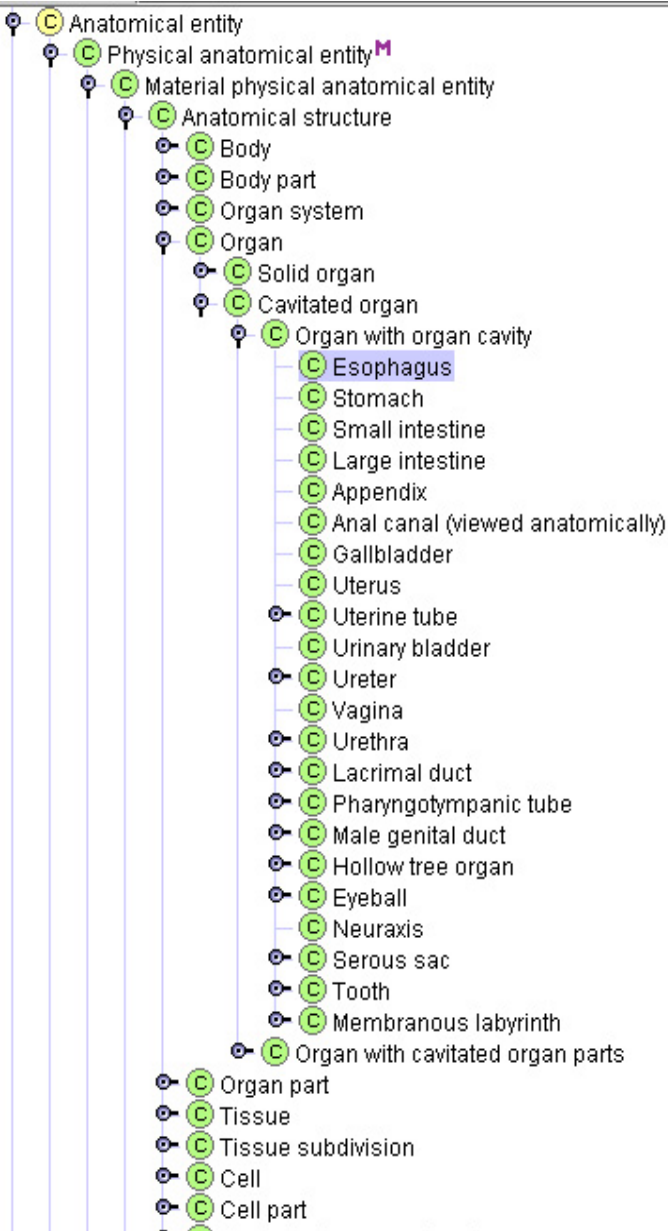
Attributed Continuous With

related object	coordinate	laterality
Pharynx	Superior	
Stomach	Inferior	

Orientation

related object	coordinate	laterality
Plane of pharyngoesophageal junction	Superior	
Plane of esophagogastric junction	Inferior	

Foundational Model of Anatomy Ontology



Preferred Name LWDAID

Synonyms Non-English Equivalents

Anatomical Structural Abstraction (ASA)
- declares the spatio-structural properties of an anatomical entity

Member Of

Part Of

Attributed Part

related part	anatomical/arbitrary	shared/unshared	partition
Wall of esophagus	Anatomical	Unshared	Partition 1
Lumen of esophagus	Anatomical	Unshared	Partition 1
Cervical part of esophagus	Arbitrary	Unshared	Partition 2
Thoracic part of esophagus	Arbitrary	Unshared	Partition 2

Attributed Continuous With

related object	coordinate	laterality
Pharynx	Superior	
Stomach	Inferior	

Orientation

related object	coordinate	laterality
Plane of pharyngoesophageal ju...	Superior	
Plane of esophagogastric junction	Inferior	

Foundational Model of Anatomy Ontology

Unifying theory of anatomy

High Level Scheme

$FMA = (At, ASA, ATA, Mk)$

where:

At = Anatomy taxonomy
ASA = Anatomical Structural Abstraction
ATA = Anatomical Transformation Abstraction
Mk = Metaknowledge
(principles, rules, axioms)

