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Stage 16, posterior-dorsal view 18 hr 15 min pf @ 23°C	Stage 17, anterior view 18 hr 45 min pf @ 23°C	Stage 17, posterior-dorsal view 18 hr 45 min pf @ 23°C	Stage 18, anterior view 19 hr 45 min pf @ 23°C
			
Stage 19, dorsal view 20 hr 45 min pf @ 23°C	Stage 19, anterior view 20 hr 45 min pf @ 23°C	Stage 20, dorsal view 21 hr 45 min pf @ 23°C	Stage 20, anterior view 21 hr 45 min pf @ 23°C
			
Stage 21, dorsal view 22 hr 30 min pf @ 23°C	Stage 21, anterior view 22 hr 30 min pf @ 23°C	Stage 22, dorsal view 24 hr pf @ 23°C	Stage 22, lateral view 24 hr pf @ 23°C
			
Stage 23, dorsal view 1 day, 45 minutes pf @ 23°C	Stage 23, lateral view 1 day, 45 minutes pf @ 23°C	Stage 24, dorsal view 1 day, 2 hr 15 min pf @ 23°C	Stage 24, lateral view 1 day, 2 hr 15 min pf @
			
Stage 25, dorsal view 1 day, 3 hr 30 min pf @ 23°C	Stage 25, lateral view 1 day, 3 hr 30 min pf @ 23°C	Stage 26, dorsal view 1 day, 5 hr 30 min pf @ 23°C	Stage 26, lateral view 1 day, 5 hr 30 min pf @

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http://xenbase.org/anatomy/static/xenbasefate.jsp

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Xenbase [2.4]

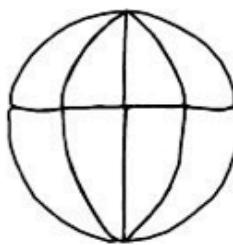
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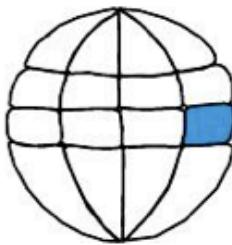
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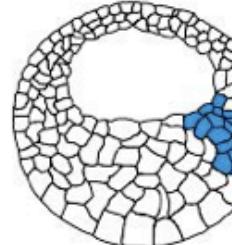
Fate maps: forward

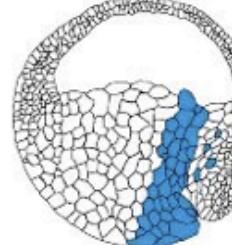
(a [reverse map](#) page is also available)

Move cursor over early blastomeres to see their future positions, [click](#) on blastomeres to read about their later fate.


ventral-----dorsal
animal up, vegetal down


32-cell / stage 6


stage 8


stage 10.5

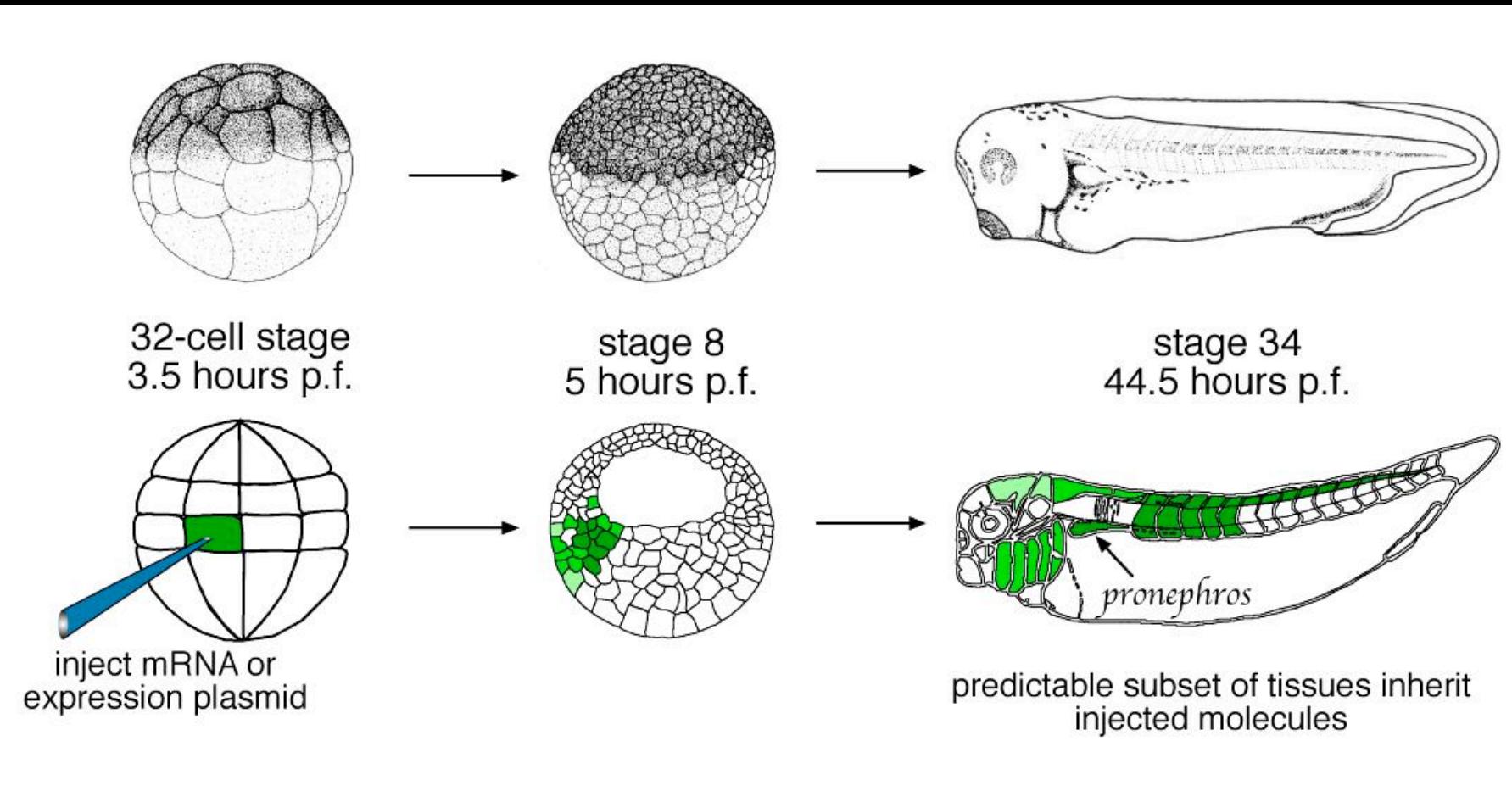
Fate mapping data and images based upon the work of Moody 1987a, 1991 and Bauer et al. 1994.

