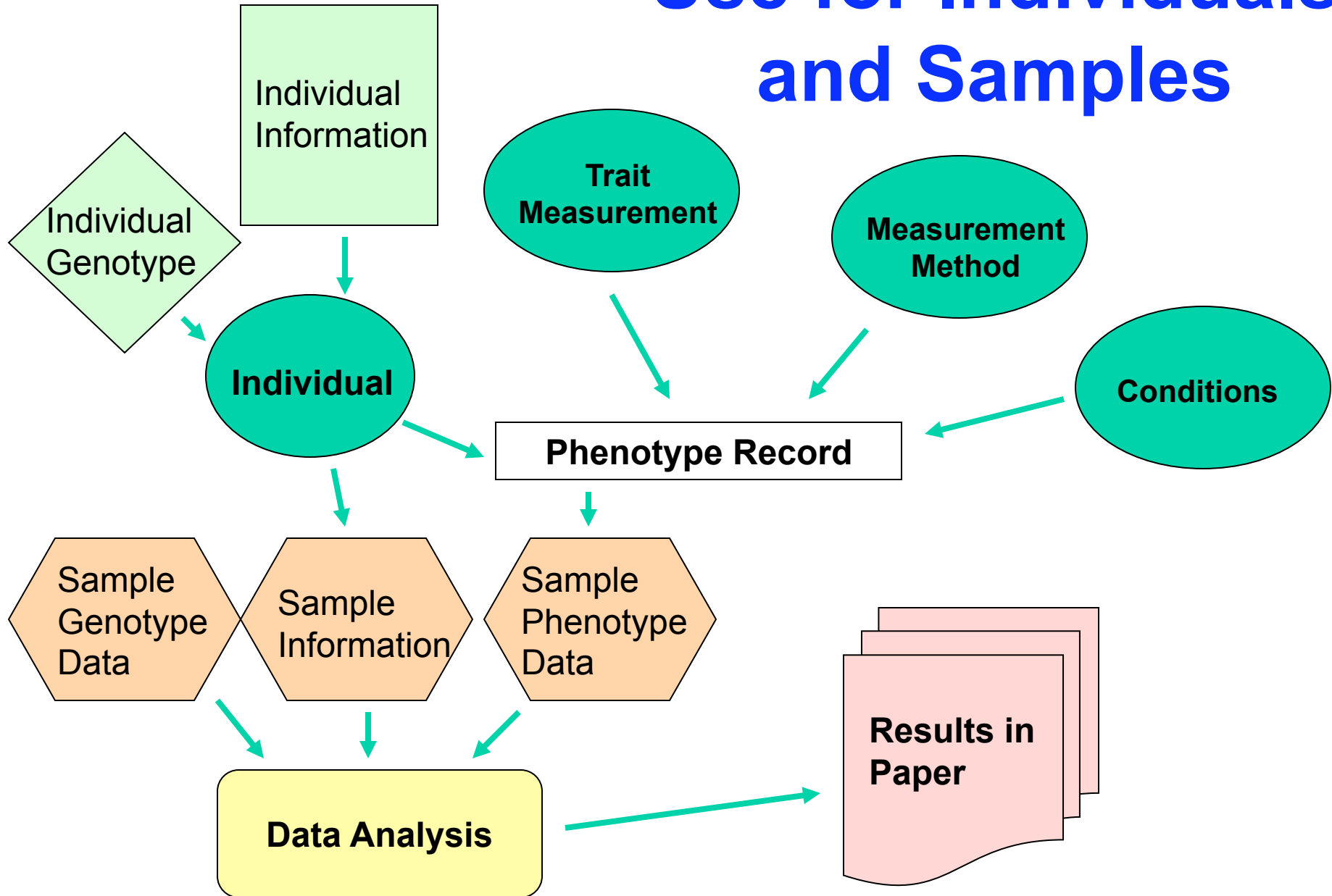


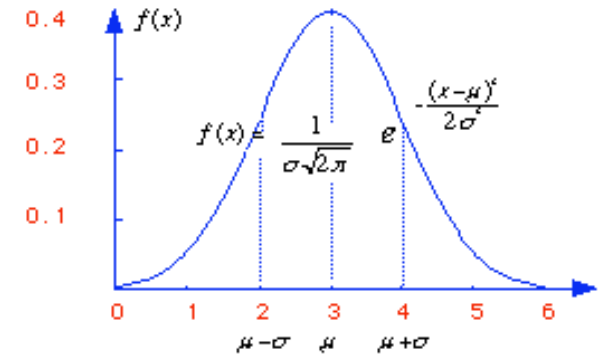
Global Perspectives on Animal Trait Ontology

