

Creation and unification of development and life stage ontologies for animals

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et évolution



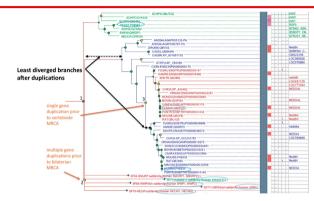




The necessity for multi-species comparative approaches

Multi-species data integration

PANTHER: GO annotations using phylogenetic trees

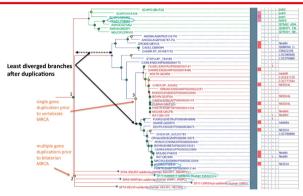


Mi H., et al., Nucleic Acids Research, 2010

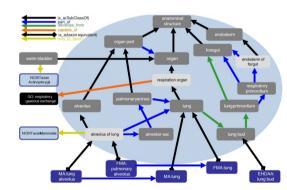
Multi-species data integration

PANTHER: GO annotations using phylogenetic trees

UBERON: comparative anatomy



Mi H., et al., Nucleic Acids Research, 2010



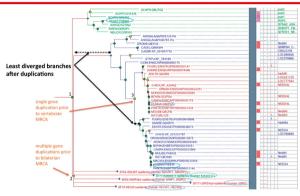
Mungall C.J., et al., Genome Biol, 2012

Multi-species data integration

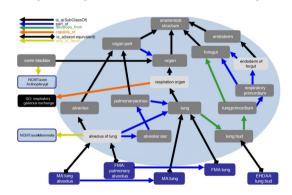
PANTHER: GO annotations using phylogenetic trees

UBERON: comparative anatomy

Need for comparative developmental and life staging



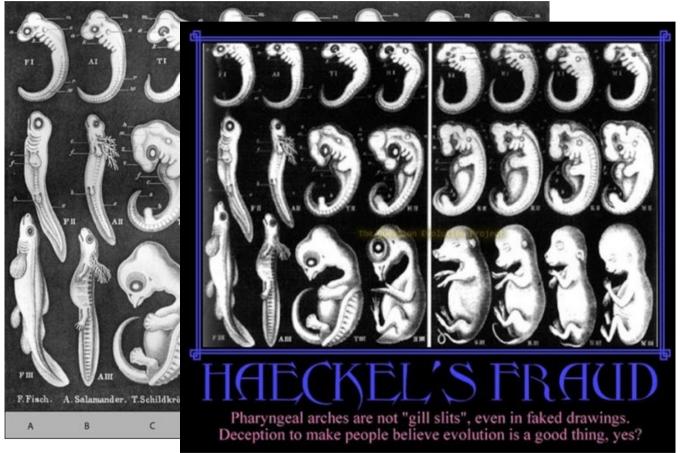
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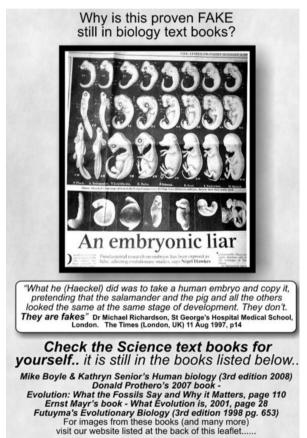


Mungall C.J., et al., Genome Biol, 2012



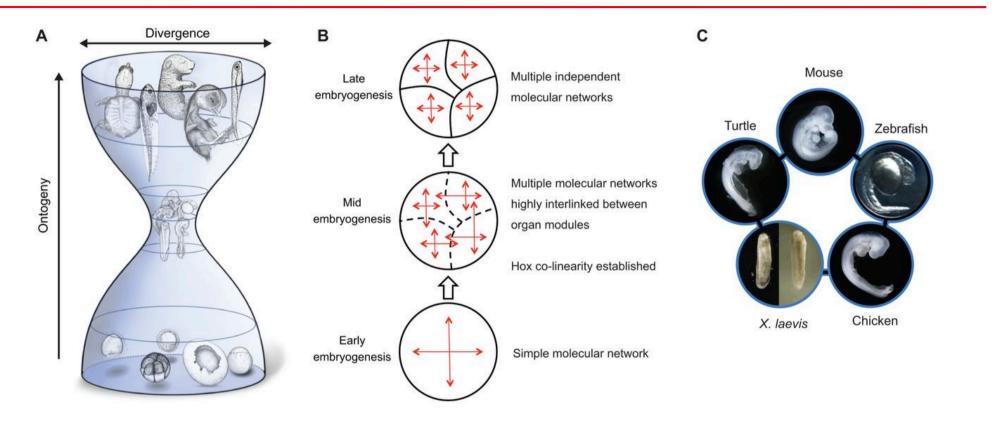
Haeckel, Natürliche Schöpfungsgeschichte, 1868



