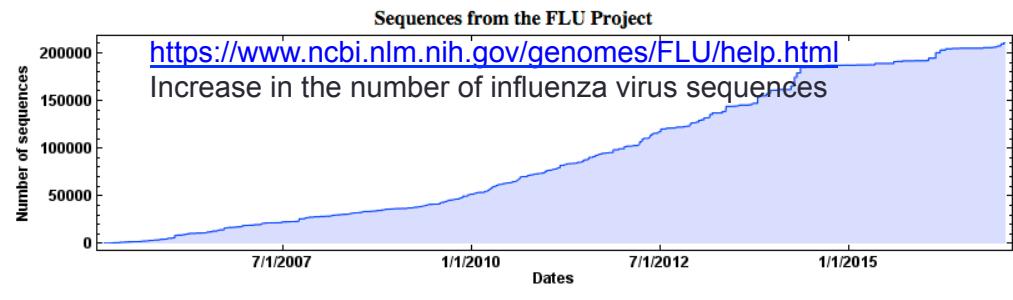
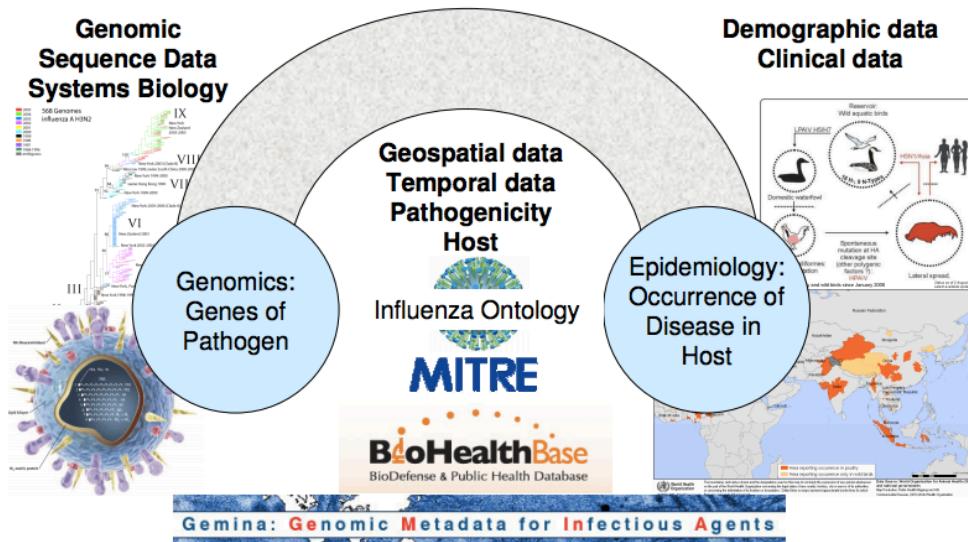


# Influenza Ontology



## Research Question: Bridging the Gap - Connecting Genomics and Epidemiology



Developed to support two use cases:

- Influenza Research
- Influenza Surveillance

Bridging the gap between

- genomic (molecular level)
- epidemiology level, (population level)

The ontology links “bird” and “avian” and the name of the species and type (duck, chicken”). This enables queries over GenBank data for “all cases of avian influenza.”

# Influenza Ontology

## Development Process

- Identify the right **collaborators**
- Collect **metadata terms**
- Identify **resources** for that include these terms
- **Regularize** metadata
  - Generate a **controlled vocabulary** (terms)
- **Validate** subset with BioHealthBase CEIRS data
- **Iterate, review** with community, publish
- **Integrate** (deploy) Influenza ontology into workflow



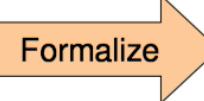
## Influenza Ontology First Draft

Status: We have just started the formalization step.

### Initial steps:

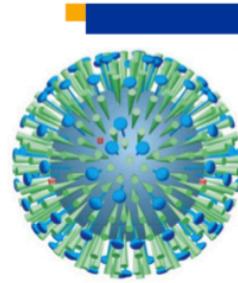
- Collect metadata terms
- Map and align terms
- Group related information
- Identify and define relationships
- Identify external ontologies

Category 1	Category 2	Category 3	Category 4	Category 5
Data 1	Data 2	Data 3	Data 4	Data 5
Data 6	Data 7	Data 8	Data 9	Data 10
Data 11	Data 12	Data 13	Data 14	Data 15
Data 16	Data 17	Data 18	Data 19	Data 20



Excel Spreadsheet

MITRE



### Formalize:

- Normalize terms into a CV
- Issue unique identifiers
- Instantiate class hierarchy
- Define properties and values
- Link to external ontology terms

The OBO-Edit interface displays a hierarchical tree of ontology terms on the left, with a detailed view of term properties and definitions on the right. A large orange arrow points from the 'Formalize' section above to this screen.