J.M. Reecy, C. Park, Z.-L. Hu, I. Hulsegege,
H. Van Der Steen, J.-F. Hocquette



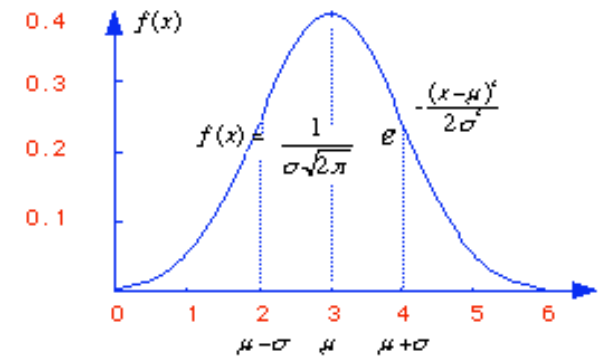
Use for Individuals and Samples



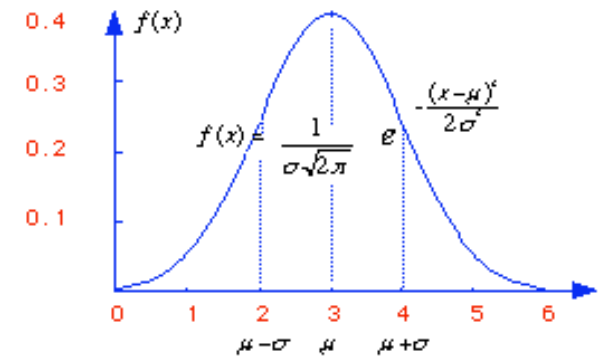


**"What's in a name? That
which we call a rose
By any other name would
smell as sweet."**

Romeo and Juliet (II, ii, 1-2)

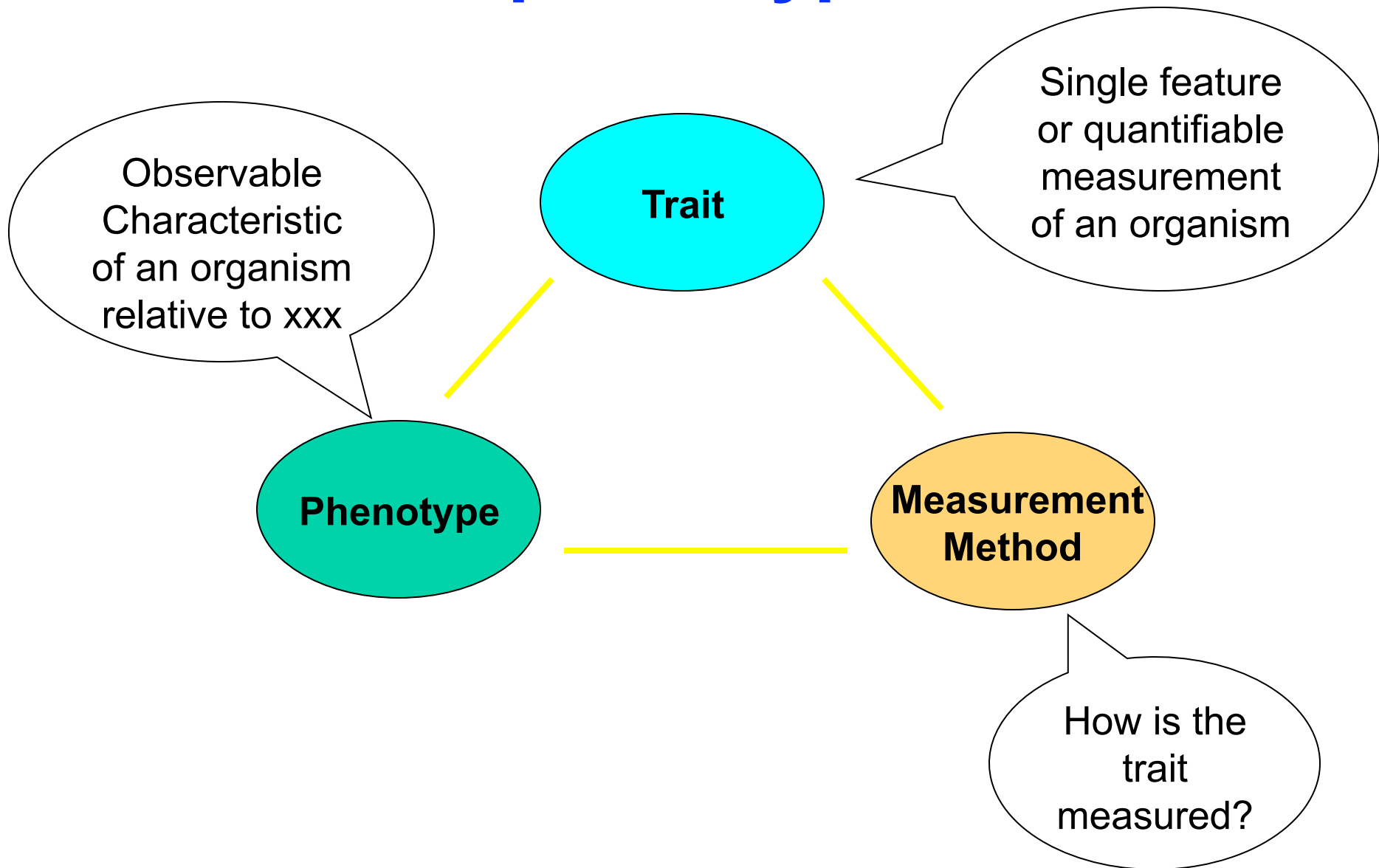


Have you ever struggled with trait nomenclature when comparing publications?

