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Skeletal Anatomy Jamboree

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Purpose of jamboree:

- Re-evaluate skeletal ontology for fishes (TAO), focussing at level of cell, tissue, and skeletal element
- Coordinate high-level terms with other vertebrate anatomy ontologies
- Goal to improve usefulness and accuracy of reasoning that is based on this ontology

Ontology important for:

- answering specific questions
- discovery, data mining
- clear communication
- interoperability across databases

*need to keep in mind what it is we want to know and communicate, i.e., use



Phenoscape

2007

- Goal: To prototype a curated, ontology-based evolutionary ‘phenotype’ database that maps to genetic databases
- Foster devo-evo synthesis
- Enable data-mining and discovery for broad scale evolutionary patterns

The ZFIN website homepage features a header with the ZFIN logo and navigation links for Research, General Information, and ZIRC. The main content area includes sections for Genes / Markers / Clones, BLAST at ZFIN, Gene Expression Antibodies, Morphology / Morphants / Transgenics, Anatomy, Publications, Community, and Data. A sidebar on the right provides links to the Zebrafish International Resource Center, Genomics, Zebrafish Programs, News, and a photo of a zebrafish.

Zebrafish (zfin.org)

The Tree of Life (ToL) web project page for Ostariophysan fishes shows a photograph of several fish in a tank. Below the image is a phylogenetic tree diagram with the following branches: Genypterygii (with Chanoides), Otophysi (with Cypriniformes, Characiformes (Tetras, piranhas, hatchetfish), Gymnotiformes, and Siluriformes).

Ostariophysan fishes

Search for:

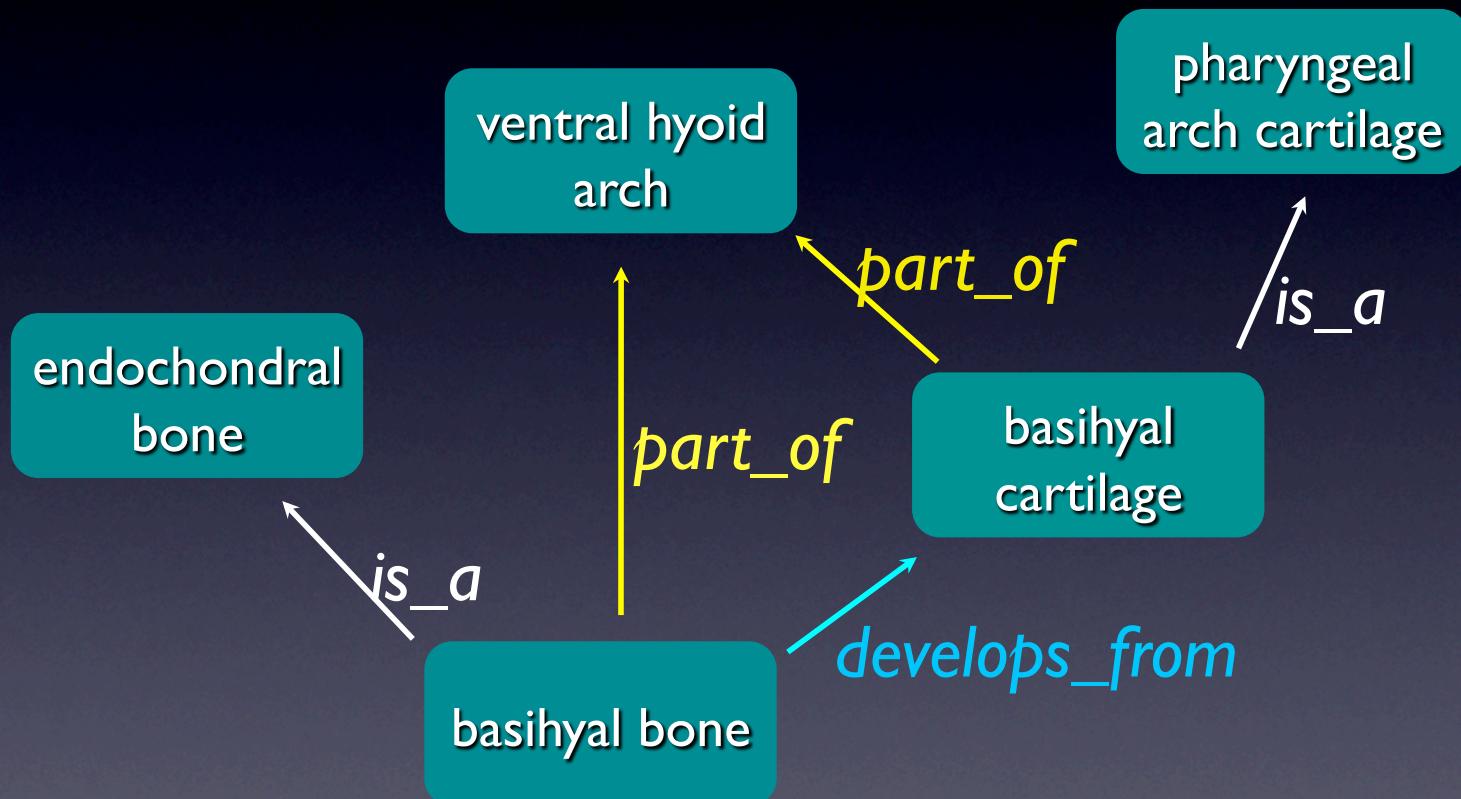
- Search for candidate genes underlying evolutionary morphology
- Search for taxa with particular morphologies
- Aggregate morphological data across studies