OBO-Edit: Ontology Editing Tool

9

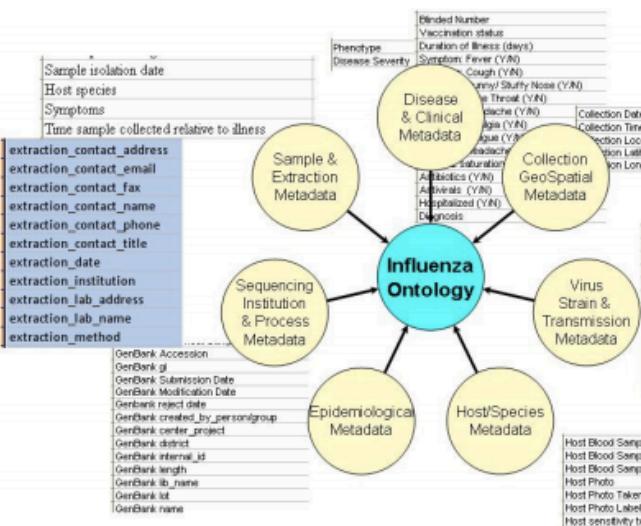
# Application Ontology

## Influenza Ontology



## Reuse of existing ontologies & metadata standards

200 controlled vocabulary terms covering several fields



**OBI – Ontology of Biomedical Investigations**

**EnvO – Environmental Ontology (habitat of pathogen)**

**GAZ – Gazetteer (geographic locations)**

**FMA – Foundational Model of Anatomy**

**DC – Dublin Core (publication metadata)**

**PATO – Phenotype**

**SO – Sequence Ontology (sequence features)**

**Cell – Cell Ontology (types of cells)**

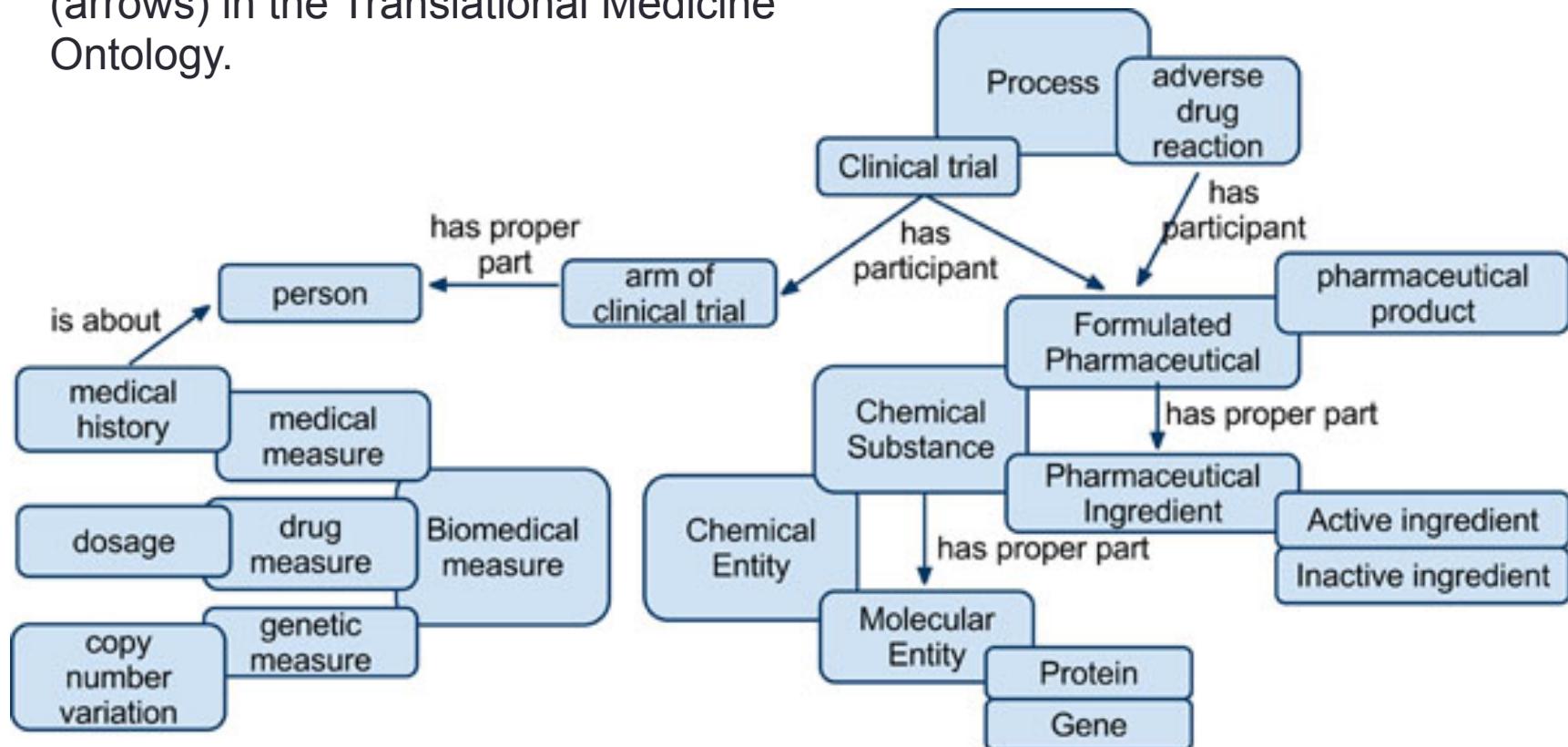
**DO – Disease Ontology**

**IDO – Infectious Disease Ontology**

# Translational Medicine Ontology



Overview of selected types, subtypes (overlap) and existential restrictions (arrows) in the Translational Medicine Ontology.



