# PODD - Towards An EXTENSIBLE, DOMAIN-AGNOSTIC SCIENTIFIC DATA MANAGEMENT SYSTEM

YUAN-FANG LI (liyf@itee.uq.edu.au)

ERESEARCH LAB, SCHOOL OF ITEE
THE UNIVERSITY OF QUEENSLAND, AUSTRALIA

# PODD - TOWARDS AN EXTENSIBLE, DOMAIN-AGNOSTIC SCIENTIFIC DATA MANAGEMENT SYSTEM

AN ONTOLOGY-DRIVEN APPROACH IN A PHENOMICS SETTING

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  - High-throughput & high-resolution processes
- Data evolution
  - Changes in model & data

# DATA MANAGEMENT REQUIREMENTS

### DATA MANAGEMENT REQUIREMENTS

- Collection
- Distribution & sharing
- Access control
- Archival & versioning
- Discovery & analysis
- Repurposing

### PODD GOALS

AN EXTENSIBLE & DOMAIN-INDEPENDENT DATA MANAGEMENT ARCHITECTURE

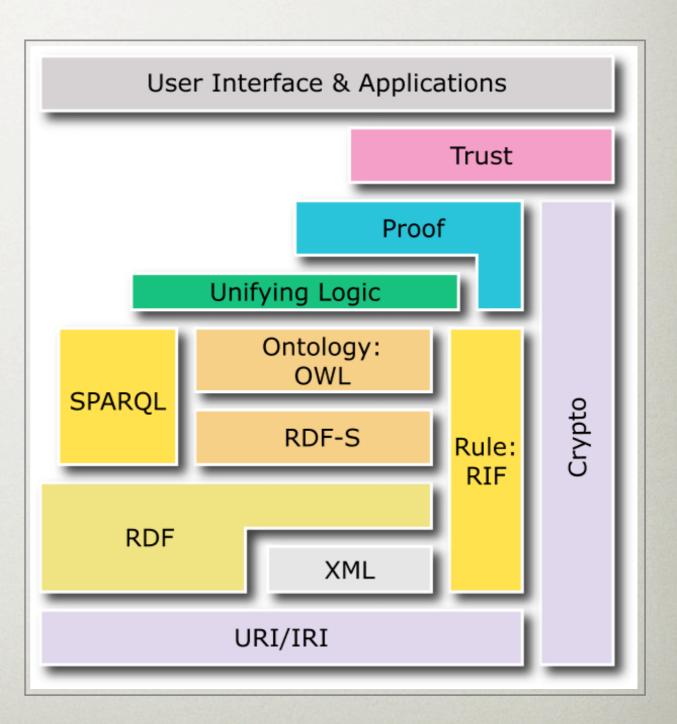
### RELATED MODELS & SYSTEMS

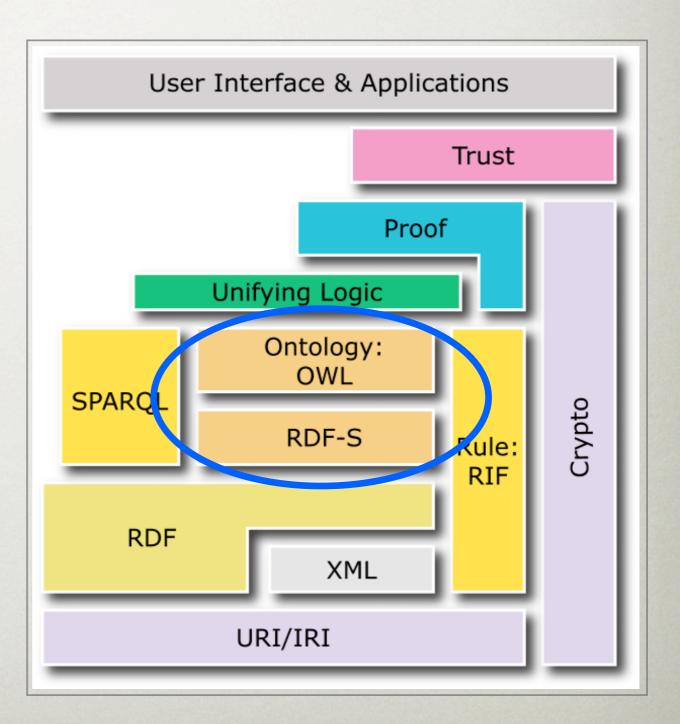
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  - Functional Genomics Experiment Model (FuGe)
    - UML & database based
  - Ontology for Biomedical Investigations (OBI)
    - 26,000+ OWL classes & 10,000+ axioms