Requirements:

1. Ontologies
2. Curation
3. Database

Ontology: Teleost Anatomy Ontology

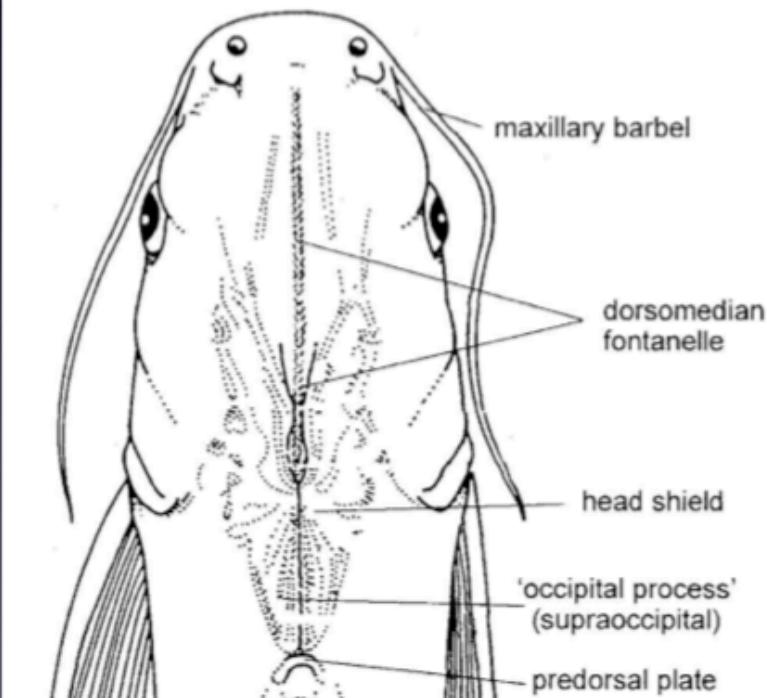


Dahdul, W. M., J. G. Lundberg, P. E. Midford, J. P. Balhoff, H. Lapp, T. J. Vision, M. A. Haendel, M. Westerfield, and P. M. Mabee. 2010. The Teleost Anatomy Ontology: Anatomical representation for the comparative synthesis of fish.

Curation: ‘free-text’ descriptions into machine-readable form using ontologies

47. *Shape of posterior dorsomedian fontanelle.* An elongate-rectangular posterior fontanelle of most catfishes appears to be the plesiomorphic condition (see also Tilak 1963, 1964, 1965a; Lundberg 1982; Arratia 1987; Grande 1987). An irregularly-shaped or rounded

State 0 = frontal broad anteriorly and moderately narrow posteriorly, anterior space reduced (adults) and arms moderately wide; 1 = frontal moderately broad posteriorly, anterior space moderately enlarged; 2 = frontal broad posteriorly, anterior arms narrow, space enlarged.