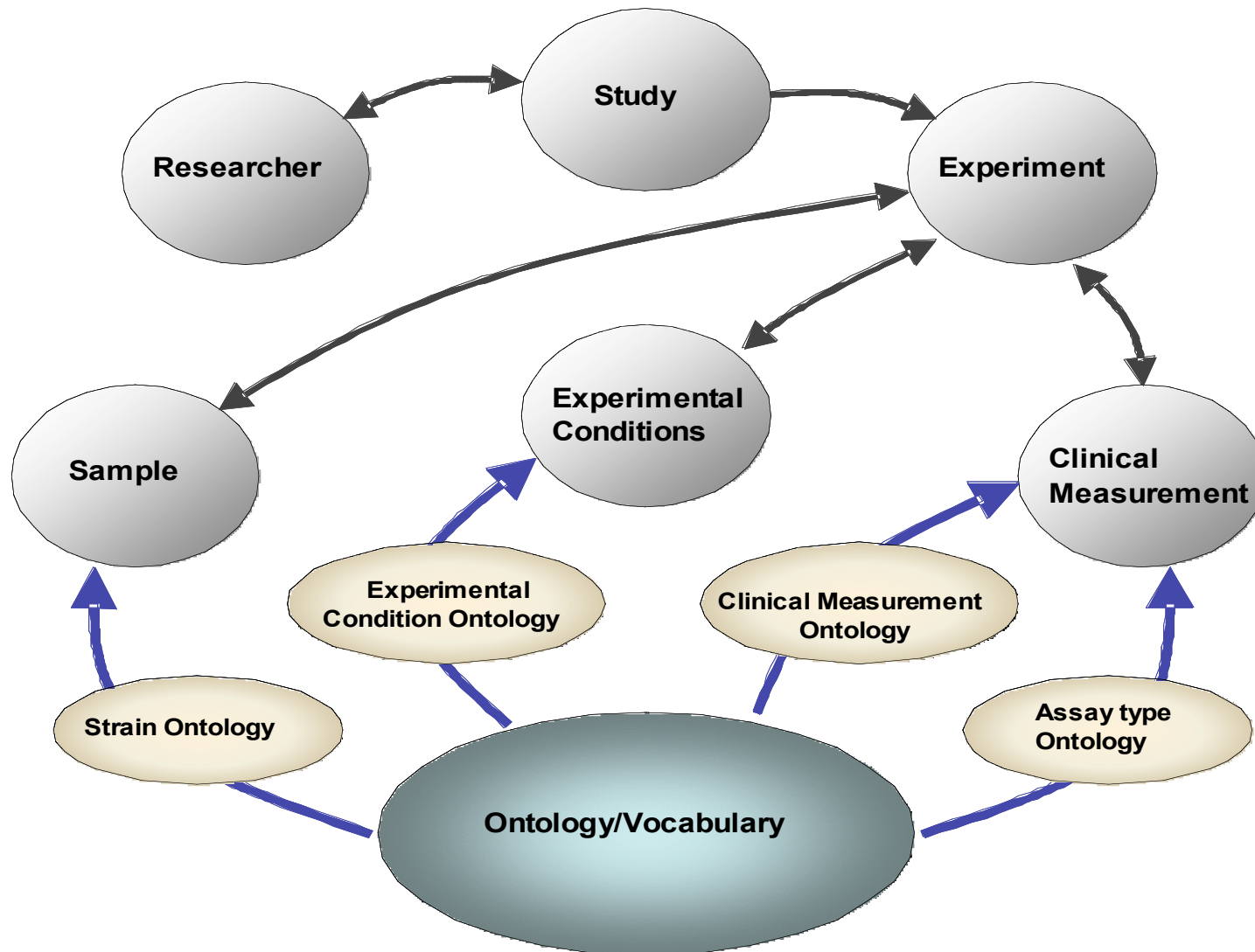


- **Why do we need an ATO**
 - To facilitate organization/comparison of data
 - Standardize nomenclature/facilitate knowledge transfer within and across species
 - Facilitate data retrieval
 - Facilitate data analysis
 - Enhance education

What is a phenotype/Trait?