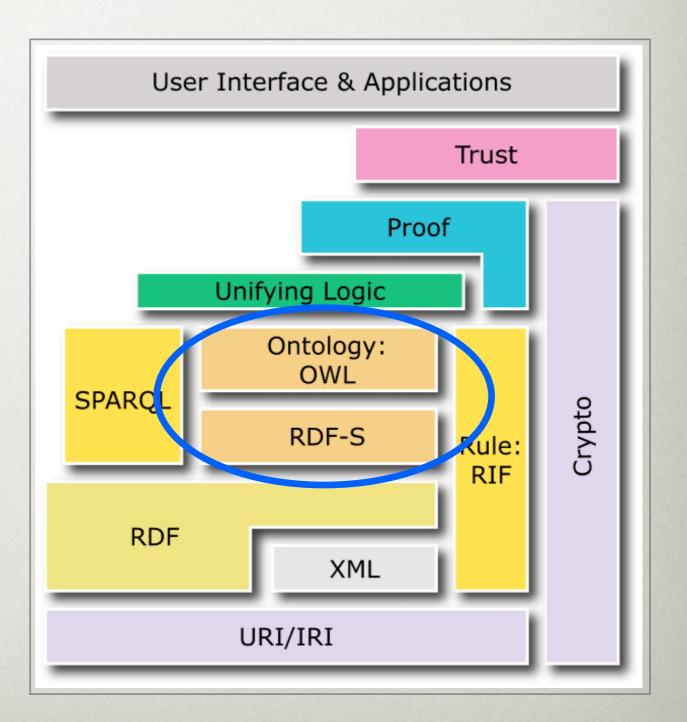
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- Systems
  - VIVO: ontology-based institutional research repository
  - PhonemicDB: a multi-organism phenotype-genotype database
  - Fedora Commons, Apache JackRabbit: digital content repository systems



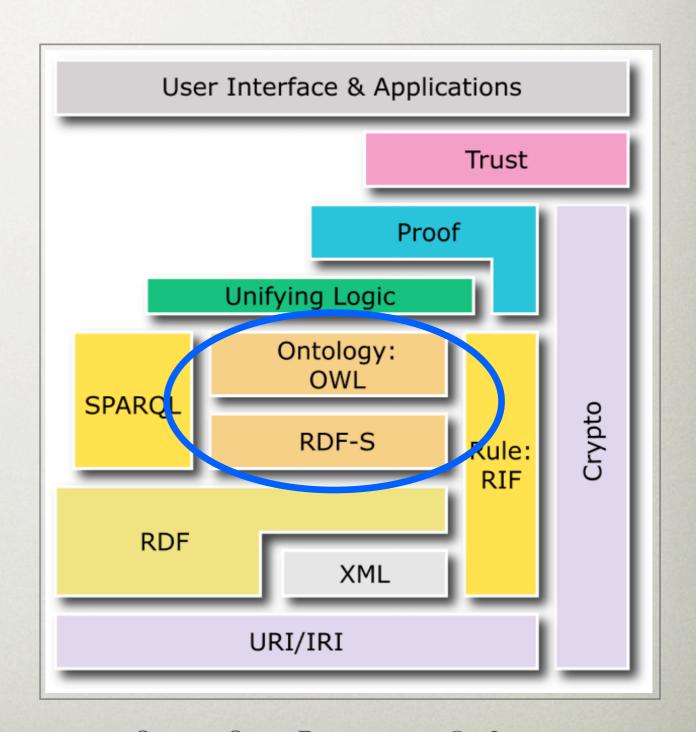


- Expressed in OWL (& RDF Schema)
  - Provides syntax & semantics enables reasoning
  - Expressivity vs decidability