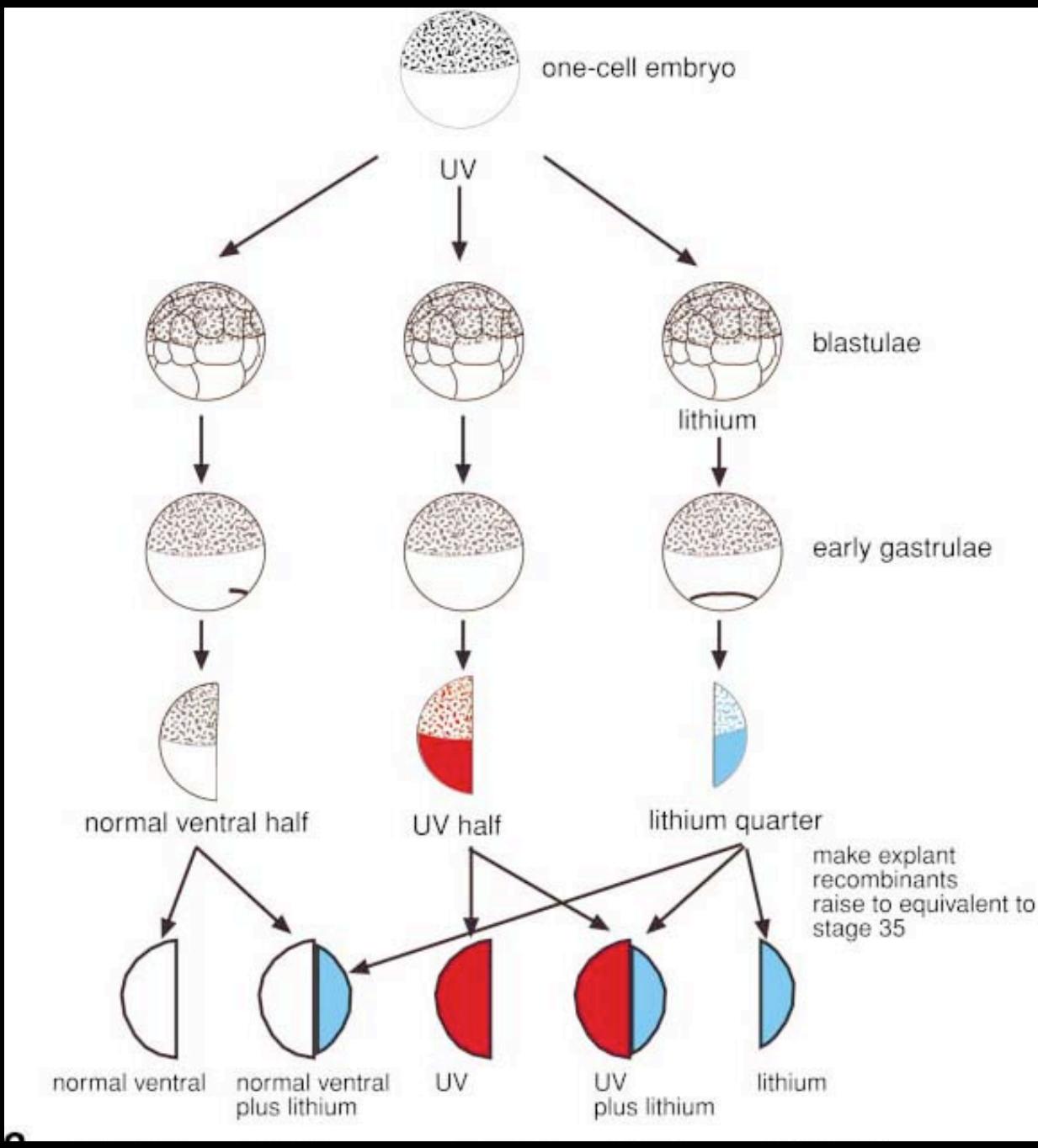
[Technical notes](#)

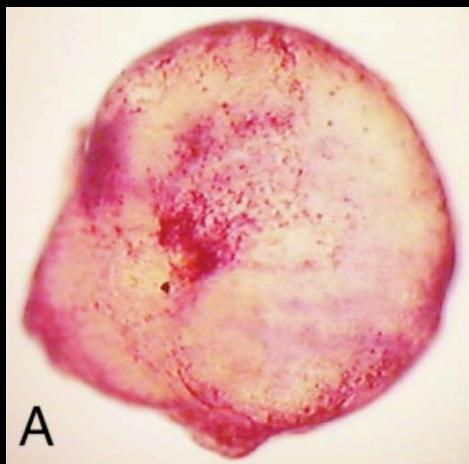
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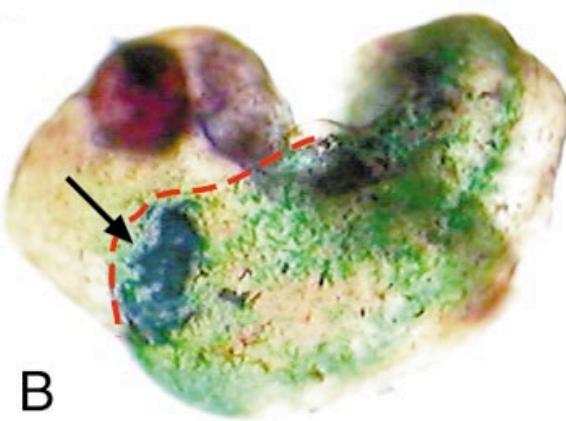






