

# Machine learning with biomedical ontologies

*Presented by Sarah Alghamdi, Azza Althagafi, Robert Hoehndorf, Maxat Kulmanov, Sumyyah Toonsi, Fernando Zhapa-Camacho*

# Learning Outcomes

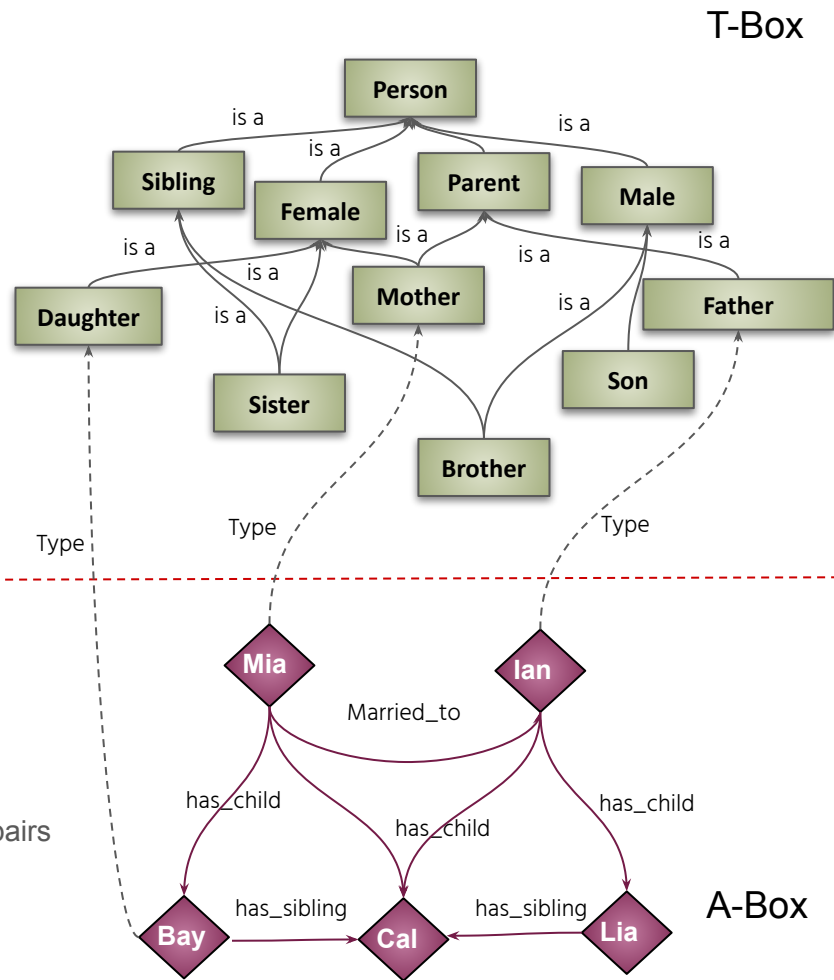
- Introduce Ontologies and the description logic
- Discuss unsupervised machine learning methods that can “embed” from one structure to another
- Introduce different methods that use ontologies in machine learning models
- Introduce mOWL, a software library for machine learning with ontologies
- Incorporate mOWL in Biomedical data analysis using different approaches

# Preliminaries: ontologies

- “An ontology is an **explicit specification** of a **conceptualization**”  
... [Gruber 1993](#)
- “An ontology is an explicit **formal** specification of a **shared** conceptualization” ... [Borst 1997](#)
- “An ontology is a **logical theory** designed in order to capture the **intended models** corresponding to a certain conceptualization and to **exclude the unintended ones**” ... [Guarino 2009](#)

# Preliminaries: ontologies

- Ontology consist of :  
 $O=\{C,R,I,I,-\}$
- T-Box
  - Set of terminological Component
- A-Box
  - Set of assertions using T-Box
    - Concept assertions
    - Relation Assertion
- Metadata
  - representation of a resource in terms of attribute name-value pairs
    - Definition
    - Labels
    - ...