49. *Laminar bone over the anterior vertebrae.* The laminar bone is usually continuous medially in ariids (except in *Galeichthys* and *Ancharius*) and is more extensive in larger individuals, an ontogenetic change evidenced in most taxa. However, the excavation of the laminar bone posteromedially and the overlapping of the transverse process bases laterally is variable.

I consider that a minimal cover over the ventral processes is plesiomorphic in ariids as exposed transversely. A 'cover' as exposed transversely is defined as a median excavation on the laminar bone in ariids extends over the ventral processes. Some ariids possess an apophysis, a laminar shelf, such as a high and acute in *Batilus armiger* or double keel

?	0	0	?	3	0	?	0	?	0	0	?	1	0	?
1	?	?	?	0	?	?	0	1	0	0	2	0	2	0
0	?	?	?	0	?	?	0	?	0	0	2	1	2	1
0	?	?	?	0	?	?	0	1	?	?	2	1	2	0
1	?	0	?	2	0	?	0	1	0	?	2	1	2	?
1	?	?	?	0	0	?	0	?	2	1	1	1	1	1
3	?	?	?	0	?	?	0	1	0	1	2	1	0	2
3	?	?	?	0	?	?	0	1	0	?	2	1	0	1
?	?	?	?	0	1	?	0	0	0	0	1	1	1	?

Curated 4,627 characters in 3,449 species from 51 papers

TAXON	# PAPERS	# SPECIES	# CHARACTERS
Cypriniforms	8	676	794
Siluriforms	20	1724	2,110
Characiforms	10	754	1,156
Gymnotiforms	1	116	231
Gonorynchiforms	3	41	467
Clupeiforms	5	200	439
Euteleosts	3	145	582
Total	51	3,656	5,779
TOTAL ANNOTATED			

Database: Phenoscape KB

kb.phenoscape.org

486,714 evolutionary
phenotype assertions

16,426 phenotype
statements about
3,539 genes

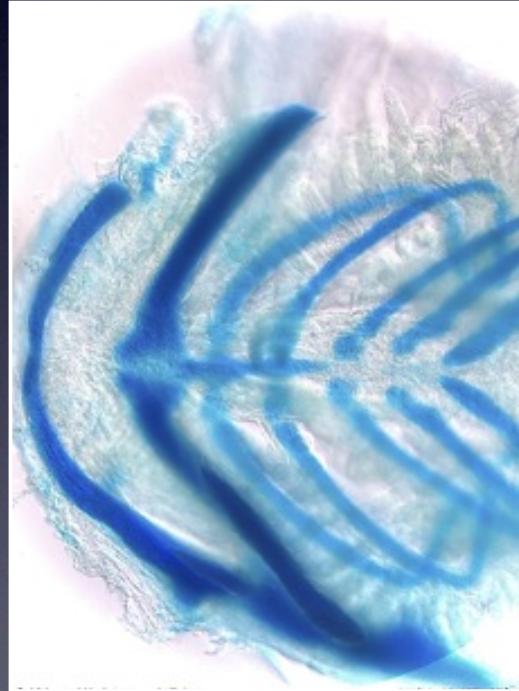


25 July 2009

Kb enables:

- Selection of candidate genes for evolutionary morphology
- Broad scale view, patterns of morphology across taxa
- E.g.,

I. Evolutionary phenotype: Loss of skeletal ‘tongue’ **(basihyal)** in all catfishes



Ictalurus punctatus

What genes are involved in the development of the basihyal?

The Phenoscope Knowledgebase is currently in BETA testing - we would greatly value your feedback as we continue its development.