Complex Experimental Measurement Data





The Open Biomedical Ontologies

Ontologies


Resources

Participate














About

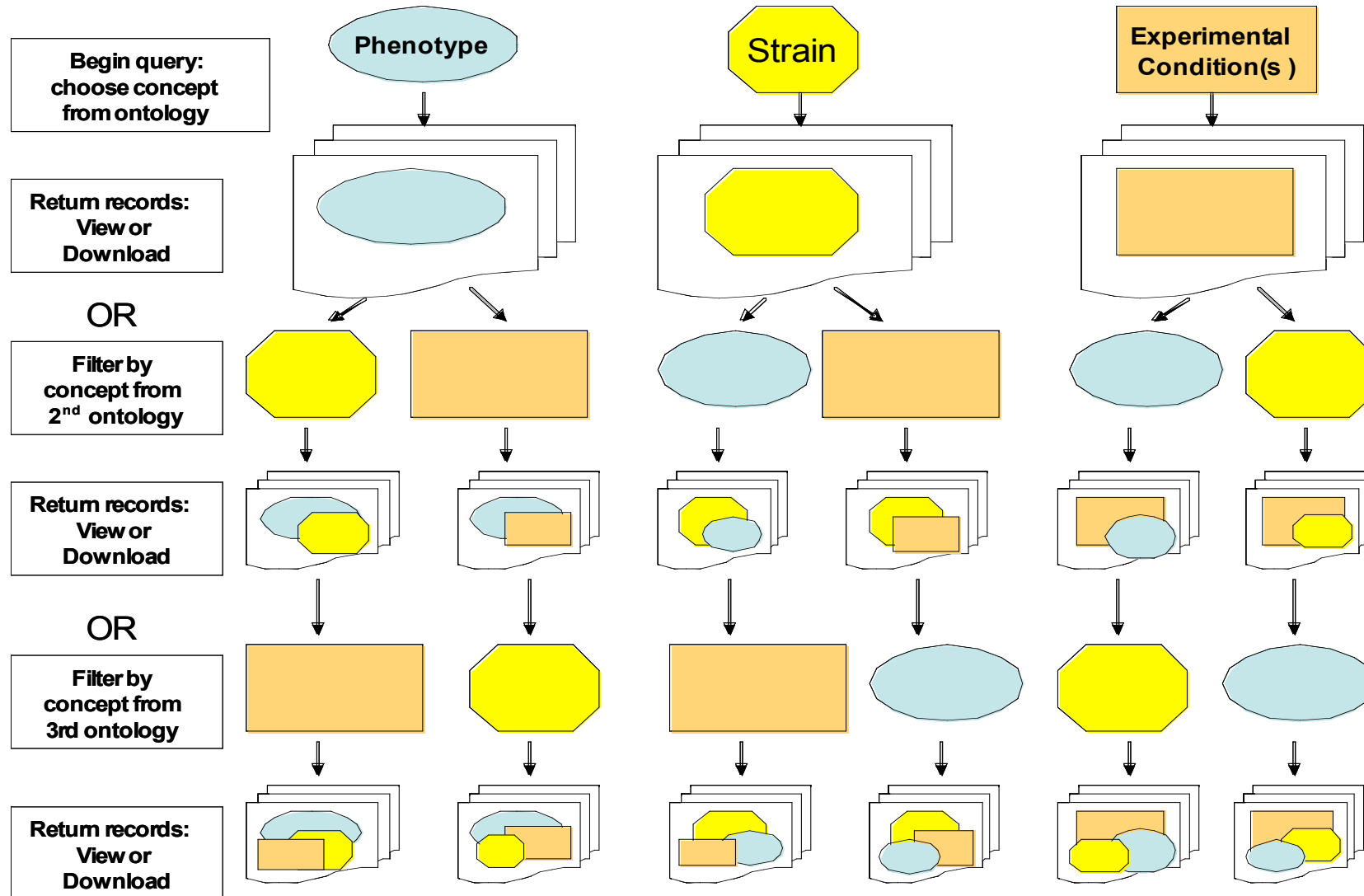
The OBO Foundry is a collaborative experiment involving developers of science-based ontologies who are establishing a set of principles for ontology development with the goal of creating a suite of orthogonal interoperable reference ontologies in the biomedical domain. The groups developing ontologies who have expressed an interest in this goal are listed below, followed by other relevant efforts in this domain.

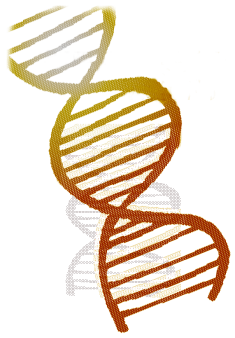
In addition to a listing of OBO ontologies, this site also provides a statement of the OBO Foundry principles, discussion fora, technical infrastructure, and other services to facilitate ontology development. We welcome feedback and encourage participation.

Click any column header to sort the table by that column. The  link to the term request trackers for the listed ontologies.

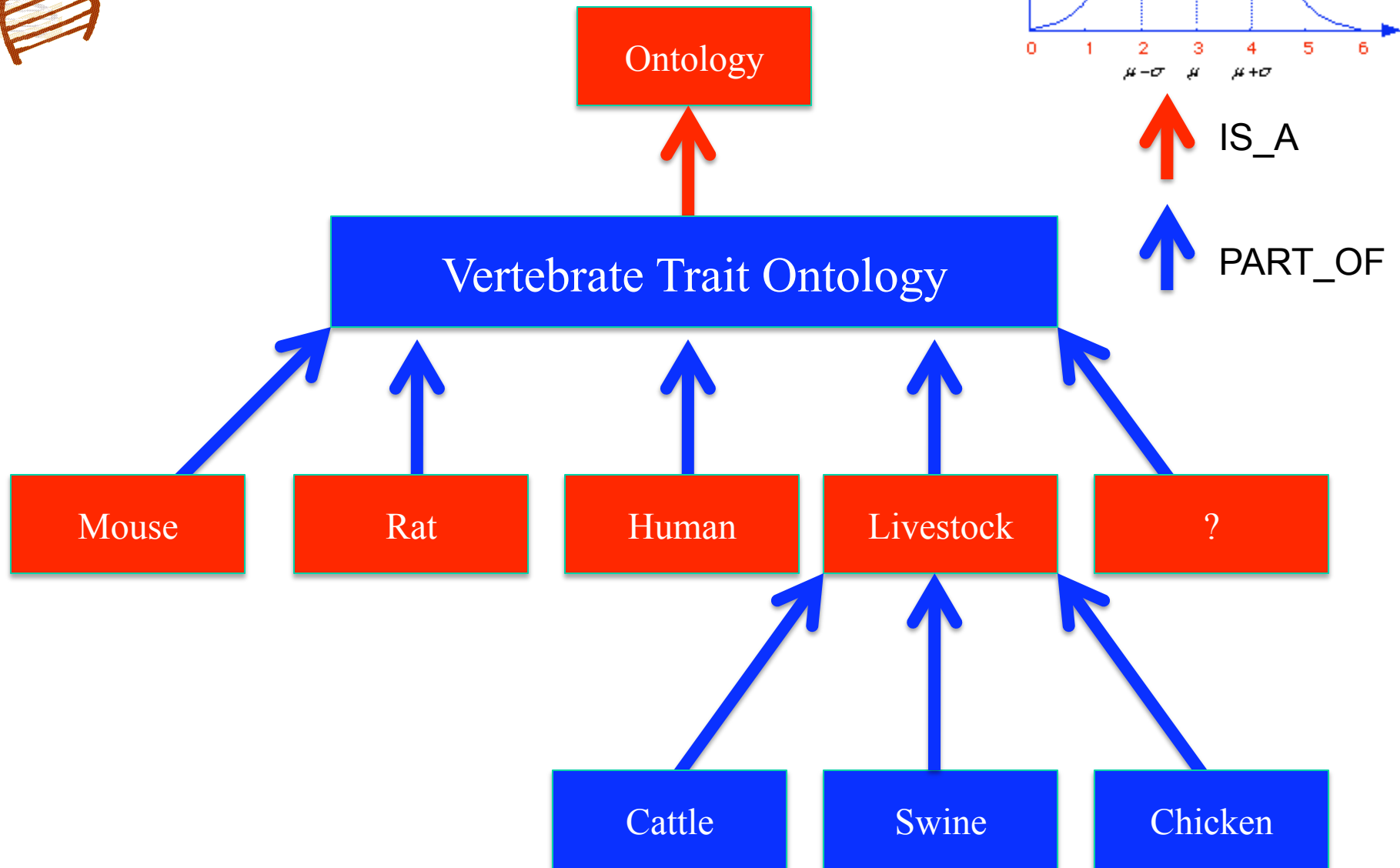
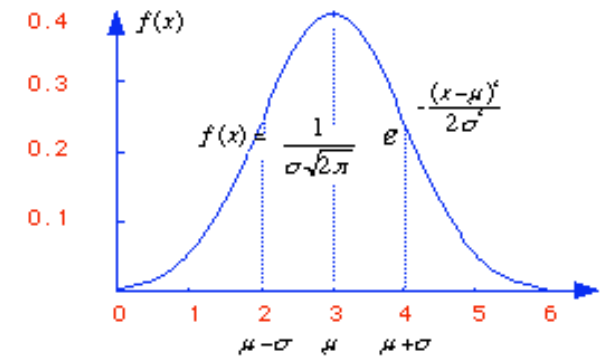
OBO Foundry candidate ontologies