# Preliminaries: ontologies

- Description Logic (DL) is used to formally and explicitly represent ontologies

Name	DL syntax	Semantics
Top concept	$\top$	$\Delta^{\mathcal{I}}$
Bottom concept	$\perp$	$\emptyset$
Concept	$C$	$C^{\mathcal{I}} \subseteq \Delta^{\mathcal{I}}$
Concept disjunction	$C_1 \sqcup C_2$	$C_1^{\mathcal{I}} \cup C_2^{\mathcal{I}}$
Concept conjunction	$C_1 \sqcap C_2$	$C_1^{\mathcal{I}} \cap C_2^{\mathcal{I}}$
Concept negation	$\neg C$	$\Delta^{\mathcal{I}} \setminus C^{\mathcal{I}}$
Universal restriction	$\forall R.C$	$\{x \in \Delta^{\mathcal{I}} \mid \forall y \in \Delta^{\mathcal{I}} ((x, y) \in R^{\mathcal{I}} \wedge y \in C^{\mathcal{I}})\}$
Existential restriction	$\exists R.C$	$\{x \in \Delta^{\mathcal{I}} \mid \exists y \in \Delta^{\mathcal{I}} ((x, y) \in R^{\mathcal{I}} \rightarrow y \in C^{\mathcal{I}})\}$
Subclass of	$C_1 \sqsubseteq C_2$	$C_1^{\mathcal{I}} \subseteq C_2^{\mathcal{I}}$
Subproperty of	$R_1 \sqsubseteq R_2$	$R_1^{\mathcal{I}} \subseteq R_2^{\mathcal{I}}$
Equivalent class	$C_1 \equiv C_2$	$C_1^{\mathcal{I}} = C_2^{\mathcal{I}}$
Equivalent property	$R_1 \equiv R_2$	$R_1^{\mathcal{I}} = R_2^{\mathcal{I}}$

## Concepts , Roles

*Person*  $\sqsubseteq \top$

*Female*  $\sqcap$  *Male*  $\sqsubseteq \perp$

*Female*  $\sqcup$  *Male*  $\sqsubseteq \top$

*Female*  $\equiv \neg$  *Male*

*Parent*  $\equiv \exists$  *has\_child*. *Person*

*Son*  $\sqsubseteq$  *Male*  $\sqcap \exists$  *child\_of*. *Person*

*Mother*  $\sqsubseteq$  *Female*  $\sqcap$  *Parent*

*Sibling*  $\sqsubseteq \exists$  *has\_sibling*. *Person*

*has\_brother*  $\sqsubseteq$  *has\_sibling*

# Preliminaries: ontologies

- Description Logic (DL) is used to formally and explicitly represent ontologies

DL Syntax	Manchester Syntax
$C \sqcap D$	C and D
$C \sqcup D$	C or D
$\neg C$	not C
$\exists R.C$	R some C
$\forall R.C$	R only C
$(\geq nR.C)$	R min n C
$(\leq nR.C)$	R max n C
$(= nR.C)$	R exactly n C
$\{a\} \sqcup \{b\} \sqcup \dots$	{a b ...}

## Concepts , Roles

*Person*  $\sqsubseteq \top$

*Female*  $\sqcap$  *Male*  $\sqsubseteq \perp$

*Female*  $\sqcup$  *Male*  $\sqsubseteq \top$

*Female*  $\equiv \neg$  *Male*

*Parent*  $\equiv \exists$  *has\_child*. *Person*

*Son*  $\sqsubseteq$  *Male*  $\sqcap \exists$  *child\_of*. *Person*

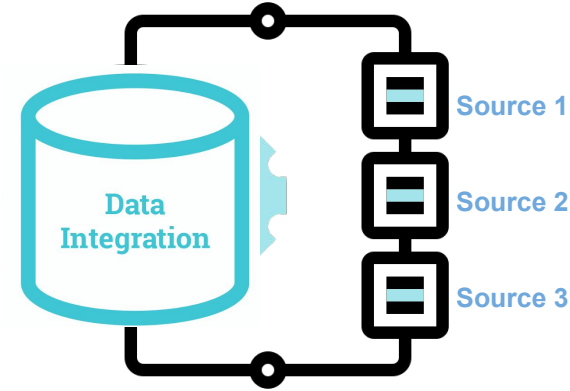
*Mother*  $\sqsubseteq$  *Female*  $\sqcap$  *Parent*

*Sibling*  $\sqsubseteq \exists$  *has\_sibling*. *Person*

*has\_brother*  $\sqsubseteq$  *has\_sibling*

# How Ontologies are used in Databases

- Annotations and data integration
  - Ontologies play a crucial role in facilitating data integration across databases due to their usage of standard identifiers for classes and relations



# How Ontologies are used in Databases

- Annotations and data integration

## GAF fields

The annotation flat file format is comprised of 17 tab-delimited fields.