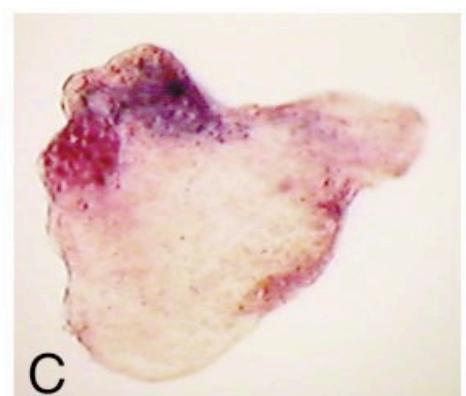
A

UV half embryo



B

recombinant



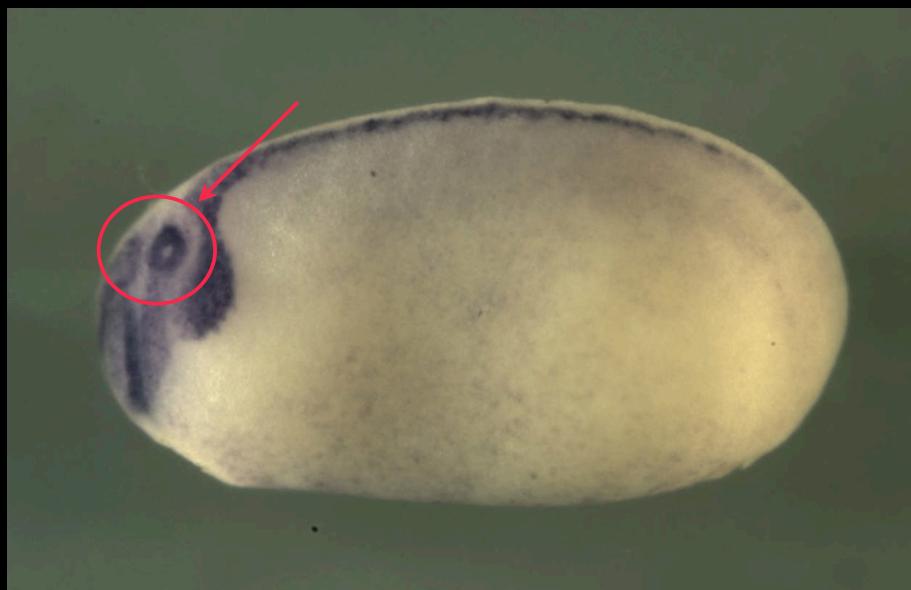
C

lithium quarter embryo

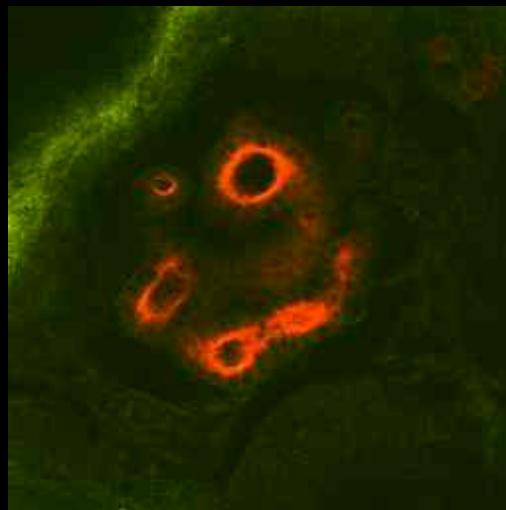
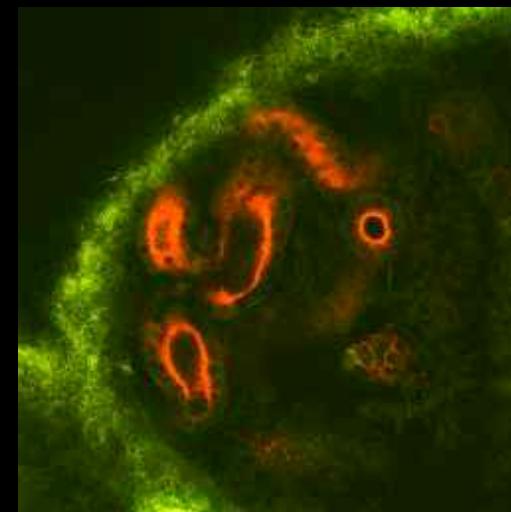
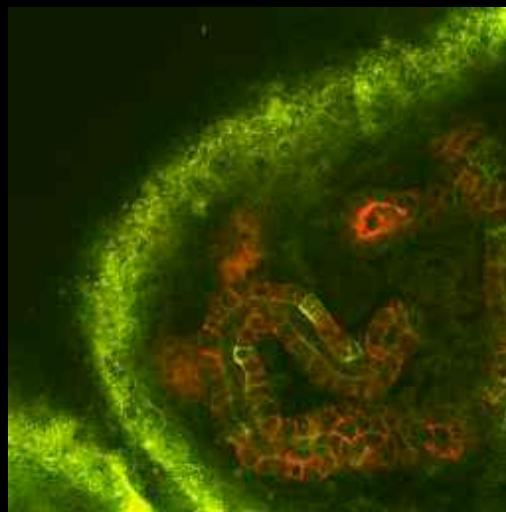
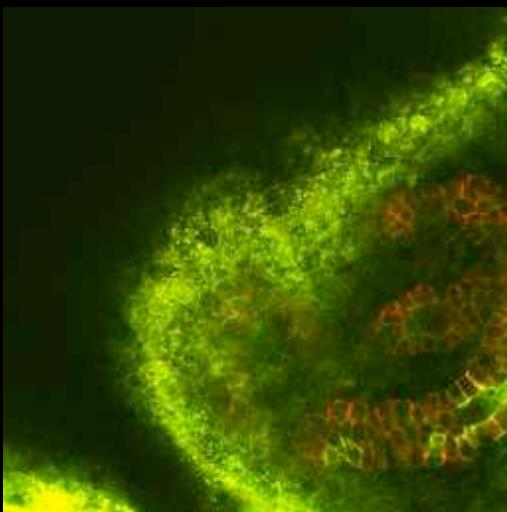
gene A