New search (entity, taxon, gene): Go

Home Provide Feedback About Contributors

entity term: basihyal element

Properties	
ID:	TAO:0001891
Ontology:	Teleost Anatomy
Synonyms:	None
Definition:	None
Relationships (about)	
is a type of:	skeletal element
is part of:	ventral hyoid arch
has subtype:	basihyal bone , basihyal cartilage
More information	
View at NCBO Bioportal	

Phenotypes

Entity	Quality	Zebrafish Genes	Taxa
anterior side of basihyal element	composition : composition of	0	19
anterior side of basihyal element	structure	0	19
basihyal bone	shape : triangular	0	239
basihyal bone	composition : ossified	1	0
basihyal bone	size : decreased length, decreased size, increased length	2	196
basihyal bone	count : absent, present	2	35
basihyal cartilage	count : absent, present	5	37
basihyal cartilage	quality : malformed	1	0
basihyal cartilage	size : decreased length, decreased size	3	0
basihyal cartilage	position : mislocalised laterally, mislocalised posteriorly	2	0
fossa of basihyal bone	size	0	27

Results of query: list of **basihyal element**
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Mutations in *brpf1* result in basihyal cartilage loss (zebrafish phenotype)

PHENOTYPE:

Genotype(s): *brpf1*^{b943/b943}, *brpf1*^{t20002/t20002}, *brpf1*^{t25114/t25114}

Observed In: basihyal

Stage Range : Day 5

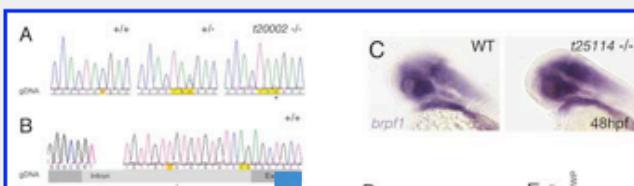
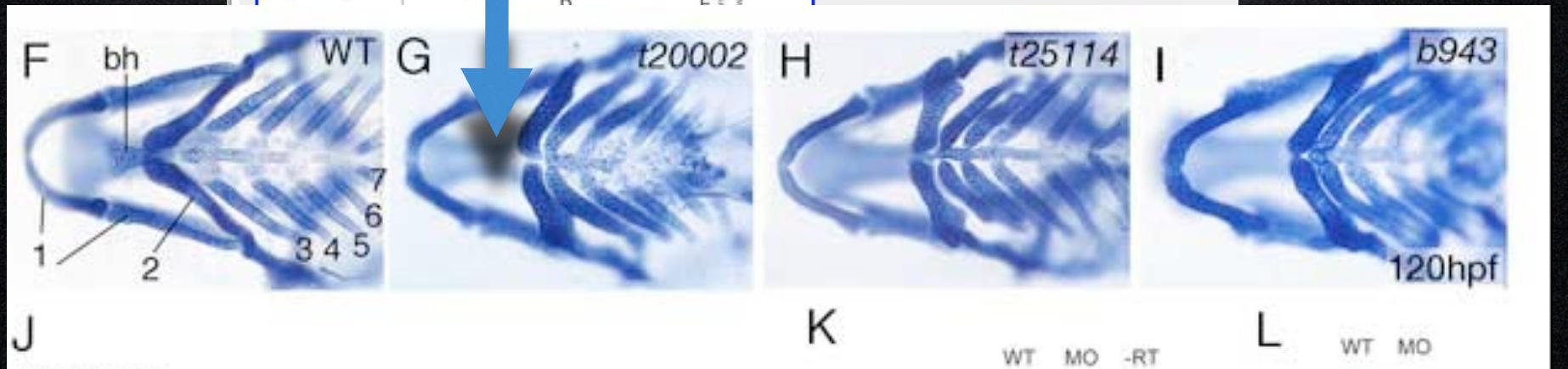


Fig. S1 Analysis of *bpfr1* mutations:

(+/+), heterozygous (+/-) and *t20002* mutant by an asterisk. (B) Sequencing profile (-/-, lower panel). The *t25114* mutation



ZFIN ID: ZDB-FIG-080604-30

[Laue et al., 2008](#) - The multidomain protein Brpf1 binds histones and is required for Hox gene expression and segmental identity. Development 135(11):1935-1946 - [Full text @ Development](#)

2. What genes are involved in the development of intramembranous bones?

The Phenoscope Knowledgebase is currently in BETA testing - we would greatly value your feedback as we continue its development.

Search (entity, taxon, gene):

Enter entity terms (e.g. basithyal bone), taxonomic names (e.g. Ictaluridae), or gene names or symbols (e.g. cadherin 8, cathe).