Column	Content	Required?	Cardinality	Example
1	<a href="#">DB</a>	required	1	UniProtKB
2	<a href="#">DB Object ID</a>	required	1	P12345
3	<a href="#">DB Object Symbol</a>	required	1	PHO3
4	<a href="#">Qualifier</a>	required	1 or 2	NOTInvolved_in
5	<a href="#">GO ID</a>	required	1	GO:0003993
6	<a href="#">DB:Reference (IDB:Reference)</a>	required	1 or greater	PMID:2676709
7	<a href="#">Evidence Code</a>	required	1	IMP
8	<a href="#">With (or) From</a>	optional	0 or greater	GO:0000346
9	<a href="#">Aspect</a>	required	1	F
10	<a href="#">DB Object Name</a>	optional	0 or 1	Toll-like receptor 4
11	<a href="#">DB Object Synonym (ISynonym)</a>	optional	0 or greater	hTollTollbooth
12	<a href="#">DB Object Type</a>	required	1	protein
13	<a href="#">Taxon(Itaxon)</a>	required	1 or 2	taxon:9606
14	<a href="#">Date</a>	required	1	20090118
15	<a href="#">Assigned By</a>	required	1	SGD
16	<a href="#">Annotation Extension</a>	optional	0 or greater	part_of(CL:0000576)
17	<a href="#">Gene Product Form ID</a>	optional	0 or 1	UniProtKB:P12345-2



# How Ontologies are used in Databases

## ● Annotations and data integration

1.	UniProtKB	1.	MGI
2.	A0A024RBG1	2.	MGI:1913300
3.	NUDT4B	3.	0610009B22Rik
4.	enables	4.	enables
5.	GO:0003723	5.	GO:0001222
6.	GO_REF:0000043	6.	MGI:MGI:4834177 GO_REF:0000096
7.	IEA	7.	ISO
8.	UniProtKB-KW:KW-0694	8.	UniProtKB:P0DI82
9.	F	9.	F
10.	Diphosphoinositol polyphosphate phosphohydrolase	10.	RIKEN cDNA 0610009B22 gene
11.	NUDT4B	11.	protein_coding_gene
12.	NUDT4B	12.	taxon:10090
13.	Protein	13.	20210709
14.	taxon:9606      20221109	14.	MGI
15.	UniProt	15.	
16.		16.	
17.		17.	

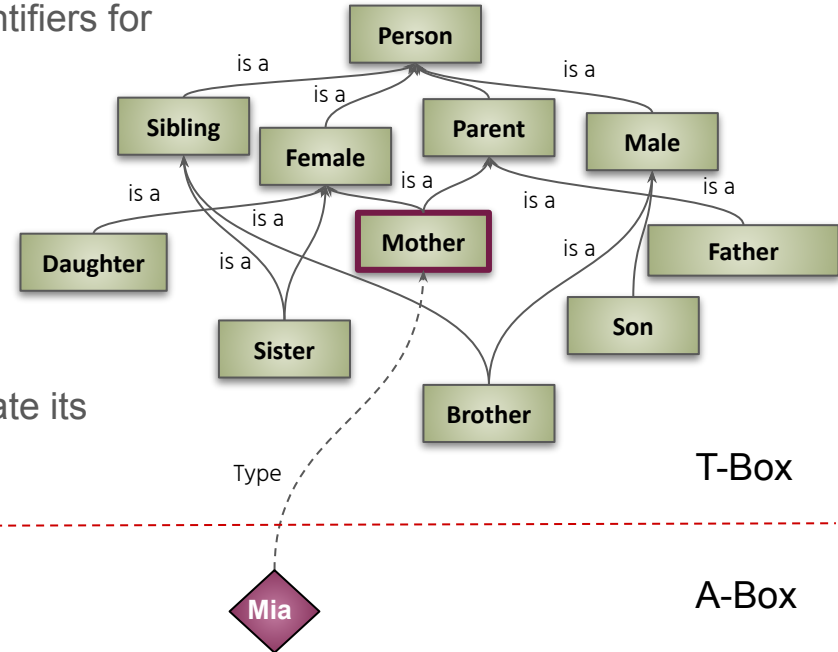
## GAF fields

The annotation flat file format is comprised of 17 tab-delimited fields.