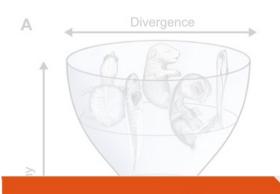


Haeckel, Natürliche Schöpfungsgeschichte, 1868

Watts et al., Theory in Biosciences, 2019



Irie N., et al., Development, 2014, adapted from Wang et al. 2013



PLOS GENETICS

The Hourglass and the Early Conservation Models—Co-Existing Patterns of Developmental Constraints in Vertebrates

Barbara Piasecka, Paweł Lichocki, Sébastien Moretti, Sven Bergmann 🚥, Marc Robinson-Rechavi 🚥 🖂

Published: April 25, 2013 • https://doi.org/10.1371/journal.pgen.1003476





Multiple molecular networks



Inter-embryo gene expression variability recapitulates the hourglass pattern of evo-devo

Jialin Liu ☑, Michael Frochaux, Vincent Gardeux, Bart Deplancke & Marc Robinson-Rechavi ☑

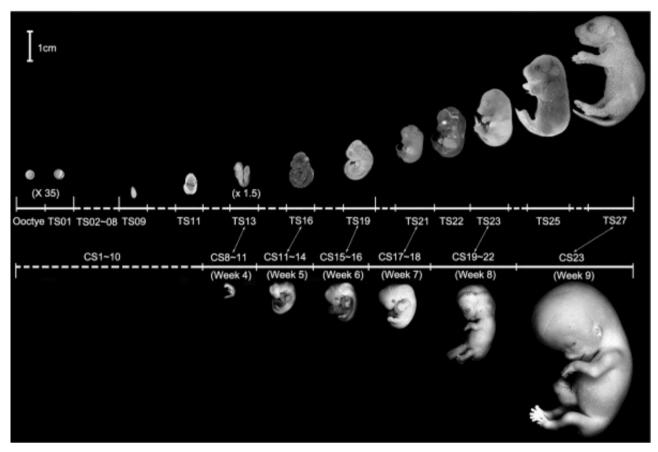
BMC Biology 18, Article number: 129 (2020) Cite this article



The hourglass model of evolutionary conservation during embryogenesis extends to developmental enhancers with signatures of positive selection

Jialin Liu^{1,2,4}, Rebecca R. Viales³, Pierre Khoueiry^{3,5}, James P. Reddington³, Charles Girardot³, Eileen E.M. Furlong³ and Marc Robinson-Rechavi^{1,2}

Irie N., et al., Development, 2014, adapted from Wang et al. 2013

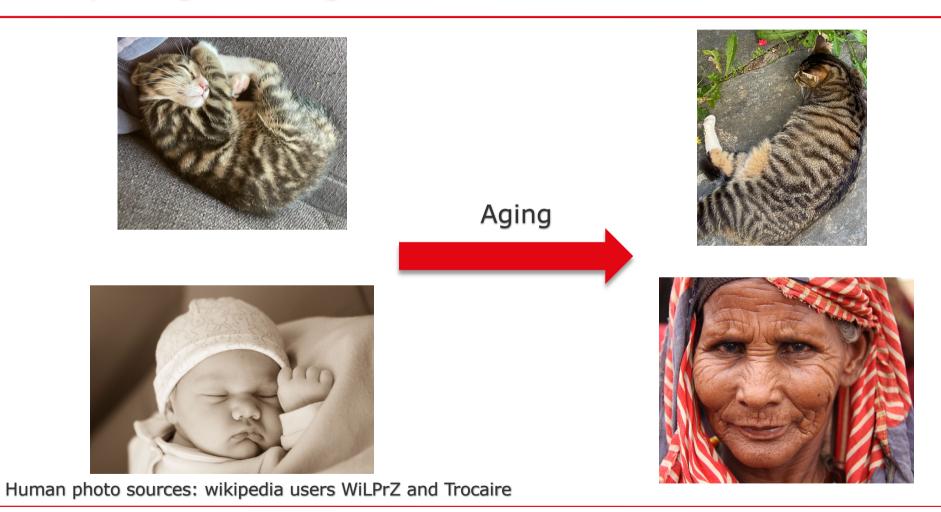


Mouse Theiler stages based on somite number and characteristics

Carnegie stages standardized 23 stages based on development of structures

Xue L., et al., BMC Genomics, 2013

Comparing life stages



Developmental stages capture phenotypic features

Ages during postembryo are not based on phenotypic features

Development is heterochronous



Multi-species core development and life stage ontology

UBERON:0000104 life cycle

UBERON:0000068 embryo stage

UBERON:0000092 post-embryonic stage -

UBERON:0000068 embryo stage

UBERON:0000106 zygote stage

UBERON:0000107 cleavage stage

UBERON:0000108 blastula stage

UBERON:0000109 gastrula stage

UBERON:0000110 neurula stage

UBERON:0000111 organogenesis stage

UBERON:0004707 pharyngula stage

UBERON:0007220 late embryonic stage

UBERON:0000092 post-embryonic stage

UBERON:0000069 larval stage

. . .