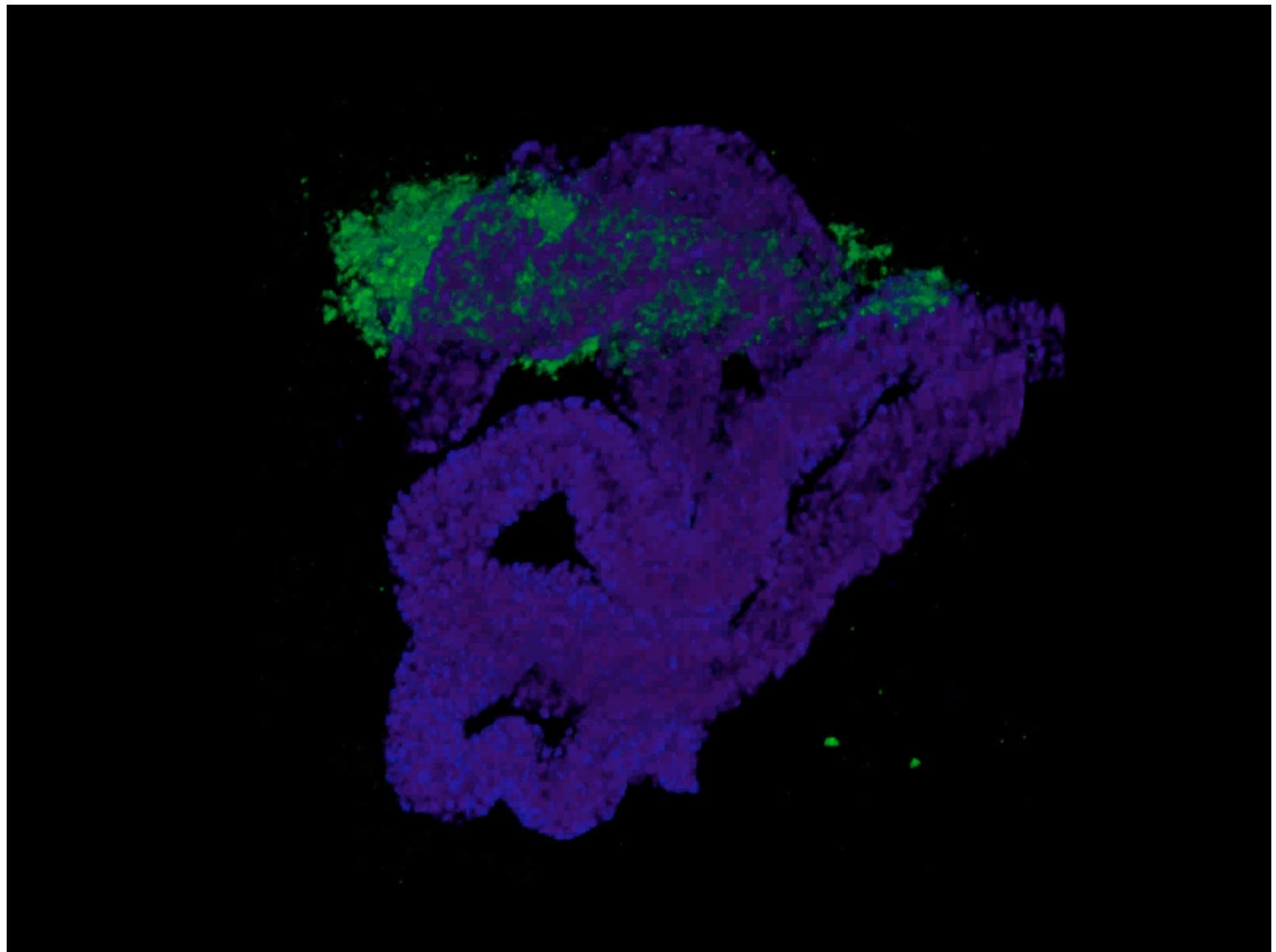


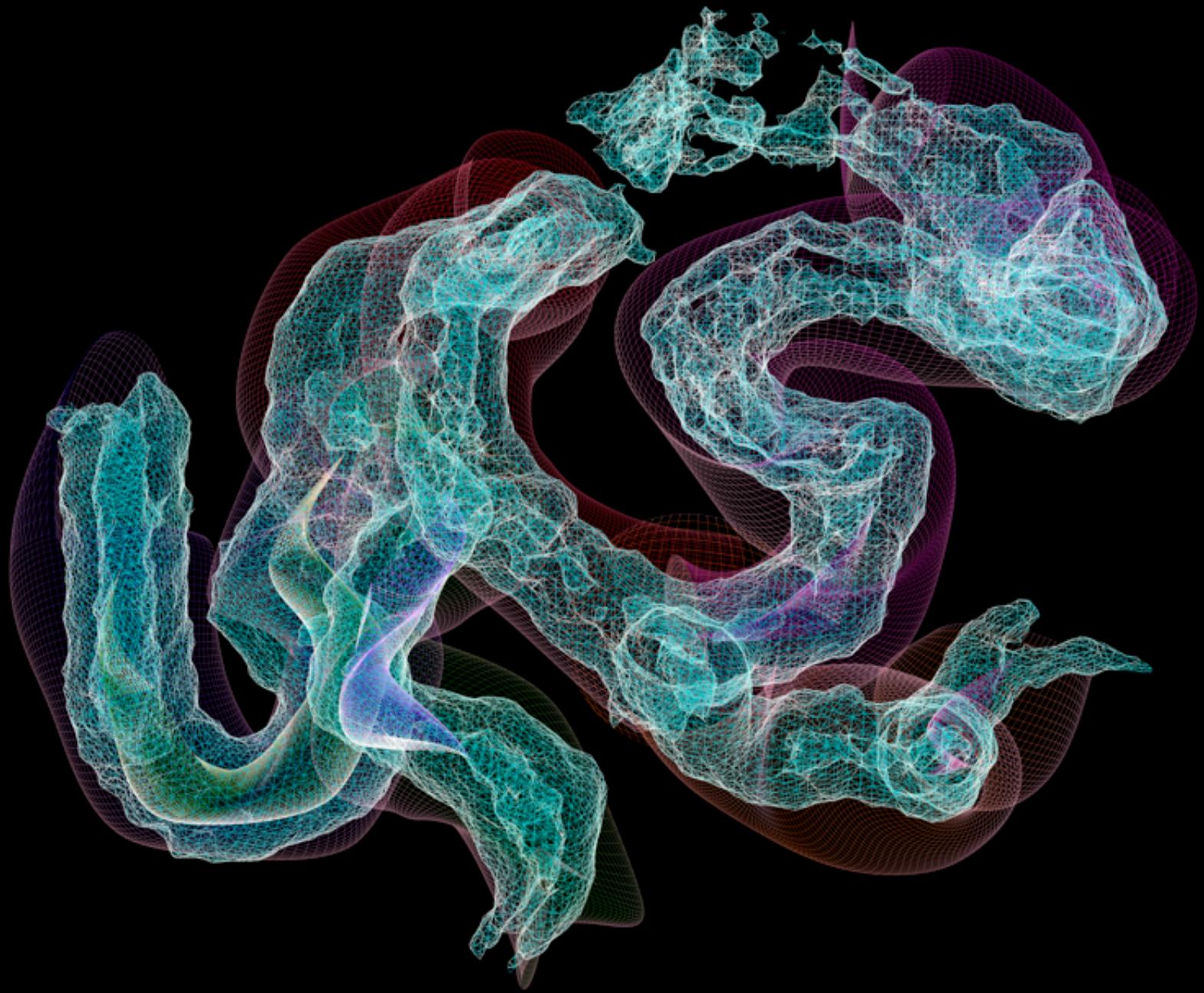
gene B

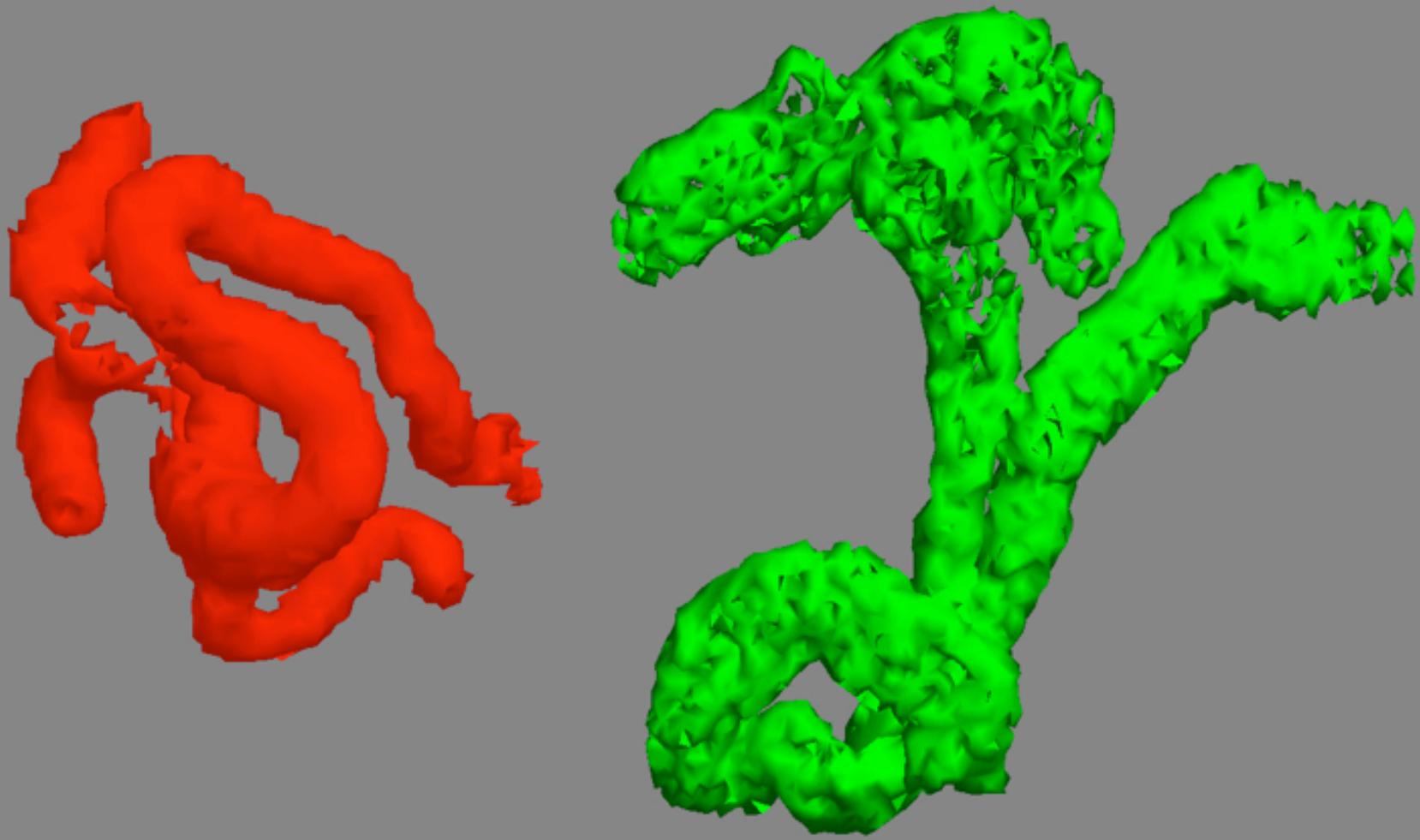








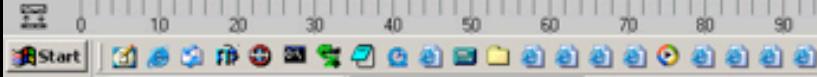