Column	Content	Required?	Cardinality	Example
1	<a href="#">DB</a>	required	1	UniProtKB
2	<a href="#">DB Object ID</a>	required	1	P12345
3	<a href="#">DB Object Symbol</a>	required	1	PHO3
4	<a href="#">Qualifier</a>	required	1 or 2	NOTInvolved_in
5	<a href="#">GO ID</a>	required	1	GO:0003993
6	<a href="#">DB:Reference (IDB:Reference)</a>	required	1 or greater	PMID:2676709
7	<a href="#">Evidence Code</a>	required	1	IMP
8	<a href="#">With (or) From</a>	optional	0 or greater	GO:0000346
9	<a href="#">Aspect</a>	required	1	F
10	<a href="#">DB Object Name</a>	optional	0 or 1	Toll-like receptor 4
11	<a href="#">DB Object Synonym (ISynonym)</a>	optional	0 or greater	hToll      Tollbooth
12	<a href="#">DB Object Type</a>	required	1	protein
13	<a href="#">Taxon(ITaxon)</a>	required	1 or 2	taxon:9606
14	<a href="#">Date</a>	required	1	20090118
15	<a href="#">Assigned By</a>	required	1	SGD
16	<a href="#">Annotation Extension</a>	optional	0 or greater	part_of(CL:0000576)
17	<a href="#">Gene Product Form ID</a>	optional	0 or 1	UniProtKB:P12345-2

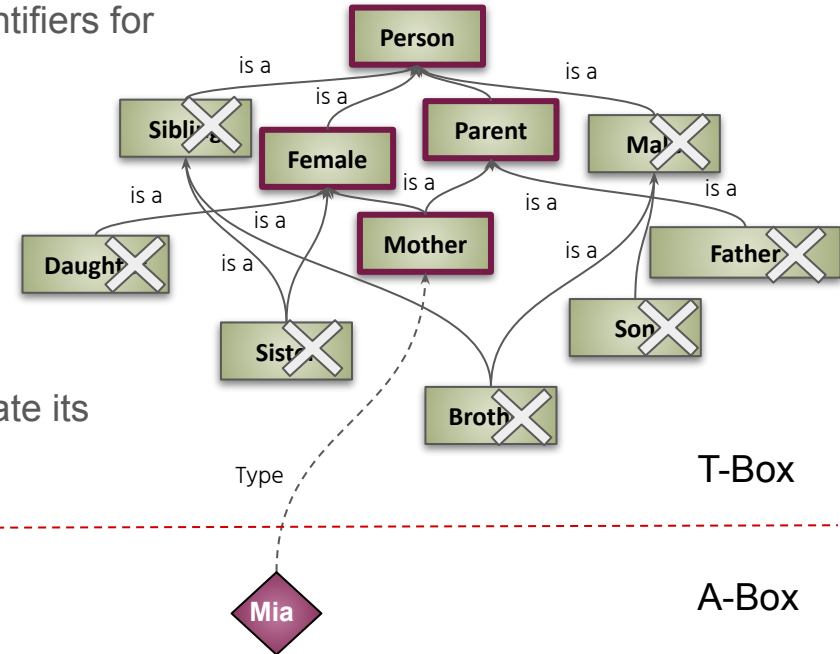
# How Ontologies are used in Databases

- Annotations and data integration
  - Ontologies play a crucial role in facilitating data integration across databases due to their usage of standard identifiers for classes and relations
- True path rule:
  - Annotation for a class is passed to its ancestors
  - Unannotated entities for a class is not used to annotate its descendants



# How Ontologies are used in Databases

- Annotations and data integration
  - Ontologies play a crucial role in facilitating data integration across databases due to their usage of standard identifiers for classes and relations
- True path rule:
  - Annotation for a class is passed to its ancestors
  - Unannotated entities for a class is not used to annotate its descendants

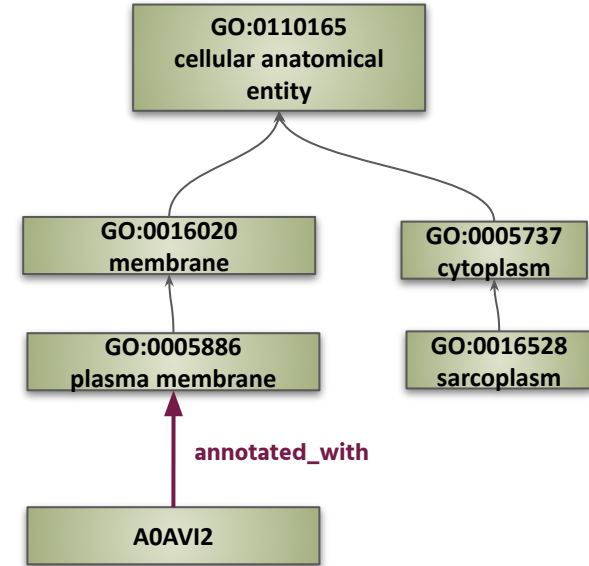


# Transforming GO Annotations to OWL axioms

Example:

Annotating protein **A0AVI2** To Gene Ontology

- Annotations to T-Box
  - A0AVI2**  $\sqsubseteq \exists$  *annotated\_with*. **GO:0005886**