Title	Domain	Prefix	File	Last changed
Amphibian gross anatomy	anatomy	AAO	amphibian_anatomy.obo	2008/06/19
Amphibian taxonomy	anatomy	ATO	amphibian_taxonomy.obo	
Ascomycete phenotype ontology	phenotype	APO	ascomycete_phenotype.obo	2009/07/10
Biological process	biological process	GO	gene_ontology_edit.obo 	2009/08/24
C. elegans development	anatomy	WBIs	worm_development.obo	2008/01/31
C. elegans gross anatomy	anatomy	WBbt	WBbt.obo 	2009/08/19
C. elegans phenotype	phenotype	WBPhenotype	worm_phenotype.obo	2009/08/19
Cell type	anatomy	CL	cell.obo 	2008/12/09
Cellular component	anatomy	GO	gene_ontology_edit.obo 	2009/08/24
Cereal plant trait	phenotype	TO	plant_trait.obo 	2009/08/18
Chemical entities of biological interest	biochemistry	CHEBI	chebi.obo 	2009/07/29
Common Anatomy Reference Ontology	anatomy	CARO	caro.obo 	2007/06/17
Dictyostelium discoideum anatomy	anatomy	DDANAT	dictyostelium_anatomy.obo 	2008/05/29
Drosophila development	anatomy	FBdv	fly_development.obo 	2008/12/11
Drosophila gross anatomy	anatomy	FBbt	fly_anatomy.obo 	2009/03/17
Environment Ontology	environment	ENVO	envo.obo 	2009/05/20
Evidence codes	experiments	ECO	evidence_code.obo	2009/03/20
Fly taxonomy	taxonomy	FBsp	fly_taxonomy.obo 	2007/04/10
Foundational Model of Anatomy (subset)	anatomy	FMA	fma2_obo.obo 	
Fungal gross anatomy	anatomy	FAO	fungal_anatomy.obo	2009/07/10





Development of Vertebrate Trait Ontology





Search ATO:

☐ Exact Match

[Advanced Query](#)

Ontology Filter:

All
feed consumption
First rib backfat

Display options:

☐ XML

☐ Flat File

[PERMALLINK](#)
[ATO HOME](#)