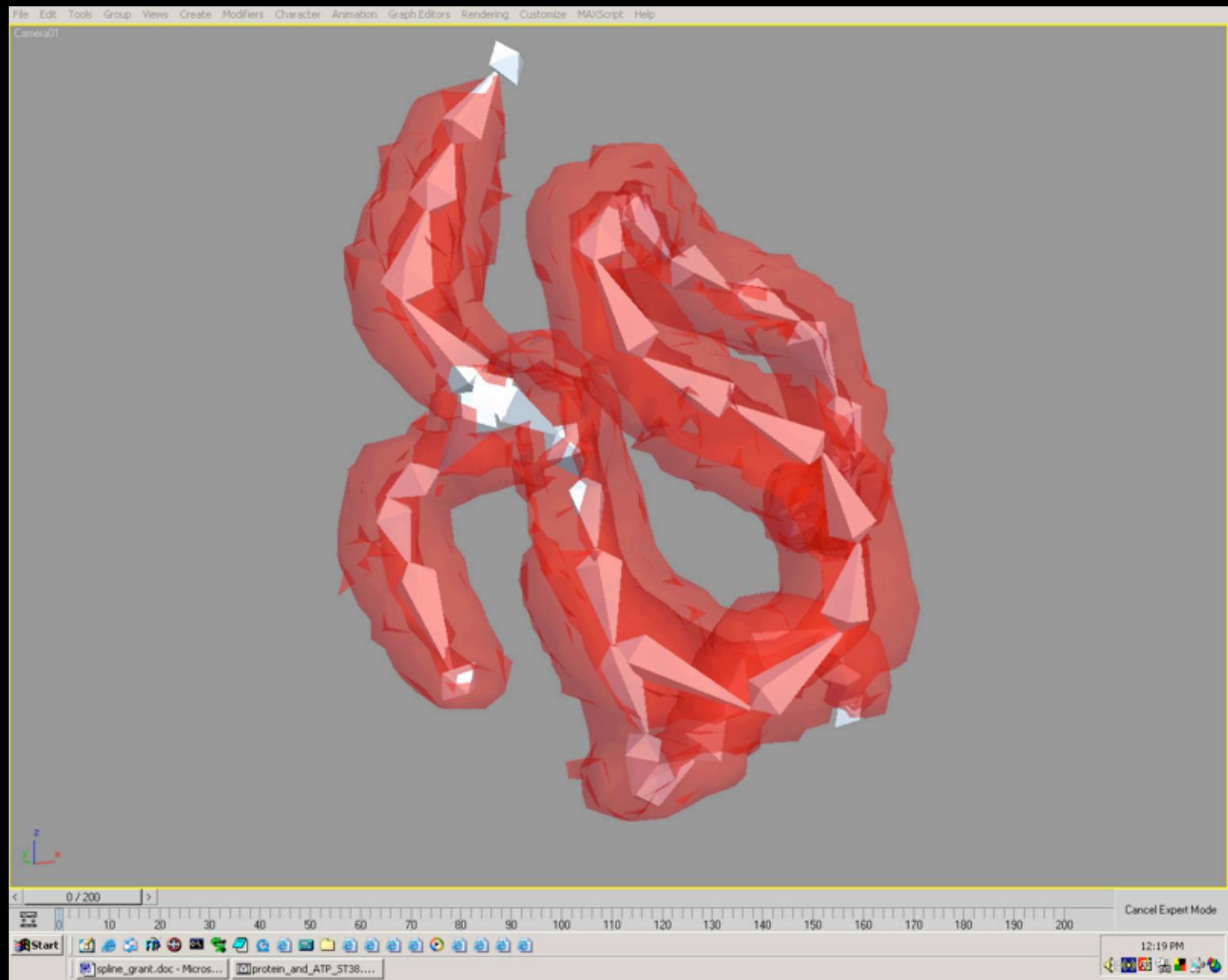
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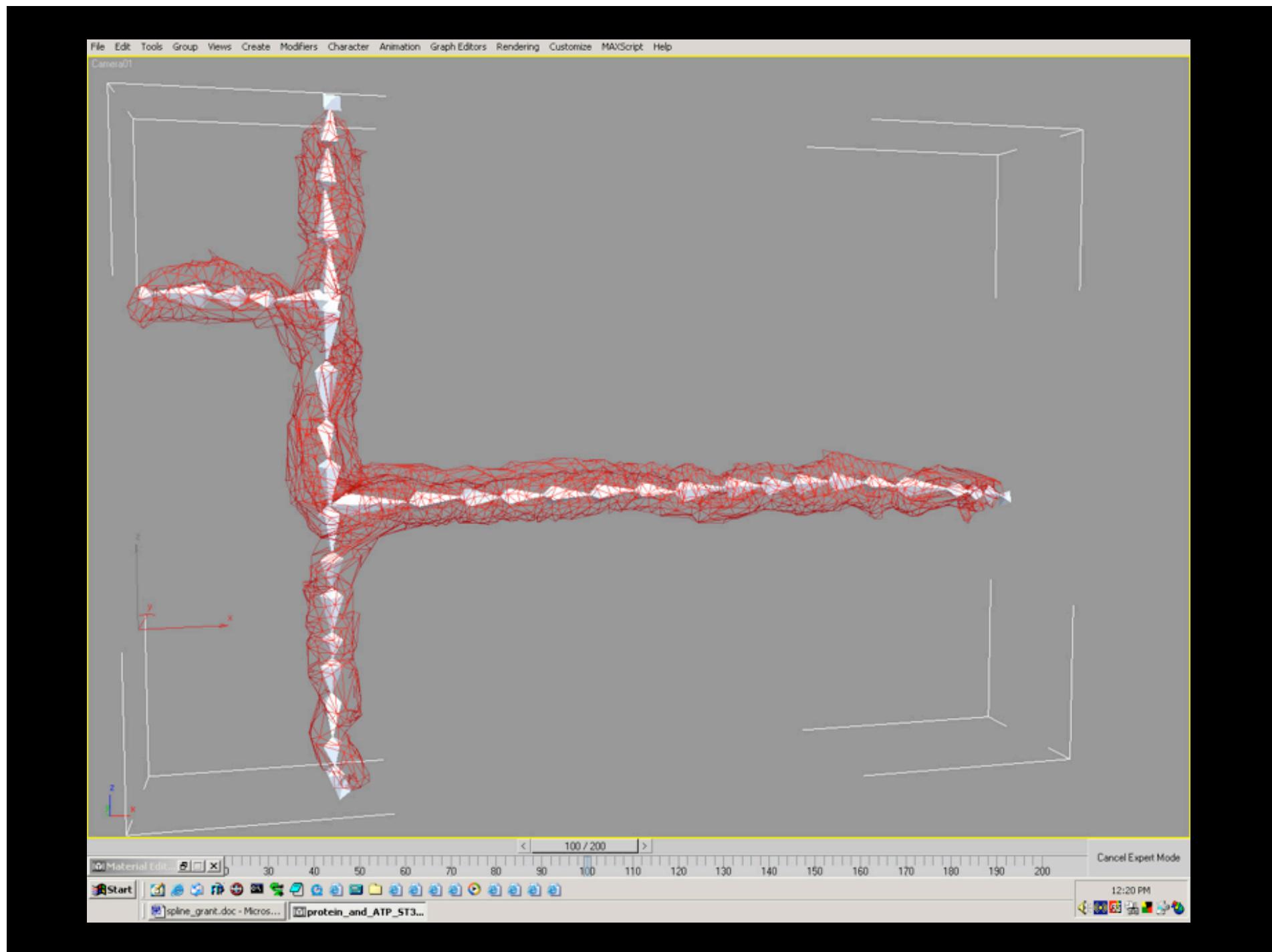