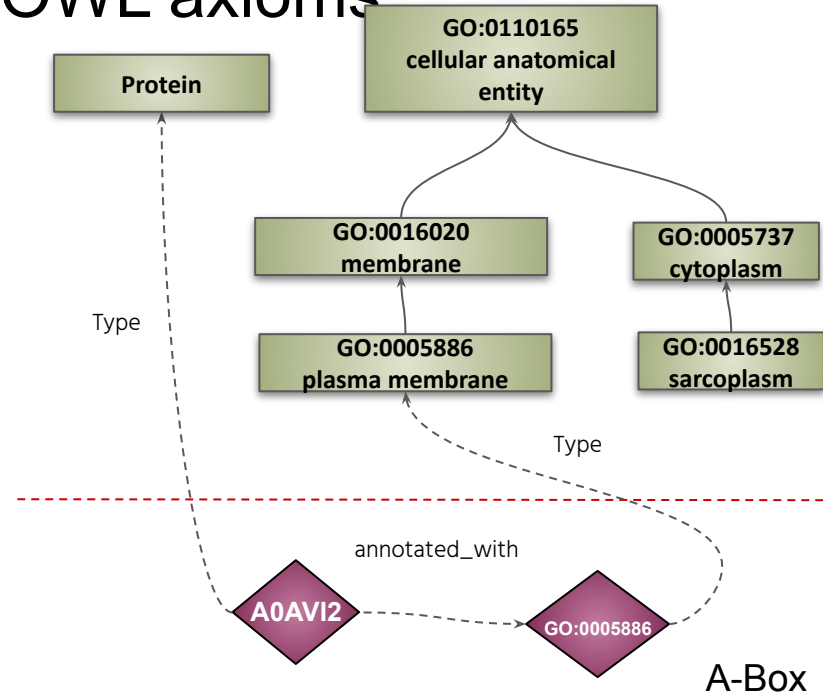


# Transforming GO Annotations to OWL axioms

Example:

Annotating protein **A0AVI2** To Gene Ontology

- Annotations to T-Box
  - A0AVI2**  $\sqsubseteq \exists$  *annotated\_with*. **GO:0005886**
- Annotations to A-Box
  - Protein(**A0AVI2**)
  - annotated\_with*(**A0AVI2**, **GO:0005886**)