UBERON:0000070 pupal stage

UBERON:0000066 fully formed stage

UBERON:0000112 sexually immature stage

UBERON:0018685 nursing stage

UBERON:0007221 neonate stage

UBERON:0035946 start of neonate stage

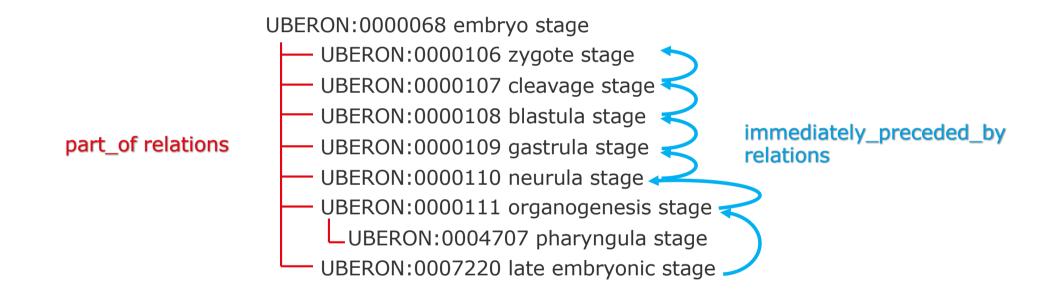
UBERON:0034920 infant stage

UBERON:0034919 juvenile stage

UBERON:0000113 post-juvenile

UBERON:0018241 prime adult stage

UBERON:0007222 late adult stage



```
UBERON:0000092 post-embryonic stage
    UBERON:0000069 larval stage
                                                      only_in_taxon Holometabola
    UBERON:0000070 pupal stage
    UBERON:0000066 fully formed stage
       UBERON:0000112 sexually immature stage
                                                     only_in_taxon Mammalia
         UBERON:0018685 nursing stage
           UBERON:0007221 neonate stage
              UBERON: 0035946 start of neonate stage
                                                     only in taxon Mammalia
           UBERON: 0034920 infant stage
         UBERON:0034919 juvenile stage
       UBERON:0000113 post-juvenile
         UBERON:0018241 prime adult stage
         UBERON:0007222 late adult stage
```

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UBERON:0035946 start of neonate stage

UBERON:0034920 infant stage ■

UBERON:0034919 juvenile stage

UBERON:0000113 post-juvenile

UBERON:0018241 prime adult stage

UBERON:0007222 late adult stage

A mammalian developmental stage that covers the period from birth until weaning.

ends at weaning

The stage of being no more dependent of the nest and/or from caregivers for subsistence while having not reach sexual maturity.