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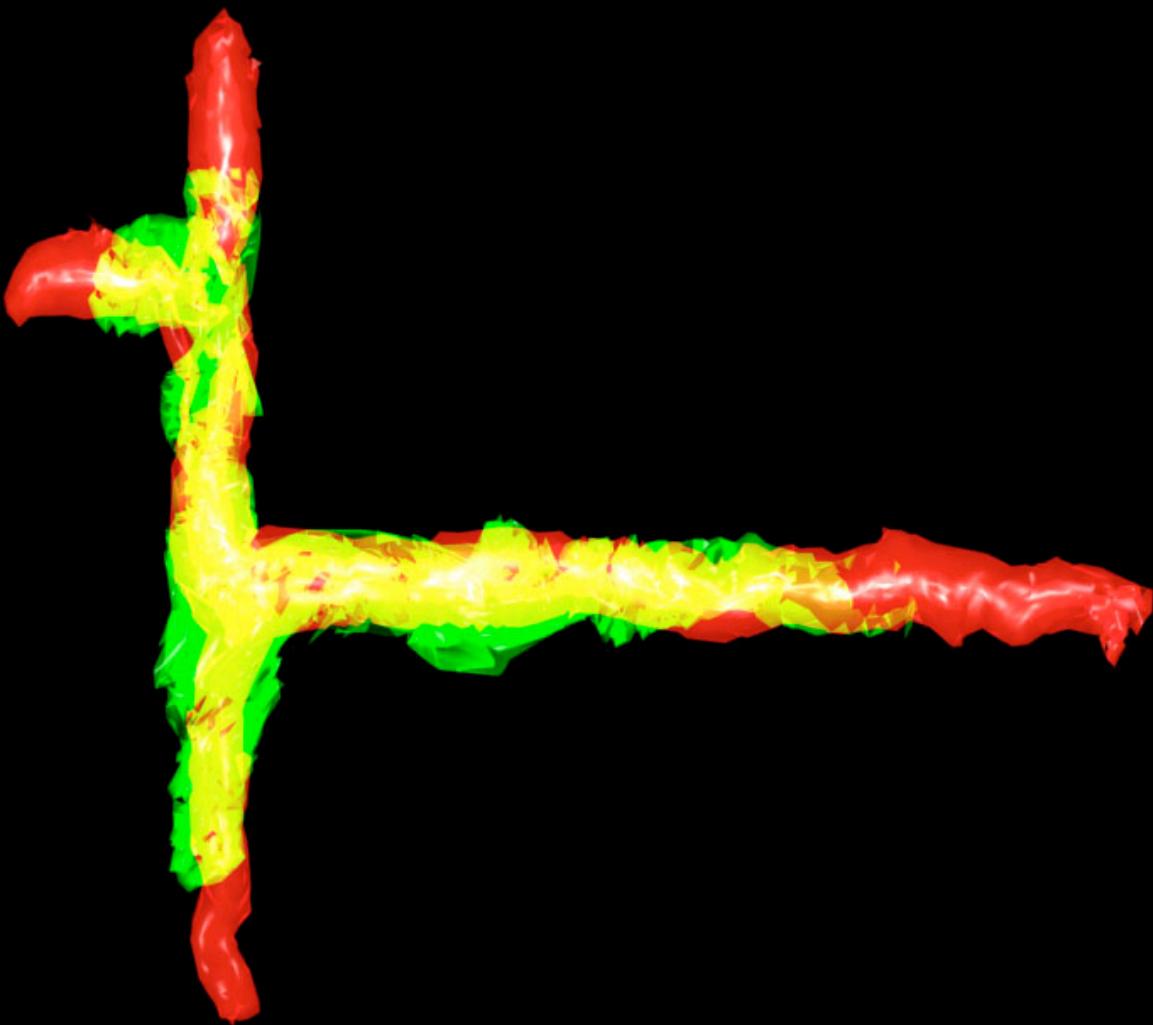
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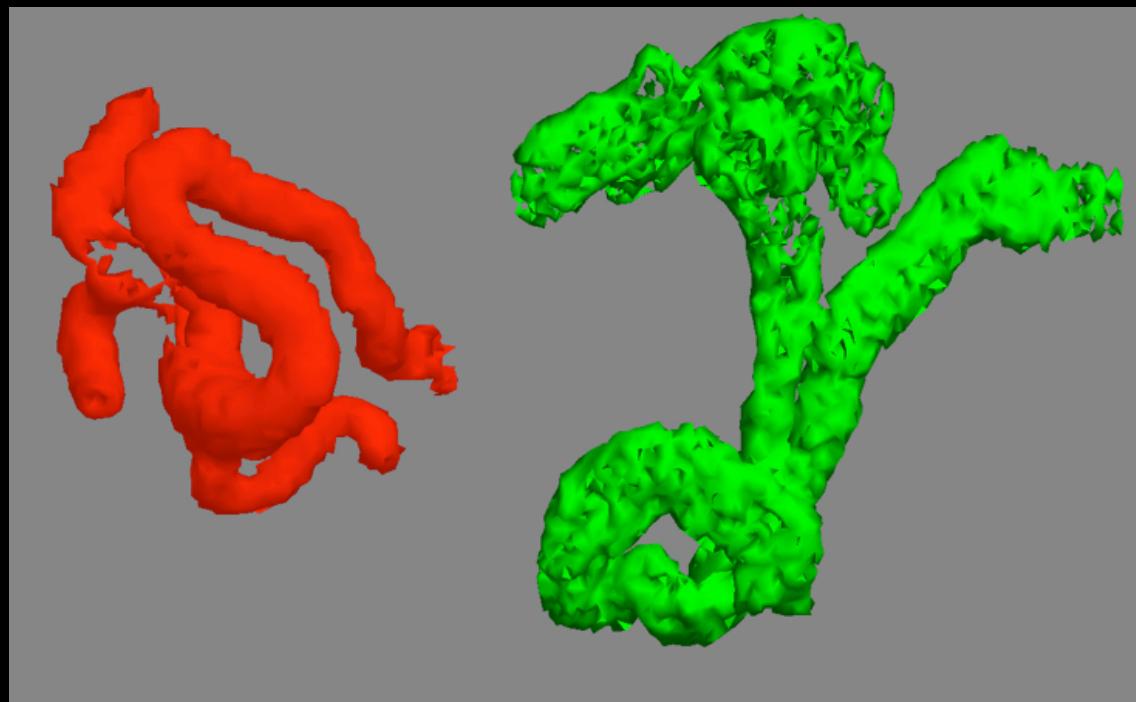