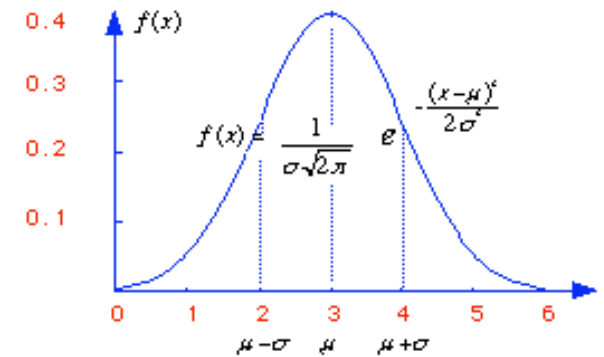
☐ all : all

☐ ① ATO:0000346 : ATO

☐ ① ATO:0000348 : Bos taurus

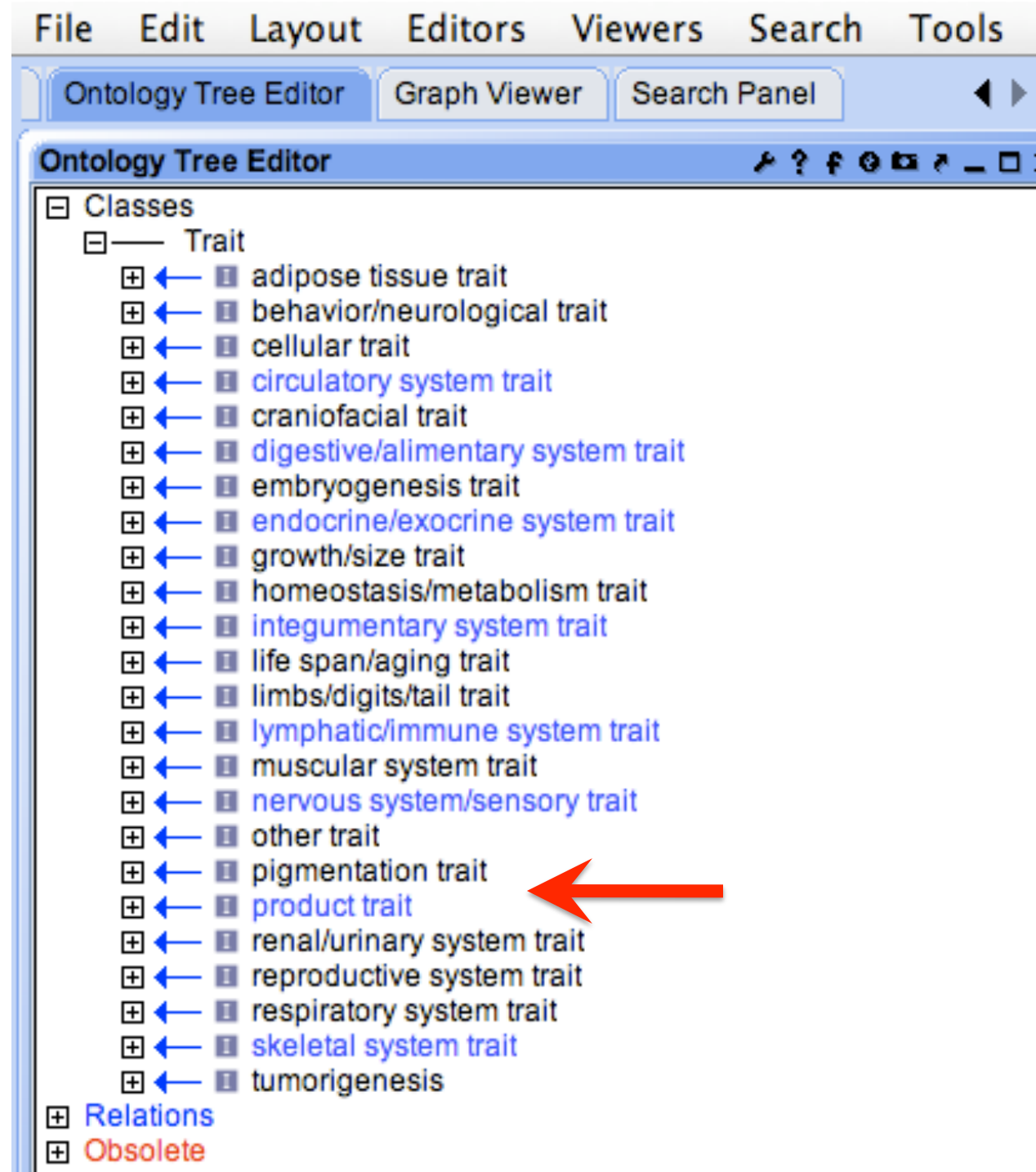
☐ ① ATO:0000482 : Gallus gallus

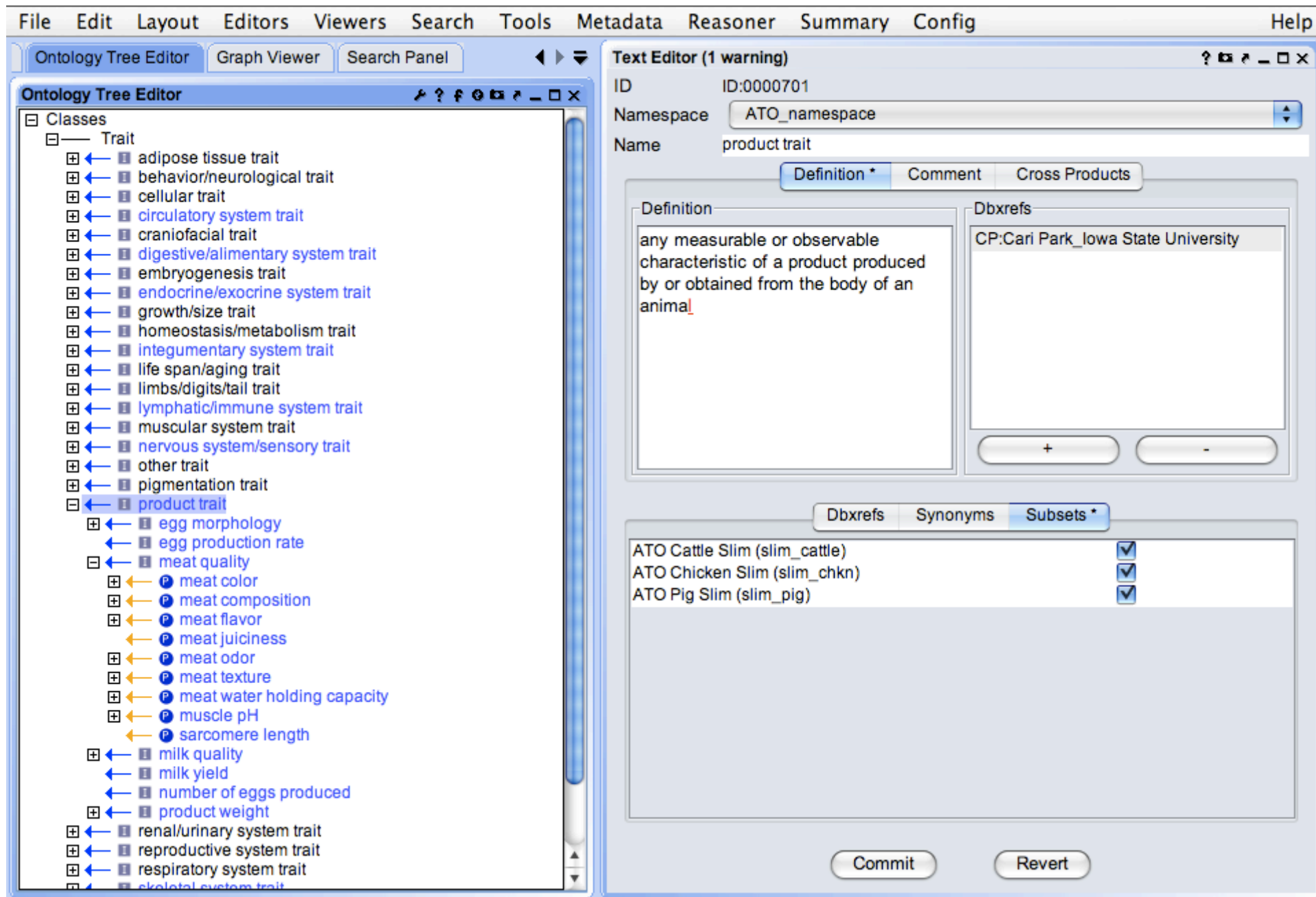
☐ ① ATO:0000347 : Sus scrofa

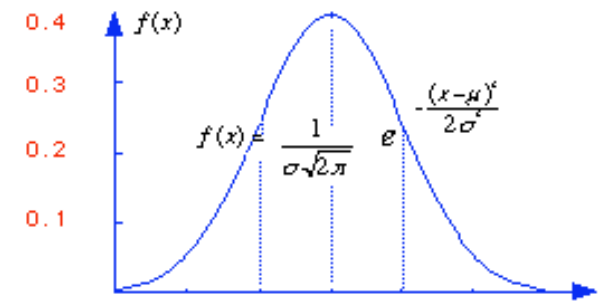




How to organize traits?







Ontology Tree Editor Graph Viewer Search Panel

Ontology Tree Editor

- [-] Trait
 - [+] [i] adipose tissue trait
 - [+] [i] behavior/neurological trait
 - [+] [i] cellular trait
 - [+] [i] circulatory system trait
 - [+] [i] craniofacial trait
 - [+] [i] digestive/alimentary system trait
 - [+] [i] embryogenesis trait
 - [+] [i] endocrine/exocrine system trait
 - [+] [i] growth/size trait
 - [+] [i] homeostasis/metabolism trait
 - [+] [i] integumentary system trait
 - [+] [i] life span/aging trait
 - [+] [i] limbs/digits/tail trait
 - [+] [i] lymphatic/immune system trait
 - [+] [i] muscular system trait
 - [+] [i] muscle morphology
 - [+] [i] eye muscle morphology