Species-specific ontologies

```
HsapDv:0000264 sexually immature stage
  HsapDv:0000260 nursing stage
     HsapDv:0000262 newborn stage
    HsapDv:0000261 infant stage
       HsapDv:0000273 1-month-old stage
       HsapDv:0000184 11-month-old stage
  HsapDv:0000265 child stage
     HsapDv:0000246 1-year-old stage
       HsapDv:0000185 12-month-old stage
       HsapDv:0000196 23-month-old stage
    HsapDv:0000270 2-4 year-old child stage
       HsapDv:0000096 2-year-old stage
       HsapDv:0000097 3-year-old stage
       HsapDv:0000098 4-year-old stage
  HsapDv:0000271 juvenile stage
    HsapDv:0000099 5-year-old stage
    HsapDv:0000108 14-year-old stage
```

```
[Term]
id: HsapDv:0000261
name: infant stage
namespace: human developmental stage
def: "Immature stage that refers to an infant who is over 28 days and is under 12 months old." [Bgee:curator "Bgee"]
comment: We consider that this pediatric stage refers to an infant who is over 28 days and under 12 months old, to follow
HPO (Human Phenotype Ontology) which considers infant as between 28 days to one year of life, based on the ethymology of
the word (infant), meaning not speaking, see HP:0003593 (Infantile onset) and see Peter Robinson's comment
[[HPO:probinson] at https://github.com/obophenotype/human-phenotype-ontology/issues/5119#issuecomment-578882539.
synonym: "infantile stage" EXACT []
synonym: "under-1-year-old stage" EXACT []
xref: HP:0003593
xref: UBERON:0034920
is_a: HsapDv:0000000! life cycle stage
relationship: part_of HsapDv:0000260! nursing stage
relationship: immediately preceded by HsapDv:0000262! newborn stage
property value: start ypb "0.077" xsd:float
property value: end ypb "1.0" xsd:float
property value: start mpb "0.92" xsd:float
property_value: end_mpb "12.0" xsd:float
```

```
HsapDv:0000264 sexually immature stage
  HsapDv:0000260 nursing stage
     HsapDv:0000262 newborn stage
                                                     Based on speach acquisition
    HsapDv:0000261 infant stage
                                                     characteristic
       HsapDv:0000273 1-month-old stage
                                                     Age range
       HsapDv:0000184 11-month-old stage
  HsapDv:0000265 child stage
     HsapDv:0000246 1-year-old stage
       HsapDv:0000185 12-month-old stage
       HsapDv:0000196 23-month-old stage
    HsapDv:0000270 2-4 year-old child stage
       HsapDv:0000096 2-year-old stage
       HsapDv:0000097 3-year-old stage
       HsapDv:0000098 4-year-old stage
  HsapDv:0000271 juvenile stage
     HsapDv:0000099 5-year-old stage
    HsapDv:0000108 14-year-old stage
```

[Term]

id: HsapDv:0000260 name: nursing stage

...

comment: The definition of an age for human weaning is difficult to define due to individual and cultural variations. The natural age of weaning in humans could be 2.5 years, with maximum 7.0 years (see https://www.health-e-learning.com/articles/A_Natural_Age_of_Weaning.pdf), but we decide to stop this period at 1 year-old, before usually language acquisition.

HsapDv:0000226 prime adult stage
HsapDv:0000266 young adult stage
HsapDv:0000268 15-19 year-old
HsapDv:0000109 15-year-old stage
...
HsapDv:0000113 19-year-old stage
HsapDv:0000237 third decade stage
HsapDv:0000114 20-year-old stage
...
HsapDv:0000123 29-year-old stage
HsapDv:0000238 fourth decade stage
HsapDv:0000124 30-year-old stage
...

HsapDv:0000133 39-year-old stage

HsapDv:0000267 middle aged stage
HsapDv:0000239 fifth decade stage
HsapDv:0000134 40-year-old tage
...
HsapDv:0000143 49-year-old stage
HsapDv:0000240 sixth decade stage
HsapDv:0000144 50-year-old stage
...
HsapDv:0000153 59-year-old stage



Alignment with Uberon

Human ontology: bridge file

[Term]

id: HsapDv:0000045

property_value: IAO:0000589 "prenatal stage (human)" xsd:string

intersection_of: UBERON:0000068

intersection_of: part_of NCBITaxon:9606

[Term]

id: HsapDv:0000262

property_value: IAO:0000589 "newborn stage (human)" xsd:string

intersection_of: UBERON:0007221

intersection_of: part_of NCBITaxon:9606

[Term]

id: HsapDv:0000192

property_value: IAO:0000589 "19-month-old stage (human)" xsd:string

relationship: only in taxon NCBITaxon:9606! human

...

Human ontology: merge with Uberon

UBERON:0000104 life cycle

UBERON:000068 embryo stage UBERON:0000106 zygote stage UBERON:0000107 cleavage stage

HsapDv:0000205 morula stage (human)

UBERON:0000108 blastula stage

HsapDv:0000007 Carnegie stage 03 (human) HsapDv:0000008 Carnegie stage 04 (human) HsapDv:0000009 Carnegie stage 05 (human) HsapDv:0000031 Carnegie stage 05a (human)

HsapDv:0000031 Carnegie stage 03a (Human) HsapDv:0000033 Carnegie stage 05b (human)

UBERON:0000109 gastrula stage

HsapDv:0000011 Carnegie stage 06 (human) HsapDv:0000034 Carnegie stage 06a (human) HsapDv:0000035 Carnegie stage 06b (human)

UBERON:0000110 neurula stage

HsapDv:0000013 Carnegie stage 07 (human) HsapDv:0000014 Carnegie stage 08 (human)