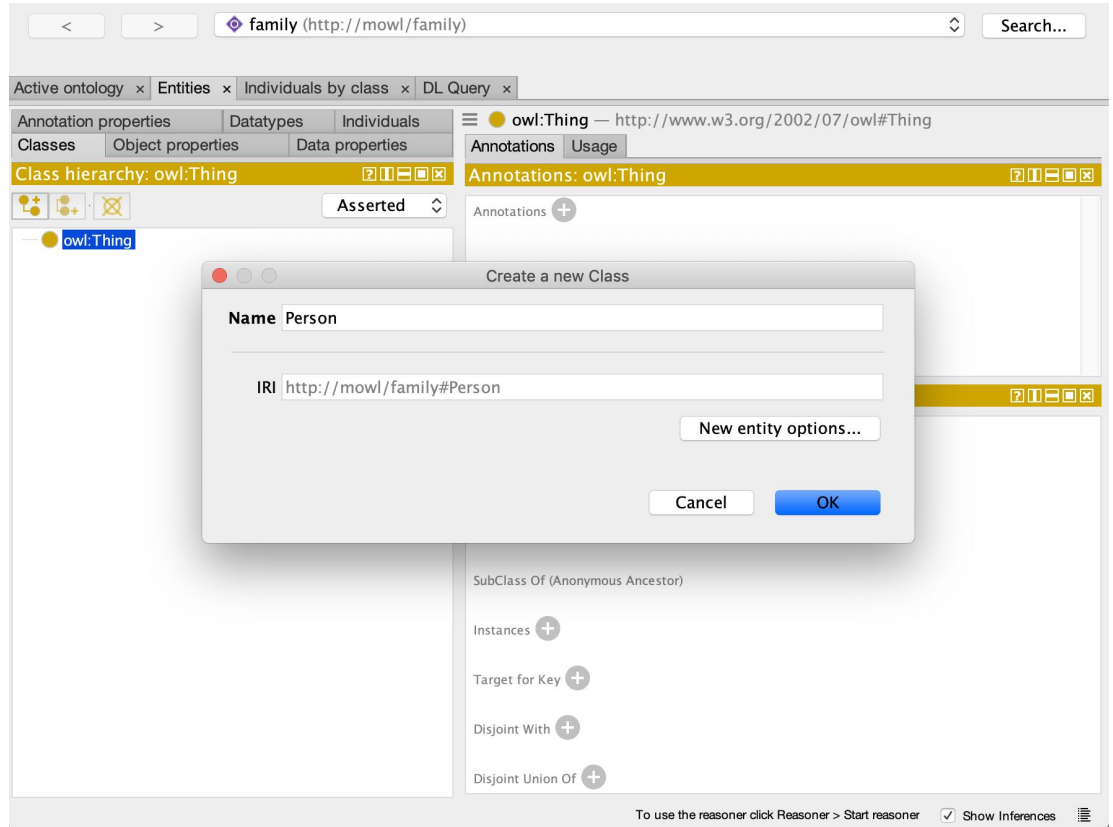


# Hands on (1):

- Creating the family ontology
  - WebProtégé
    - <https://webprotege.stanford.edu/#login>
    - Download from: <https://protege.stanford.edu/>

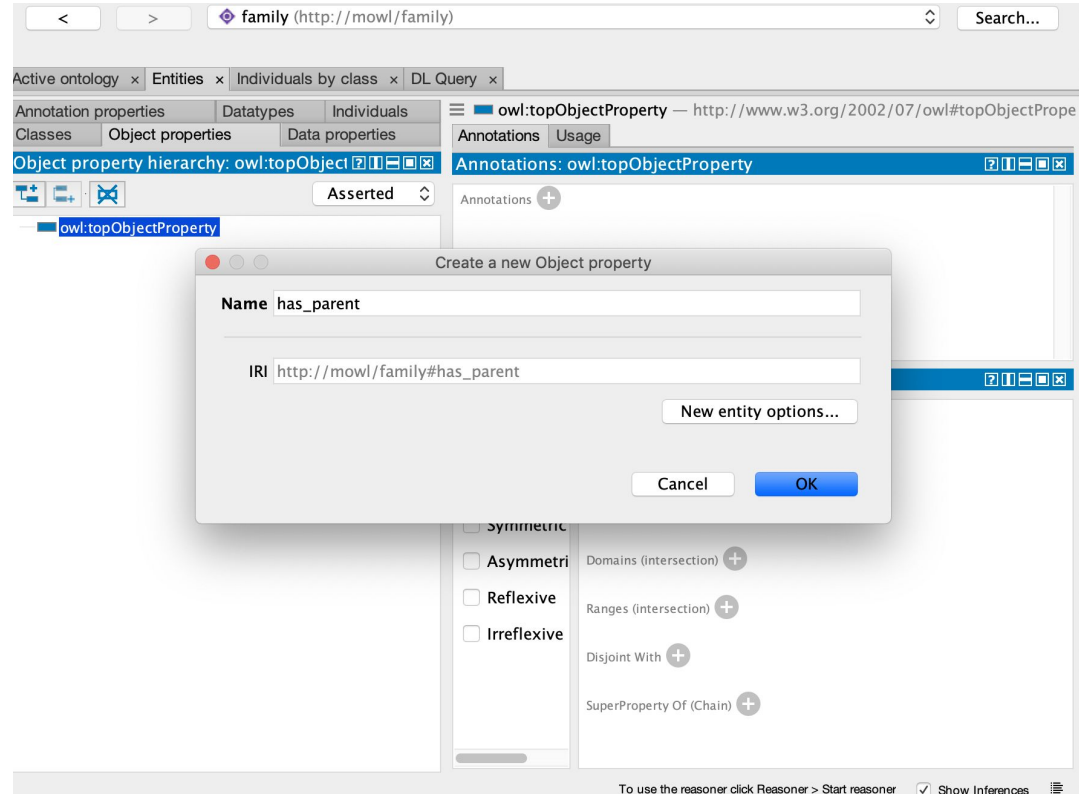
# Hands on (1):