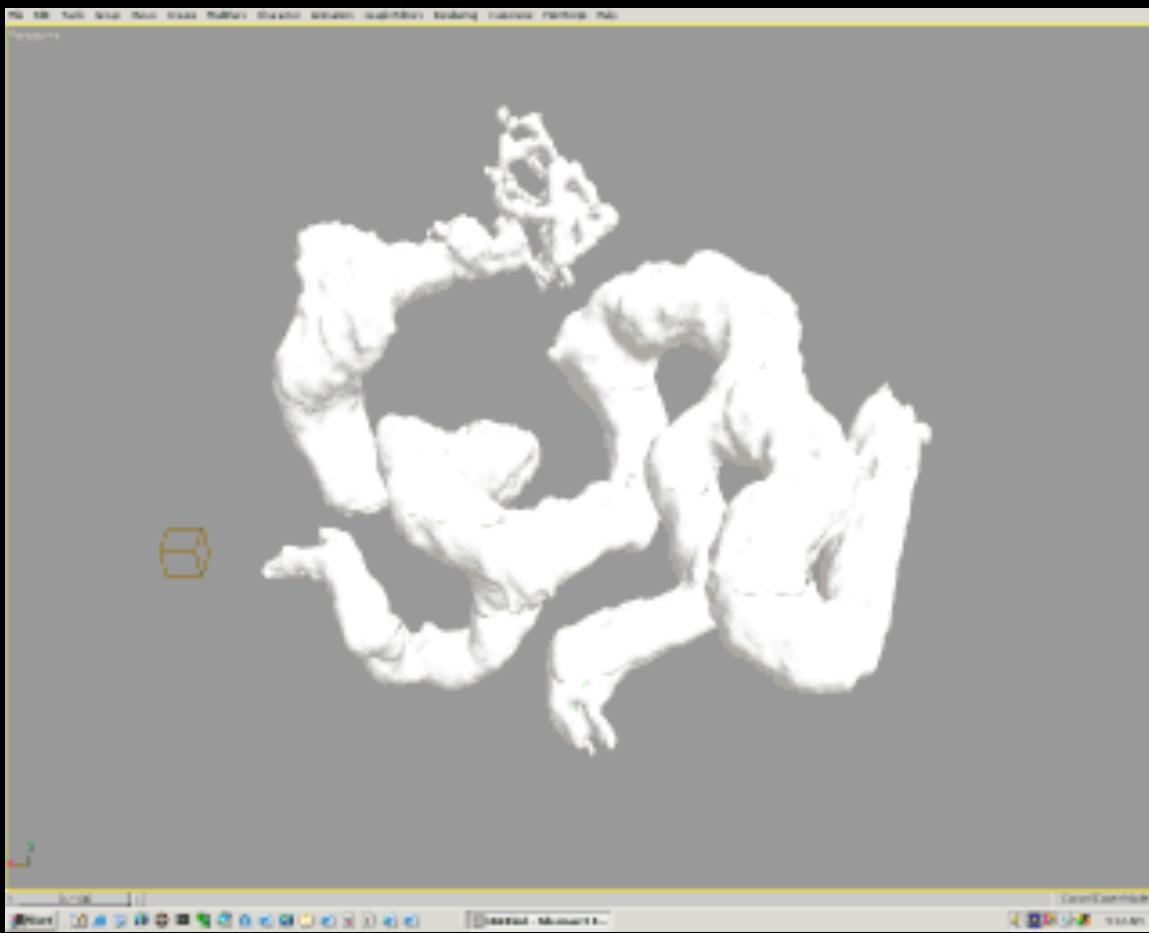


5



reverse time-line to refold





automated spline generation

gene A



ear
kidney

embryo stage: 23

gene B

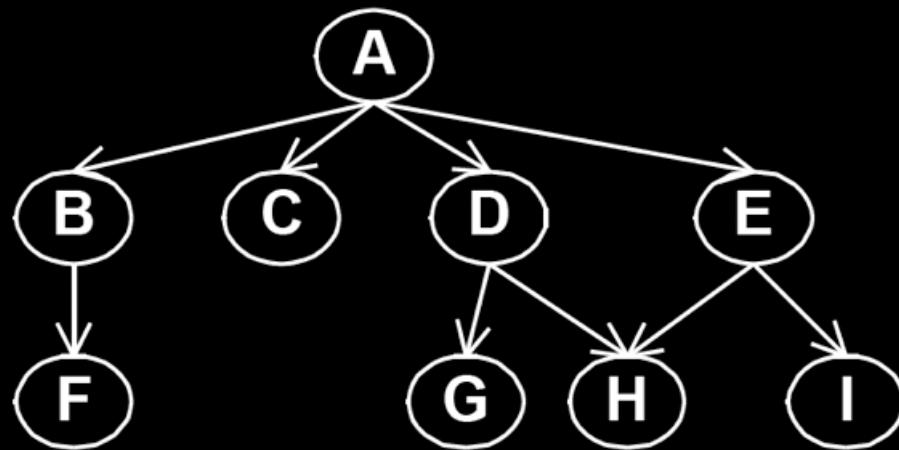


ear
branchial arches
neural crest

embryo stage: 23

Anatomical Ontology

- controlled vocabulary of terms
- map of relationships between terms



[Summary](#) [Expression](#) [Gene Literature \(21\)](#) [Interactants \(105\)](#)

XB-FEAT-483692

Gene Symbol: pax8

Gene Name: paired box 8

Anatomy terms

oviduct, whole organism

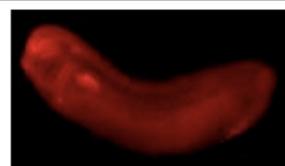
Anatomy stages

NF stage 15 to adult

Community Submitted Images



NF stage 23



unspecified stage

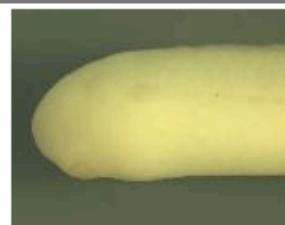
XDB3 & Axel DB Images



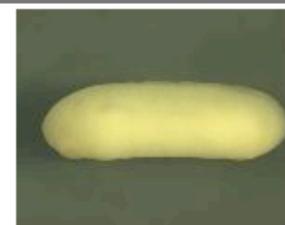
NF stage 28



NF stage 28



NF stage 28



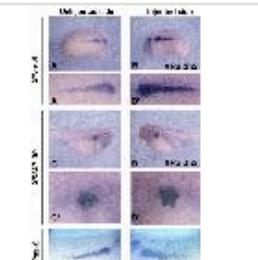
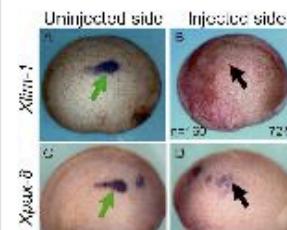
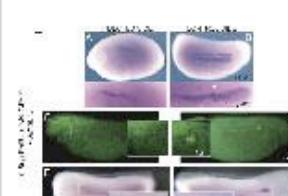
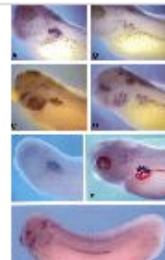
NF stage 28



NF stage 28

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G

XB-ART-12317

Dev Biol. October 1, 1999; 214 (1): 46-59.

Synergism between Pax-8 and lim-1 in embryonic kidney development.

[Carroll TJ](#) , [Vize PD](#) .

Abstract

Pax genes encode a family of highly conserved DNA-binding transcription factors. These proteins play key roles in regulating a number of vertebrate and invertebrate developmental processes. Mutations in Pax-6 result in eye defects in flies, mice, and humans, and ectopic expression of this gene can trigger development of ectopic compound eyes in flies. Likewise, mutation of other Pax genes in vertebrates results in the failure of specific differentiation programs. Pax-1 causes skeletal defects; Pax-2, kidney defects; Pax-3 or Pax-7, neural crest defects; Pax-4, pancreatic beta-cell defects; Pax-5, B-cell defects; Pax-6, thyroid defects; and Pax-9, tooth defects. Although this class of genes is obviously required for the normal differentiation of a number of distinct organs, they have not previously been demonstrated to be capable of directing the embryonic development of organs in vertebrates. In this report, it is demonstrated that Pax-8 plays such a role in the establishment of the Xenopus embryonic kidney, the pronephros. However, in order to efficiently direct cells to form pronephric kidneys, XPax-8 requires cofactors, one of which may be the homeobox transcription factor Xlim-1. These two genes are initially expressed in overlapping domains in late gastrulae, and cells expressing both genes will go on to form the kidney. Ectopic expression of either gene alone has a modest effect on pronephric patterning, while coexpression of XPax-8 plus Xlim-1 results in the development of embryonic kidneys of up to five times normal size and also leads to the development of ectopic pronephric tubules. This effect was synergistic rather than additive. XPax-2 can also synergize with Xlim-1. The expression profile of this gene indicates that it normally functions later in pronephric development than does XPax-8. Together these data indicate that the interaction between XPax-8 and Xlim-1 is a key early step in the establishment of the pronephric primordium.