[Home](#) [Provide Feedback](#) [About](#) [Contributors](#)

entity term: intramembranous bone

Properties	
ID:	TAO:0001644
Ontology:	Teleost Anatomy
Synonyms:	<i>None</i>
Definition:	Bone that forms directly within mesenchyme, and does not replace other tissues.
Relationships (about)	
is a type of:	bone
develops from:	mesenchyme
has subtype:	dermal bone , membrane bone
More information	
View at NCBO Bioportal	

Phenotypes: intramembranous bone

This table lists phenotypes, categorized by type of quality, which inhere in intramembranous bone, its subtypes, or parts.

Entity	Quality	Zebrafish Genes	Taxa
abdominal scute	relational structural quality : relational structural quality towards rib	0	13
abdominal scute	count : absent, count of 2; 1 pair, count of 4; 2 pairs, count of 6 or 8; 3 or 4 pairs, present	0	399
accessory vomerine tooth plate	count : absent, present	0	412
actinotrichium of lepidotrichium	position : mislocalised	1	0
anal fin lepidotrichium	quality : aplastic growth	2	0
anal fin lepidotrichium	shape : undulate	1	62
anal fin lepidotrichium	position	0	27
anal fin lepidotrichium	relational structural quality : relational structural quality towards anal fin distal radial, relational structural quality towards anal fin proximal radial	0	155
anal fin lepidotrichium	size : decreased thickness, decreased width, increased thickness, size towards anal fin lepidotrichium	0	220
anal fin lepidotrichium	structure	0	66
anal fin lepidotrichium	relational spatial quality : relational spatial quality towards vertebral column	0	27
anal fin lepidotrichium	count : count of 16 (low), count of 2, count of 20.1-25.5, count of hihg, count of ≤13 and 30 more	0	451
anal fin lepidotrichium 1	structure	0	27
anal fin lepidotrichium 3	shape	0	74
anal fin lepidotrichium 3	size	0	74
anal fin lepidotrichium of female	size	0	74

Results of query: list of intramembranous bones
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TAO:‘Intramembranous bone’

List sorted by genes

The Phenoscope Knowledgebase is currently in BETA testing - we would greatly value your feedback as we continue its development.

PHENOSCAPE BETA

entity term: intramembranous bone

Properties

ID: TAO:0001644
Ontology: Teleost Anatomy
Synonyms: None
Definition: Bone that forms directly within mesenchyme, and does not replace other tissues.

Relationships (about)

is a type of: [bone](#)
develops from: [mesenchyme](#)
has subtype: [dermal bone](#), [membrane bone](#)

More information

[View at NCBO Bioportal](#)

Phenotypes: intramembranous bone

This table lists phenotypes, categorized by type of quality, which inhere in intramembranous bone, its subtypes, or parts.

Entity	Quality	Zebrafish Genes	Taxa
scale	shape : round	17	395
opercle	size : decreased size, decreased thickness, decreased width, increased width, size towards head and 1 more	12	233
scale	position : inverted, spatial pattern	10	0
scale	quality : deciduous (generic), discontinuous, malformed, non-deciduous (any body part), occurrence quality	6	184
scale	size : decreased size	5	231
scale	count : absent, count of 0, count of 0-3, count of 4-5, count of 6 and 5 more	5	262
scale	relational spatial quality : angular placement, relational spatial quality towards process of parietal bone, relational spatial quality towards supraoccipital crest	3	19
opercle	shape : rectangular, round, surface feature shape, triangular and 1 more	3	1276
caudal fin lepidotrichium	quality : aplastic growth, malformed	3	0
branchiostegal ray	count : count of 3 or 4, count of 5-6, count of 7, count of >7, count of mean: 17 (range: 16-19) and 20 more	3	956
lepidotrichium	quality : abnormal, normal	2	0
gill raker	count : absent, count of 15-22, count of 26-31, count of 32-39, count of ≤11 and 4 more	2	589
gill raker	size : decreased size, size towards gill filament	2	429
entopterygoid	count : absent, absent of , present, present of	2	368
branchiostegal ray 3	shape	2	0

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3. How do skeletal regions vary across taxa?