- Protégé:
  - Add new classes



# Hands on (1):

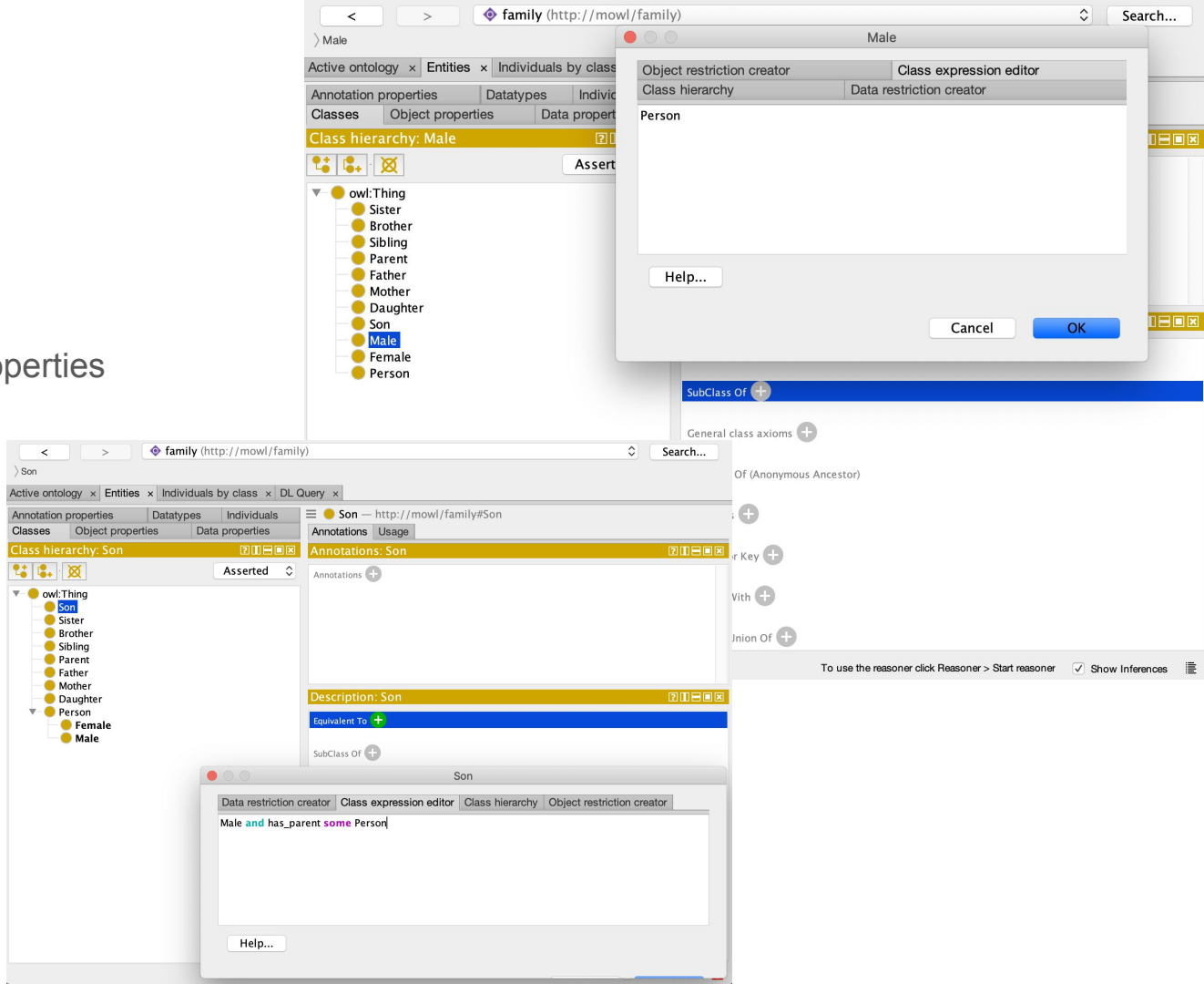
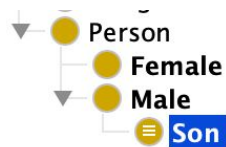
- Protégé:
  - Add new classes
  - Add new object properties