Exploring the FMA

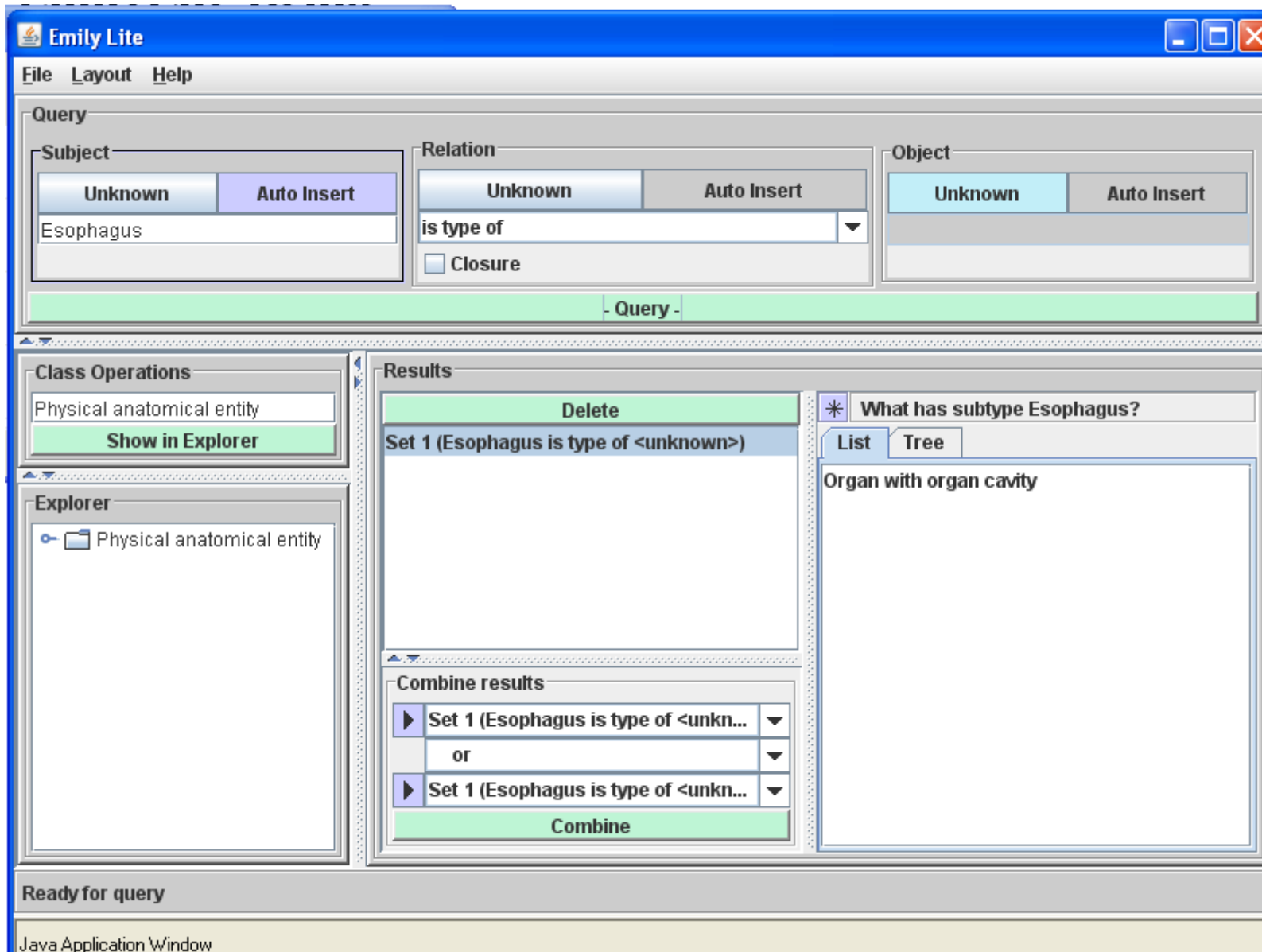
- The Foundational Model Explorer (FME)