The Translational Medicine Ontology and Knowledge Base: driving personalized medicine by bridging the gap between bench and bedside

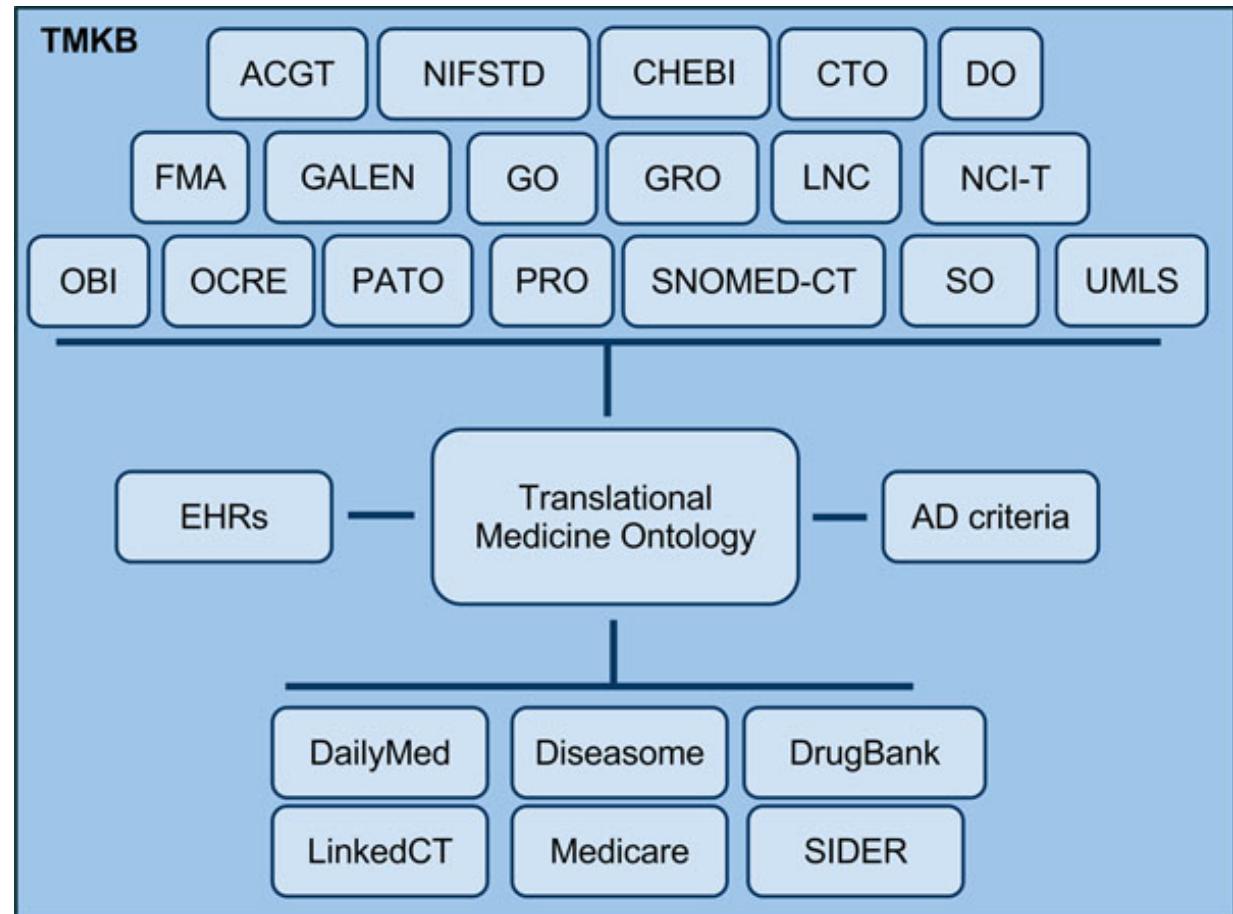
Luciano et al. Journal of Biomedical Semantics 2011, 2(Suppl 2):S1 <http://www.jbiomedsem.com/content/2/S2/S1>

# Translational Medicine Knowledge Base



Translational  
Medicine Ontology  
with mappings to  
ontologies and  
terminologies listed  
in the NCBO  
BioPortal.

The TMO provides a global schema for Indivo-based electronic health records (EHRs) and can be used with formalized criteria for Alzheimer's Disease. The TMO maps types from Linking Open Data sources.



# Best Practices



## Semantic Web Methodology & Technology Development Process