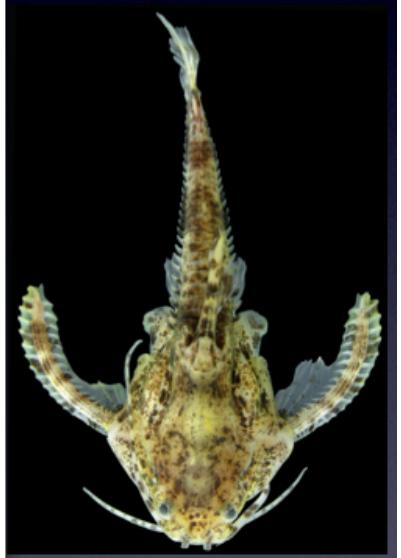
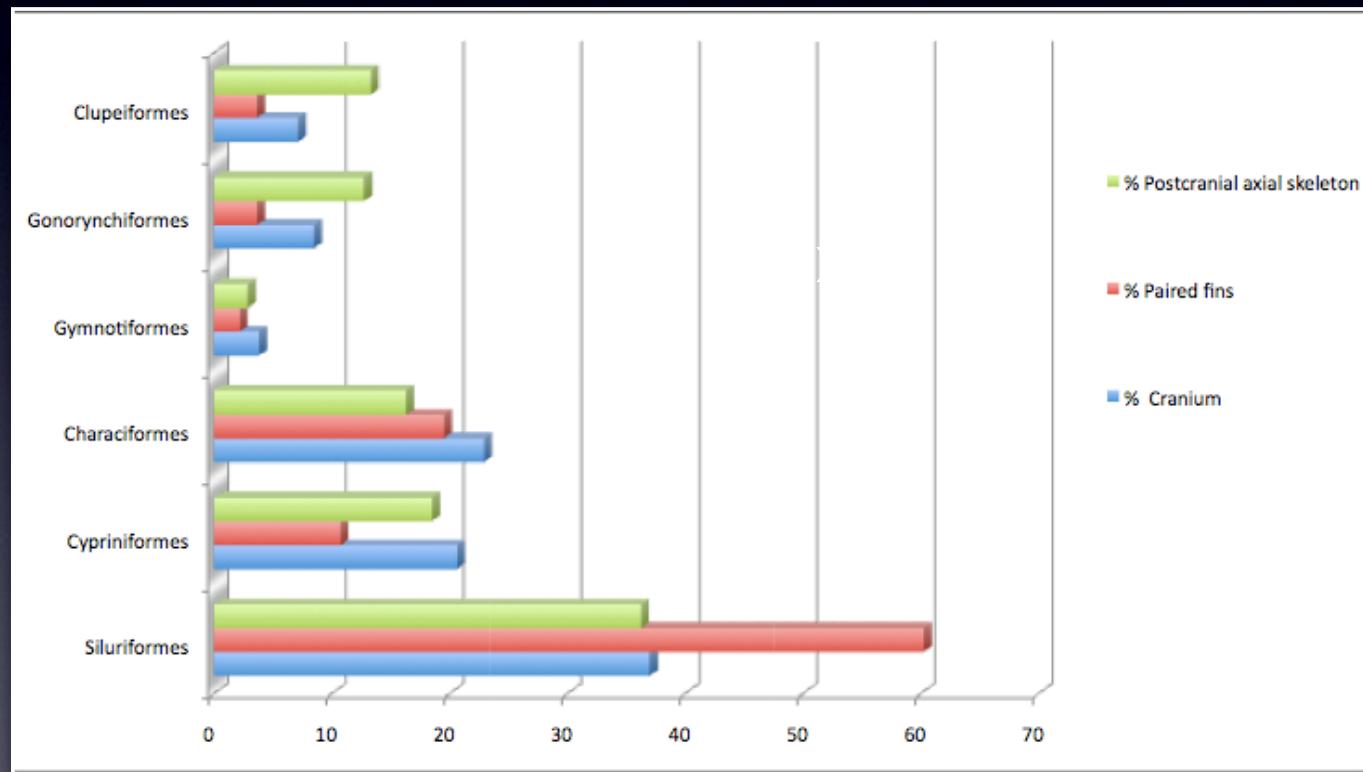


Image from Sabaj-Perez

TAO terms: 'Postcranial axial skeleton', 'Paired fins',
'Cranium'

Other use cases:

- Give me all the bones that come from neural crest vs. mesoderm
- Is there an enrichment in the neural crest vs. mesodermally derived bones in a particular clade?