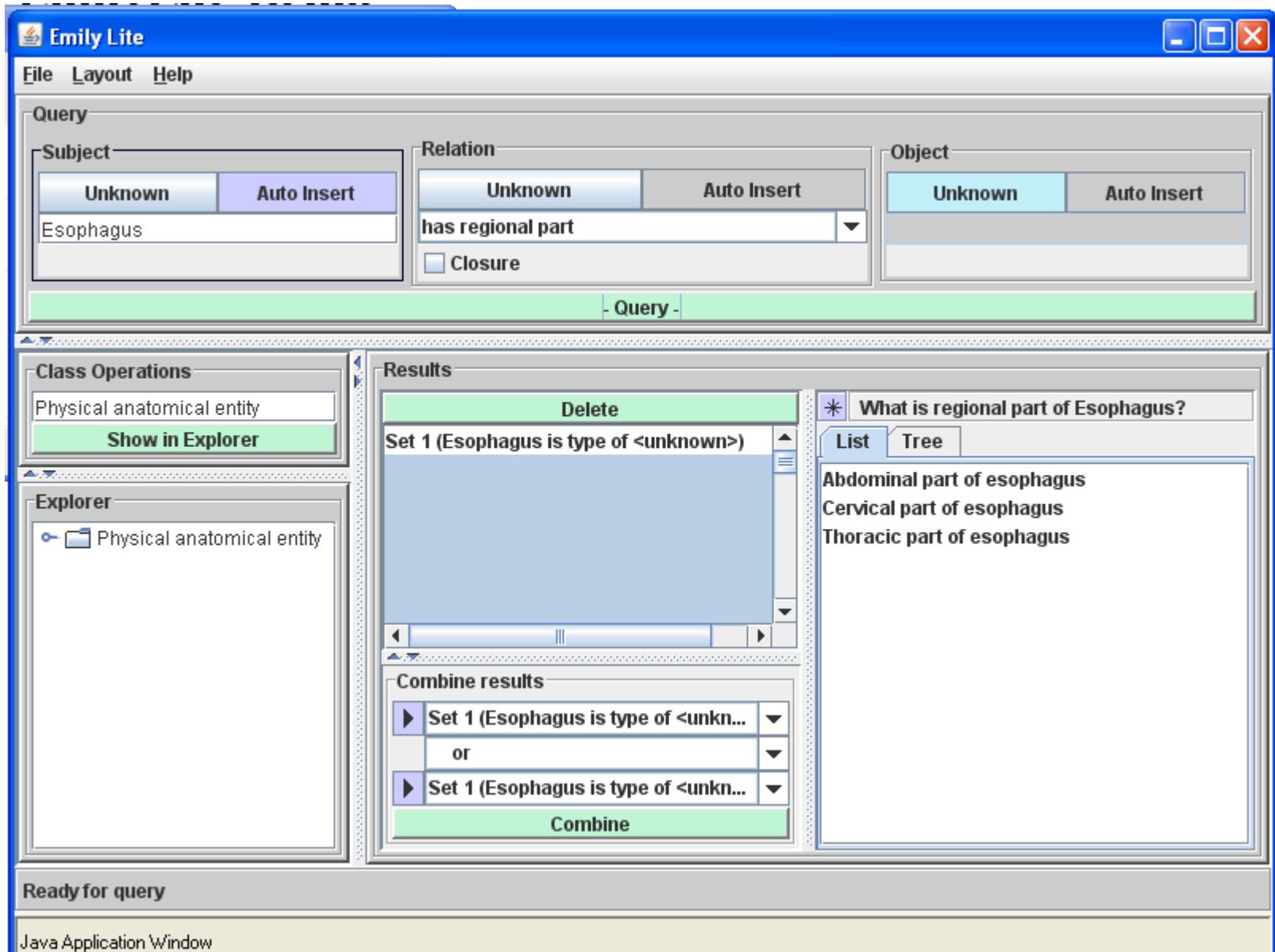
<http://fme.biostr.washington.edu/FME/index.html>

allows browsing through the frames following links.