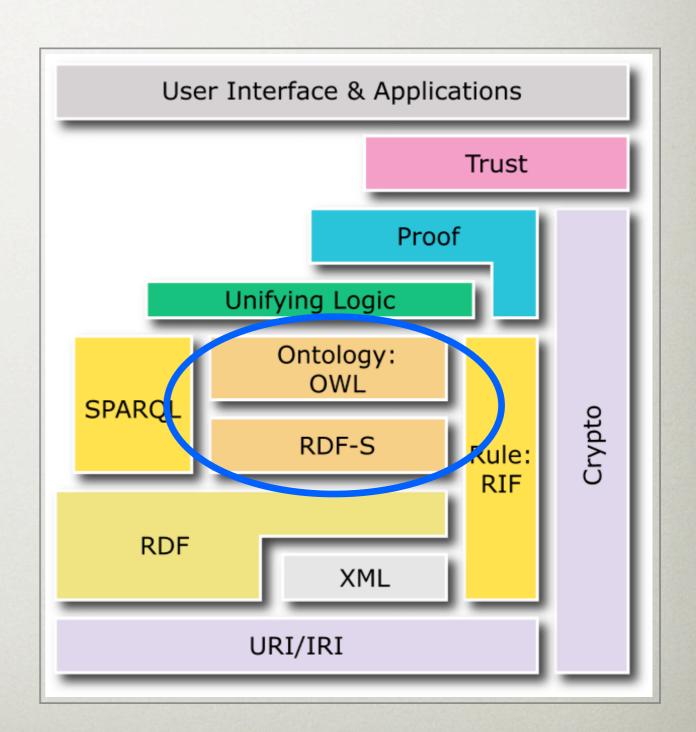


Source: Steve Bratt, <steve@w3.org>

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  - Provides syntax & semantics enables reasoning
  - Expressivity vs decidability
- Designed to be open & interoperable
  - Facilitates sharing, reuse & integration
- Maturing technology stacks
  - APIs, reasoners, triple stores, query engines



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- Basics: ontologies as domain models for scientific experiments data
  - Domain-independent & domain-specific ontologies

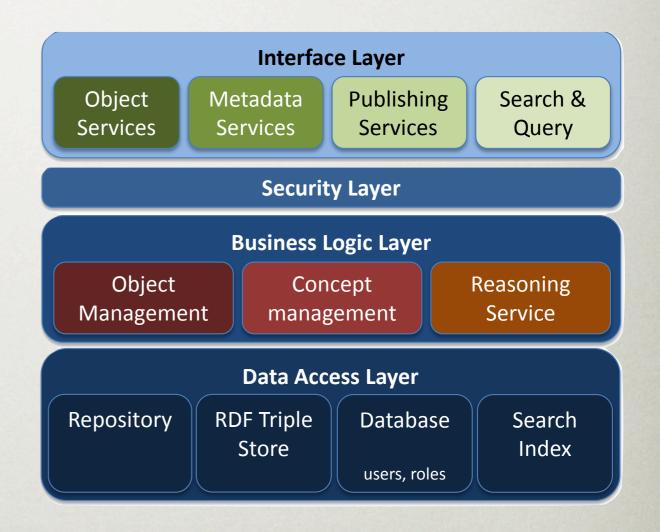
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- Ontologies & RDF central to all operations in the data lifecycle
- Aims: improved extensibility & data integration

### THE PODD SYSTEM ARCHITECTURE

- PODD: Phenomics Ontology
   Driven Data System
- Ontologies the core of the architecture
- Objects represented semantically
  - Semantics (metadata) captured in RDF
- Repository operations on RDF:
  - Ingestion, retrieval, update, query & search, export



- Extensibility through inheritance & versioning
- Integration through ontology alignment/ mapping

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Domain concepts	OWL classes
Attributes & relations	OWL restrictions
Domain objects	OWL individuals
Comments, descriptions	OWL/RDF annotations

Models scientific experiments

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- Organizes data logically
  - Represented as metadata objects
  - Parent-child relationships
  - References relationships

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- Phenomics ontology: domain specific

#### BASE

```
PODDConcept \sqsubseteq \top
\top \sqsubseteq \forall contains.PODDConcept
isContainedBy \sqsubseteq (\neg contains)
PODDConcept \sqsubseteq \leq 1 \ isContainedBy
\top \sqsubseteq refersTo.PODDConcept
```

```
Project \sqsubseteq = 1 \; hasProjectPlan \; \sqcap
\geq 1 \; hasInvestigation \; \sqcap
= 1 hasStartDate \; \sqcap
\leq 1 \; hasPublicationDate \; \sqcap
```