UBERON:0000111 organogenesis stage

HsapDv:0000016 Carnegie stage 09 (human)
HsapDv:0000017 Carnegie stage 10 (human)
HsapDv:0000018 Carnegie stage 11 (human)
HsapDv:0000019 Carnegie stage 12 (human)
HsapDv:0000020 Carnegie stage 13 (human)
HsapDv:0000021 Carnegie stage 14 (human)
HsapDv:0000022 Carnegie stage 15 (human)
HsapDv:0000023 Carnegie stage 16 (human)
HsapDv:0000024 Carnegie stage 17 (human)
HsapDv:0000025 Carnegie stage 18 (human)
HsapDv:0000026 Carnegie stage 19 (human)
HsapDv:0000027 Carnegie stage 20 (human)
HsapDv:0000028 Carnegie stage 21 (human)
HsapDv:00000029 Carnegie stage 22 (human)
HsapDv:00000030 Carnegie stage 23 (human)

Integration of 21 species

Ontology nam	e Species	Number of	Number of classes
4 5		classes	linked to Uberon
AcarDv	Anolis carolinensis	20	20
BtauDv	Bos taurus	78	8
CfamDv	Canis lupus familiaris	24	24
CporDv	Cavia porcellus	24	24
DpseDv	Drosophila pseudoobscura	21	18
DsimDv	Drosophila simulans	21	18
EcabDv	Equus caballus	24	24
FcatDv	Felis catus	24	24
GgalDv	Gallus gallus	90	16
GgorDv	Gorilla gorilla	47	24
HsapDv	Homo sapiens	238	20
MdomDv	Monodelphis domestica	59	24
MmulDv	Macaca mulatta	66	24
MmusDv	Mus musculus	134	20
OanaDv	Ornithorhynchus anatinus	24	24
OariDv	Ovis aries	33	20
OcunDv	Oryctolagus cuniculus	24	24
PpanDv	Pan paniscus	56	24
PtroDv	Pan troglodytes	67	24
RnorDv	Rattus norvegicus	60	20
SscrDv	Sus scrofa	94	22
	2.50 501010	- •	



Application examples

Annotation in Human Cell Atlas



Search all filters Donor Tissue Type Specimen Method File **Genus Species** Organism Age Range Biological Sex Donor Disease Development Stage Projects Projects Projects Projects Homo sapiens 242 Age unit week month year female 162 normal 179 ☐ 1-month-old human stage Mus musculus 33 male abscess Min age e.g. 1 10-month-old Unspecified mixed acoustic neuroma human stage Max age e.g. 2 unknown acquired 10th week postaneurysmal fertilization subaráchnoid human stage hemorrhage Apply 11th week postacute kidney fertilization failure human stage acute kidney 12th week post-10 tubular necrosis fertilization analysis_protocol

Expression data in Bgee: PAX6 example



Gene: PAX6 - ENSG00000007372 - Homo sapiens (human)

Function¹

Transcription factor with important functions in the development of the eye, nose, central nervous system and pancreas. Required for the differentiation of pancreatic islet alpha cells (By similarity).

Anatomical entity	Dev. stage	Expression score	FDR	Sources
UBERON:0001812	HsapDv:0000246 ☑ 1-year-old stage (human)	99.55	0.002	AEIRFL
UBERON:0001812	HsapDv:0000270 2-4 year-old child stage (human)	99.55	0.002	AEIRFL
UBERON:0001812	UBERON:0034919	99.53	0.002	AEIRFL
<u>UBERON:0003053</u> ✓ ventricular zone	HsapDv:0000198	99.33	<= 1.00e-14	AEIRFL
CL:0000169	<u>UBERON:0000113</u>	99.28	0.002	AEIRFL

Expression data in Bgee: PAX6 expression in Euteleostomi

Anatomical entities	Conservation score ▼ 1	Max expression score ▼ ³	Genes with presence of expression	Genes with absence of expression ^{▲ 2}	Genes with no data
<u>cornea</u> ☑	1.00	99.72	3 genes	0 gene	48 genes
retinal neural layer ଔ	1.00	99.47	3 genes	0 gene	48 genes
ventricular zone	1.00	99.28	2 genes	0 gene	49 genes
g <u>erm cell</u> ☑	1.00	98.61	4 genes	0 gene	47 genes
metencephalon ଔ	1.00	98.54	24 genes	0 gene	27 genes
hindbrain 🛭	1.00	98.54	26 genes	0 gene	25 genes
<u>cerebellum</u> ☑	1.00	98.54	24 genes	0 gene	27 genes
lens of camera-type eye	1.00	98.11	5 genes	0 gene	46 genes
optic cup 🗹	1.00	97.95	3 genes	0 gene	48 genes
optic fissure	1.00	97.95	1 gene	0 gene	50 genes





https://github.com/obophenotype/developmental-stage-ontologies



http://purl.obolibrary.org/obo/uberon/composite-metazoan.owl

www.sib.swiss