- Emily Lite

allows relational queries





Emily Lite File Layout Help

Query

Subject: Unknown Auto Insert
Esophagus

Relation: Unknown Auto Insert
has regional part
 Closure

Object: Unknown Auto Insert

- Query -

Class Operations
Physical anatomical entity
Show in Explorer

Explorer
Physical anatomical entity

Results

Delete

Set 1 (Esophagus is type of <unknown>)
Set 2 (Esophagus has regional part <unkno...)

Combine results

Set 1 (Esophagus is type of <unkn...)
or
Set 1 (Esophagus is type of <unkn...)

Combine

*** What (is regional part of)+ Esophagus?**

List Tree

- Abdominal part of esophagus
- C6 segment of esophagus
- C6 segment proper of esophagus
- C7 segment of esophagus
- Cervical part of esophagus
- Esophageal segment of esophagogastric jun
- Esophageal segment of pharyngoesophagea
- T1 segment of esophagus
- T10 segment of esophagus
- T10 segment proper of esophagus
- T2 segment of esophagus
- T3 segment of esophagus

Ready for query

Java Application Window

Emily Lite File Layout Help

Query

Subject
 Unknown Auto Insert
 Esophagus

Relation
 Unknown Auto Insert
 has constitutional part
 Closure

Object
 Unknown Auto Insert

- Query -

Class Operations
 Physical anatomical entity
 Show in Explorer

Explorer
 Physical anatomical entity

Results

Delete

Set 1 (Esophagus is type of <unknown>)
 Set 2 (Esophagus has regional part <unknown>)
 Set 3 (Esophagus (has regional part)+ <unknown>)
 Set 4 (Esophagus (has constitutional part)+ <unknown>)

Combine results

▶ Set 1 (Esophagus is type of <unknown>)
 or
 ▶ Set 1 (Esophagus is type of <unknown>)