#### BASE

#### **PHENOMICS**

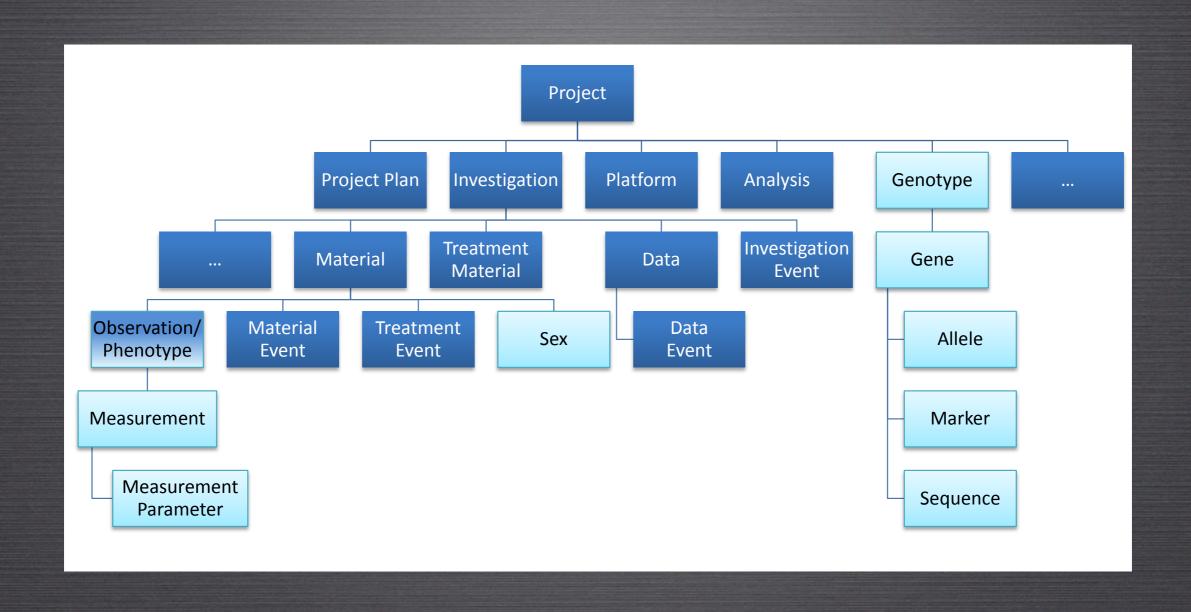
 $PODDConcept \sqsubseteq \top$   $\top \sqsubseteq \forall \ contains.PODDConcept$   $isContainedBy \sqsubseteq (\neg \ contains)$  $PODDConcept \sqsubseteq \leq 1 \ isContainedBy$ 

 $\top \sqsubseteq refersTo.PODDConcept$ 

 $Genotype \sqsubseteq PODDConcept \sqcap \ orall hasGene. Gene \sqcap \ \leq 1 \ hasEcotype \sqcap \ \leq 1 \ hasSubspecies \sqcap$ 

 $Project \sqsubseteq = 1 \; hasProjectPlan \; \sqcap$   $\geq 1 \; hasInvestigation \; \sqcap$   $= 1hasStartDate \; \sqcap$   $\leq 1 \; hasPublicationDate \; \sqcap$ 

 $Project \sqsubseteq \ \forall \ hasGenotype.Genotype \ \sqcap$   $Material \sqsubseteq \ \forall \ hasPhenotype.Phenotype \ \sqcap$   $\ \forall \ refersToGenotype.Genotype$ 



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  - OWLAPI, Pellet, Fedora Commons, Sesame, Lucene & Solr, etc.

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- Facilitates extensibility & evolution
  - Ontology reasoning instead of DB integrity constraint checking
  - Data & metadata are all versioned
- System exploration
  - Search, browsing, SPARQL querying, etc.

#### CONCLUSION

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  - ✓ An ontology-driven architecture for improving extensibility
  - √ A set of ontologies as domain models
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  - √ A set of ontologies as domain models
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- ? Future works
  - ? Ontology/vocabulary mapping
  - ? Annotation of domain objects
  - ? Workflow support

• Co-authors: Gavin Kennedy, Faith Davies, Jane Hunter (UQ)

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