What we can NOT ask for currently:

- Give me a list:
 - of all the **intramembranous bones** in a mouse or *Xenopus* or any amphibian
 - of all the **dermal bones, replacement bones** in a mouse or *Xenopus* or any amphibian



Purpose of jamboree:

- Re-evaluate skeletal ontology for fishes (TAO), focussing at level of cell, tissue, and skeletal element
- Goal to improve usefulness and accuracy of reasoning that is based on this ontology
 - to facilitate answering questions, ‘use-driven’ ontology

Re-evaluate TAO within context of other vertebrate anatomy ontologies

Teleost Anatomy
Ontology
(2371 terms; 618 skeletal)

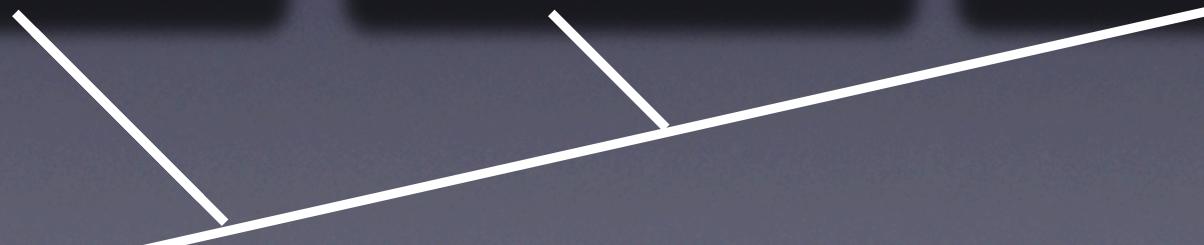
Amphibian Anatomy
Ontology

Adult Mouse Anatomy
Ontology

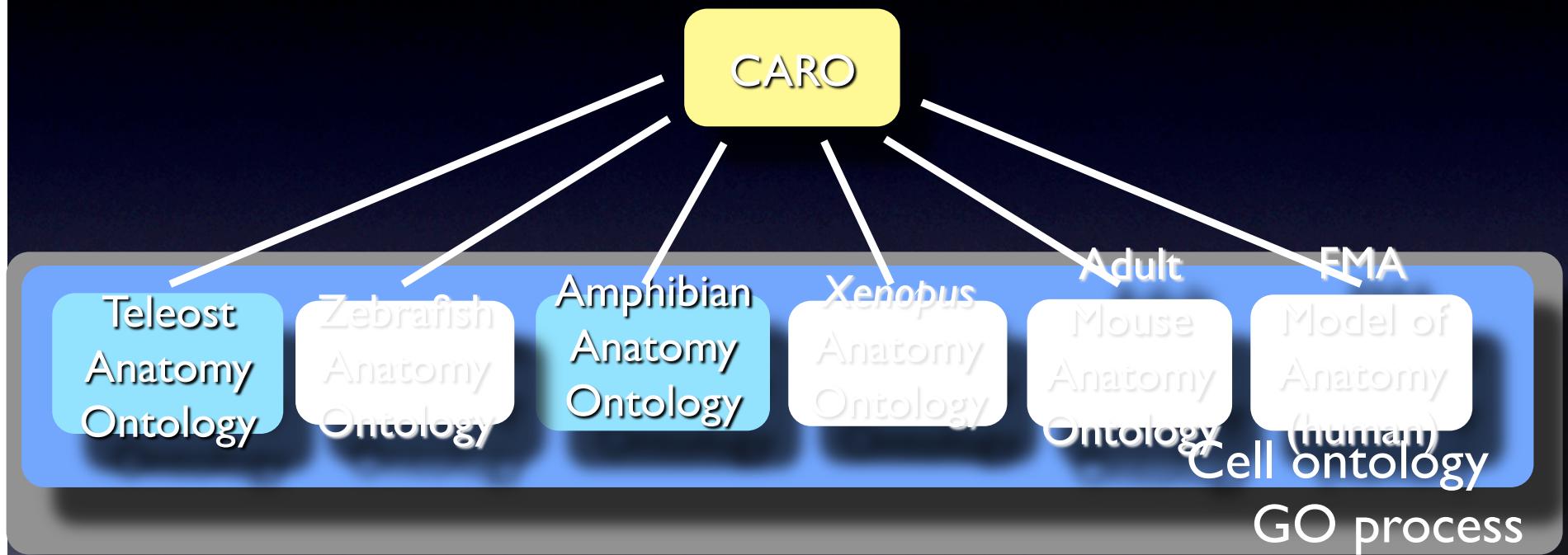
Zebrafish Anatomy
Ontology
(2196 terms; 310 skeletal)