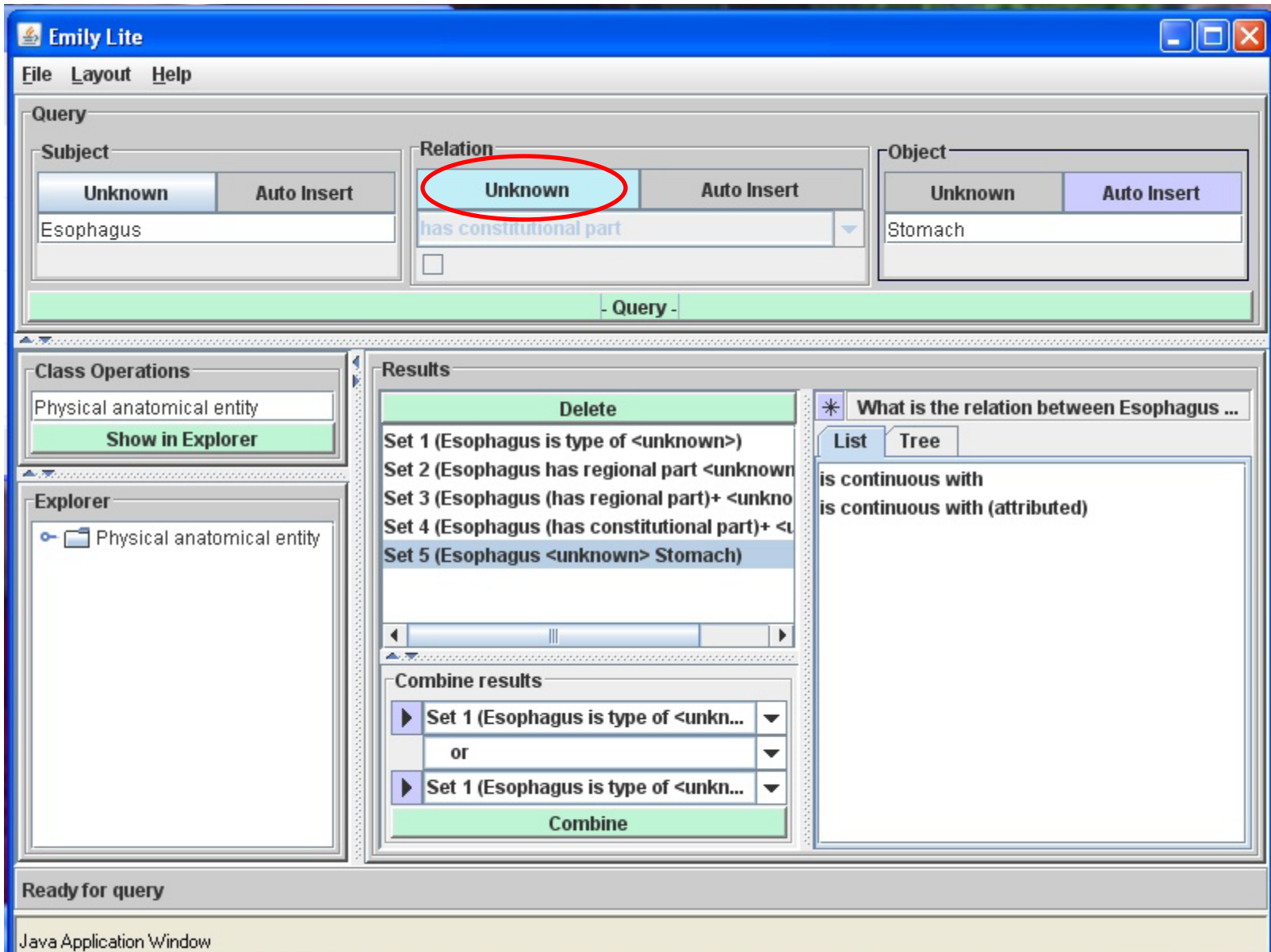
Combine

* What (is constitutional part of)+ Esophagu...
 List Tree

- Adventitia of esophagus
- Basement membrane of epithelium proper of esophagus
- Circular muscle layer of esophagus
- Dense irregular connective tissue of adventitia of esophagus
- Dense regular collagenous tissue of submucosa of esophagus
- Dense regular elastic tissue of lamina propria of esophagus
- Diffuse mucosa-associated lymphoid tissue of esophagus
- Epithelium of esophagus
- Lamina propria of esophagus
- Longitudinal muscle layer of esophagus
- Lumen of esophagus
- Mucosa of esophagus

Ready for query

Java Application Window



Emily Lite File Layout Help

Query

Subject: Unknown Auto Insert
Esophagus

Relation: Unknown Auto Insert
has constitutional part

Object: Unknown Auto Insert
Heart

- Query -

Class Operations
Physical anatomical entity
Show in Explorer

Explorer
Physical anatomical entity

Results

Delete

- Set 1 (Esophagus is type of <unknown>)
- Set 2 (Esophagus has regional part <unknown>)
- Set 3 (Esophagus (has regional part)+ <unknown>)
- Set 4 (Esophagus (has constitutional part)+ <unknown>)
- Set 5 (Esophagus <unknown> Stomach)
- Set 6 (Esophagus <unknown> Heart)

Combine results

- Set 1 (Esophagus is type of <unknown>)
- or
- Set 1 (Esophagus is type of <unknown>)

Combine

What is the relation between Esophagus and Heart?

List Tree

has regional part.is constitutional part of.has constitutional part
has regional part.is constitutional part of.has constitutional part.is nerve supply of
is type of.is type of.has subtype.has subtype
has constitutional part.is type of.has subtype.is constitutional part of

Ready for query

Java Application Window

Auto Insert

Object

Unknown	Auto Insert
Heart	

* What is the relation between Esophagus and Heart?

List **Tree**

Break down complex relations

Show set dependencies

- Esophagus
 - (is type of) Organ with organ cavity
 - (has regional part) Thoracic part of esophagus
 - (is constitutional part of) Content of mediastinum
 - (is constitutional part of) Content of superior mediastinum
 - (is constitutional part of) Intrathoracic part of chest
 - (has constitutional part) Heart
- (has constitutional part) Wall of esophagus
 - (is type of) Wall of organ
 - (has subtype) Wall of heart
 - (is constitutional part of) Heart