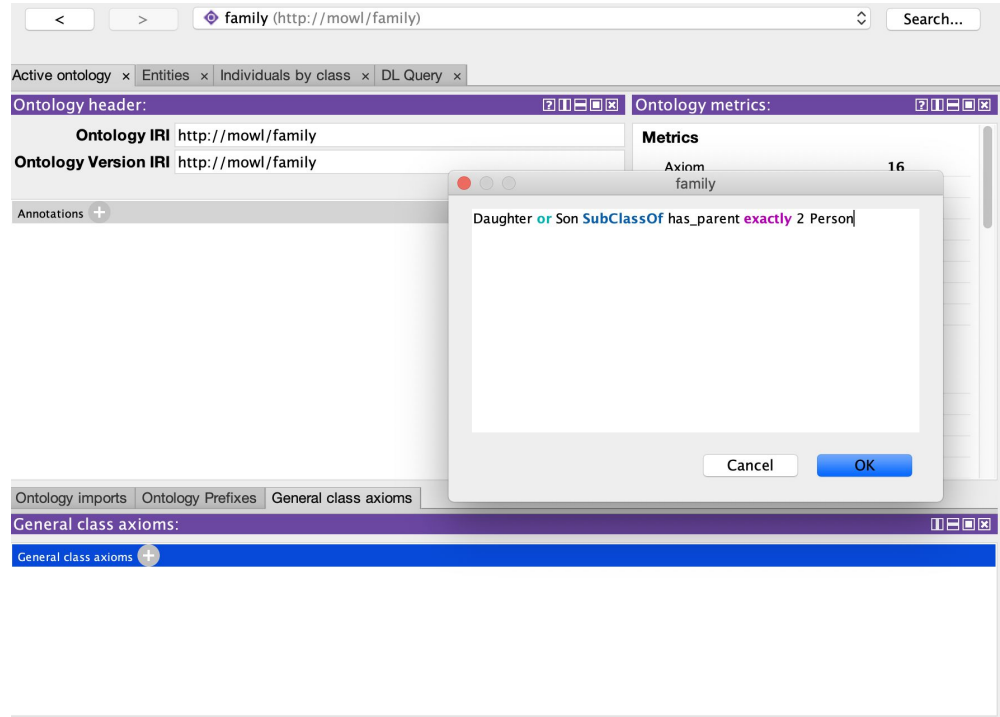
# Hands on (1):

- Protégé:
  - Add new classes
  - Add new object properties
  - Adding axioms
    - SubclassOf
    - Equivalent
    - ....



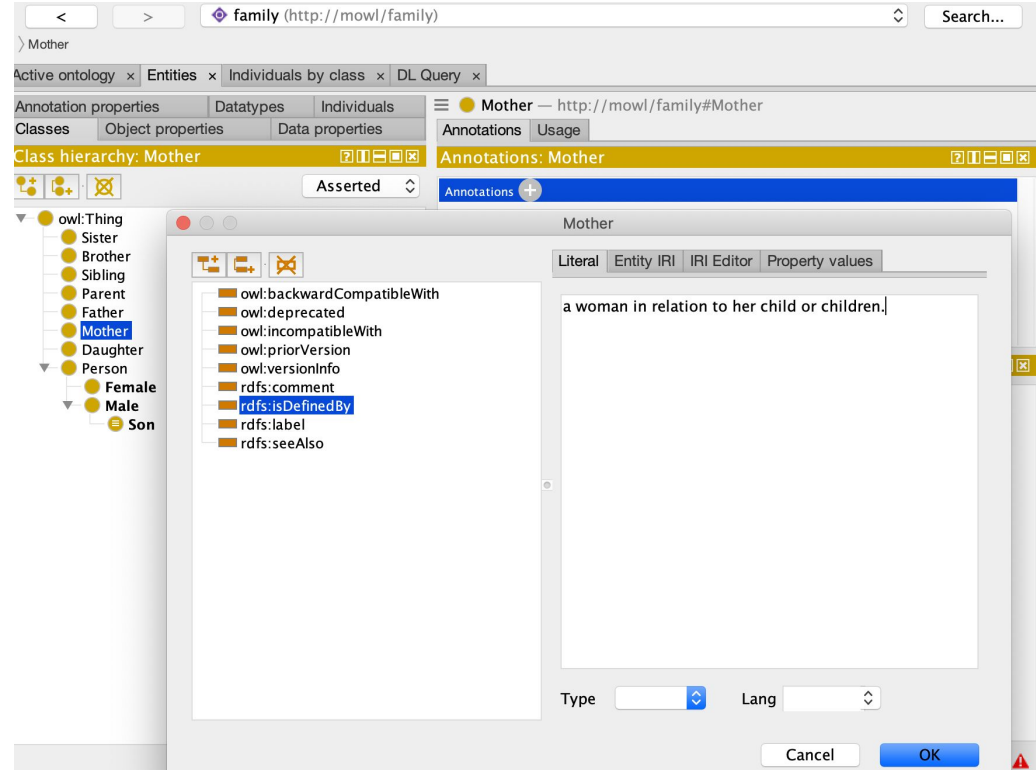
# Hands on (1):

- Protégé:
  - Add new classes
  - Add new object properties
  - Adding axioms
  - Adding GCIs



# Hands on (1):

- Protégé:
  - Add new classes
  - Add new object properties
  - Adding axioms
  - Adding GCIs
  - Adding definitions, synonyms



# Hands on (1):

- Protégé:
  - Add new classes
  - Add new object properties
  - Adding axioms
  - Adding GCIs
  - Adding definitions, synonyms
  - Using reasoners