 - [+] [i] heart muscle morphology
 - [+] [i] muscle development
 - [+] [i] muscle fiber morphology
 - [+] [i] muscle size
 - [+] [i] skeletal muscle mechanoreceptor morphology
 - [+] [i] skeletal muscle morphology
 - [+] [i] smooth muscle morphology
 - [+] [i] tendon morphology
 - [+] [i] muscle physiology
 - [i] ability to move the eye muscles
 - [i] adenosine deamination
 - [i] muscle androstenone level

Text Editor (1 warning)

ID MT:0005369

Namespace Trait.ontology

Name muscular system trait

Definition * Comment Cross Products

Definition

any measurable or observable characteristic of the contractile tissue that produces movement

Dbxrefs

CP:Carl Park_Iowa State Univer

+

-

Dbxrefs Synonyms Subsets *

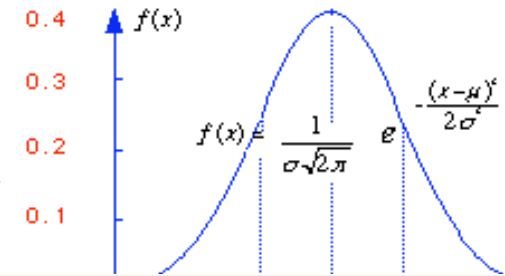
ATO Cattle Slim (slim_cattle) ☒

ATO Chicken Slim (slim_chkn) ☒

ATO Pig Slim (slim_pig) ☒



Reproduction Ontology



File Edit Ontologies Reasoner Tools Refactor Tabs View Window Help

Ontology1191324893.owl (<http://www.owl-ontologies.com/Ontology1191324893.owl>)