Xenopus Anatomy
Ontology

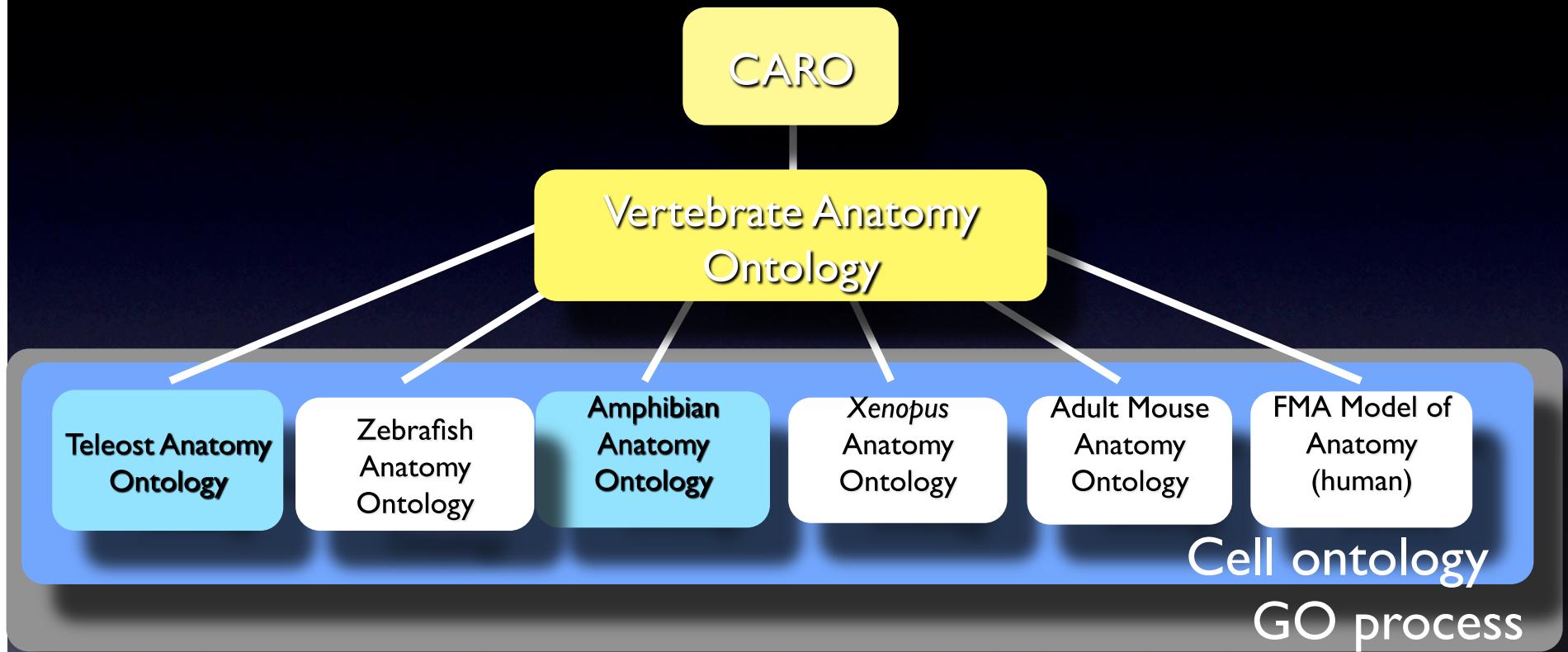
Foundational Model of
Anatomy (human)



Re-evaluate TAO in relation to other relevant ontologies



Vertebrate Anatomy Ontology





Acknowledgements

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