Active Ontology \ Entities \ Classes \ Object Properties \ Data Properties \ Individuals \ OWL Viz \ DL Query \

Asserted class hierarchy \ Inferred class hierarchy \

Asserted class hierarchy: ReproductivePerformance

- ReproductiveBehaviour
- ReproductiveFailure
- ReproductivePerformance
 - AgeAtFirstConception
 - AgeAtFirstParturition
 - AgeAtFirstService
 - AgeAtPuberty
 - CalfCrop
 - CalvingIndex
 - EaseOfParturition
 - OestrousCycle
 - ParturitionInterval
 - ParturitionPercentage
 - Rate
 - ReproductiveIndex
 - ServicesPerConception
 - VoluntaryWatingPeriod

Object property hierarchy \ Data property hierarchy \ Individuals \

Object properties:

Class Annotations \ Class Usage \

Annotations: ReproductivePerformance

Annotations +

- hasDefinition: "The final outcome of all processes involved in 1) quality and quantity of gametes of both sexes 2) pregnancy rates 3) survival rates of embryo, foetus and newborn [source: REPO]"
- hasIdentifier: "REPO:0000002"
- hasSynonym: "breeding performance"
- hasSynonym: "reproductive ability"
- hasSynonym: "reproductive capacity"
- hasSynonym: "reproductive efficiency"

Description: ReproductivePerformance

Equivalent classes +

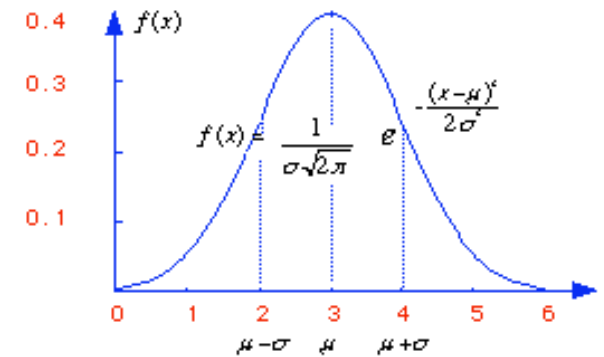
Superclasses +

- Thing

Inherited anonymous classes

Members +

Disjoint classes +



Future Cross mapping of Ontologies

Case: A

Disease

X



Phenotype

Y



Trait

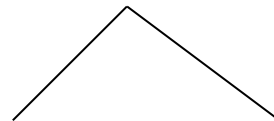
Z

Case: B

X



Y

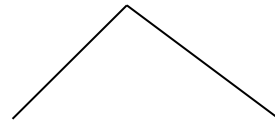


Z1

Z2

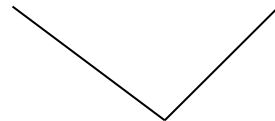
Case: C

X



Y1

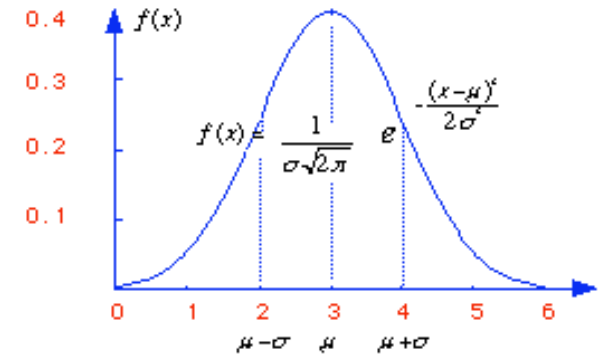
Y2



Z



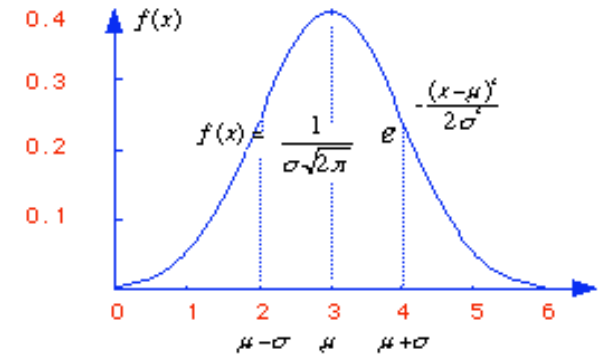
ATO Example



Disease	Phenotype(s)	Trait
1. Obesity	Increased adipose weight	Adipose weight
	Increased lipid weight	Lipid weight
2. Dwarfism	Decreased bone length	Bone length
	Abnormal bone morphology	Bone morphology
3. Mastitis	Increased somatic cell count	Somatic cell count



Summary



- Assembled a consortium
 - ISU, RGD, MGI, EADGENE, INRA
 - Experts welcome
- Trait Ontology development is on-going
 - Core curation rules have been develop
 - Higher-order terms have been identified
- Data analysis resource

Acknowledgements

Iowa State University

- Sue Lamont
- Max Rothschild
- Chris Tuggle
- Zhiliang Hu
- Eric Fritz
- Laron Hughes
- Jie Bao
- Neeraj Kaul
- Vasant Honavar



Collaborators

- Mary Shimoyama
- Anne Kwiktek
- Mindy Dwindell
- Diane Munzenmaier
- EADGENE
 - Hein Van der Steen
 - Ina Hulsegge
- INRA
 - Pierre-Yves Le Bail
 - Jean-Francios Hocquette