Pubmed Id: [10491256](#)**Article link:** [Dev Biol.](#)[delete](#)[Contact Us](#) [Need Help?](#) [Citing Xenbase](#) [About Xenbase](#)

Xenbase Gene:Expression data for pax8 [Xenopus]

<http://xenbase.org/gene/expression.do?method=displayGenePageExpression&genId=483692&geneSymbol=pax8&geneName=pax8>

Hotels by Hilton – Hotel Reservati... Xenbase Gene:Expression data for...

Community Submitted Images

NF stage 23

XDB3 & Axel DB Images

NF stage 28

Literature Images

NF stage 35 and 36

PREV

A B

C D

E F

G

NEXT

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ore [+]

Fig. 3. Gene expression during pronephric development. A) Xpax-8 (pax8) stage 36. Within the pronephros only the pronephric tubules are stained. B) Xlim-1, stage 36. Strongest staining in within the nephrostomes and the pronephric duct, with weaker staining in the pronephric tubules. C, Xwnt-4, stage 35. Strong staining is in the pronephric tubules and weaker staining in the duct. D) Xpax-2, stage 36. E) xWT1, stage 36. F) Double stain with antibodies 3G8 (blue) and 4A6 (red). G) X-ret, stage 28. Staining is strongest in the posterior portion of the pronephric duct.

Image published in [Molecular regulation of pronephric development](#).

Species	Stage(s)	Tissue	Annotation Status
Xenopus	NF stage 35 and 36	pronephric kidney early proximal tubule	Automated curation, no stage curation, no tissue curation



stage 25 lateral view



stage 25 dorsal view

Phenote 1.4

Source XDB3
Image XL044I19_RA036_2g09_r39_2
Gene pax8
Stage NF stage 25
Anatomy XAO pronephric mesenchyme Comp
Expression Qualifier uniform
Present or Absent present
Evidence inferred from in situ hybridization
Comments

Term Info

pronephric mesenchyme



ONTOLOGY xenopus_anatomy

ID XAO:0000264

Related pronephric anlage
Synonyms: pronephric bulge
pronephric primordium

Definition Portion of tissue that consists of the mesenchymal precursor to the pronephric kidney and that begins to separate from the intermediate mesoderm at NF stage 21 and epithelializes by NF stage 30.

Superclass portion of tissue

starts during NF stage 21
(start_stage)ends during NF stage 29 and 30
(end_stage)develops from intermediate mesoderm
(develops_from)develops into nephrostome
(develops_into)
pronephric kidney
pronephric duct

Source	Image	Gene	Stage	Anatomy	Expression Qualifier	Present or Absent	Evidence
XDB3	XL044I19...	pax8	NF stage 25	ear vesicle	uniform	present	inferred from in situ hybridization
XDB3	XL044I19...	pax8	NF stage 25	pronephric mesenchyme	uniform	present	inferred from in situ hybridization

New

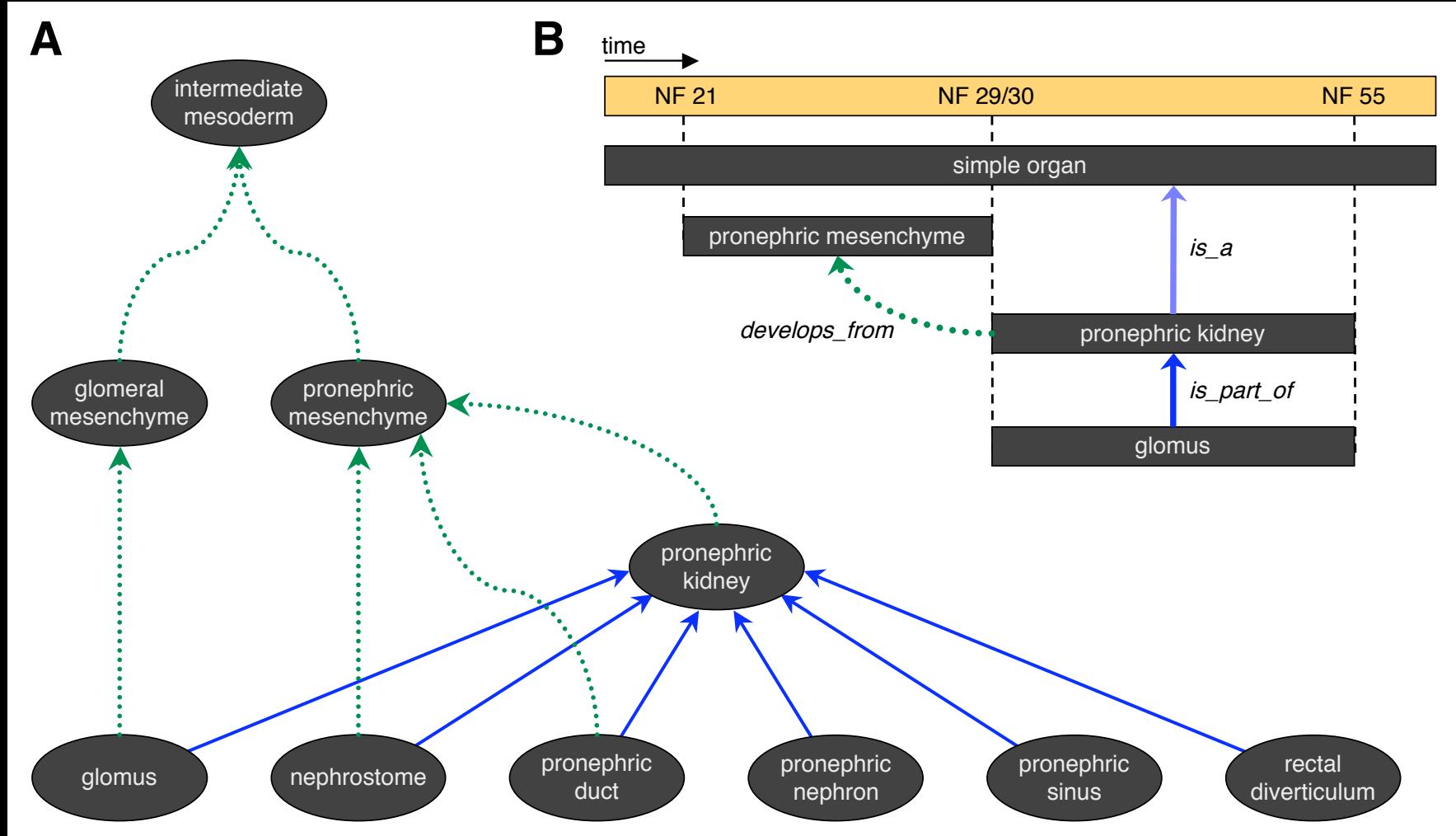
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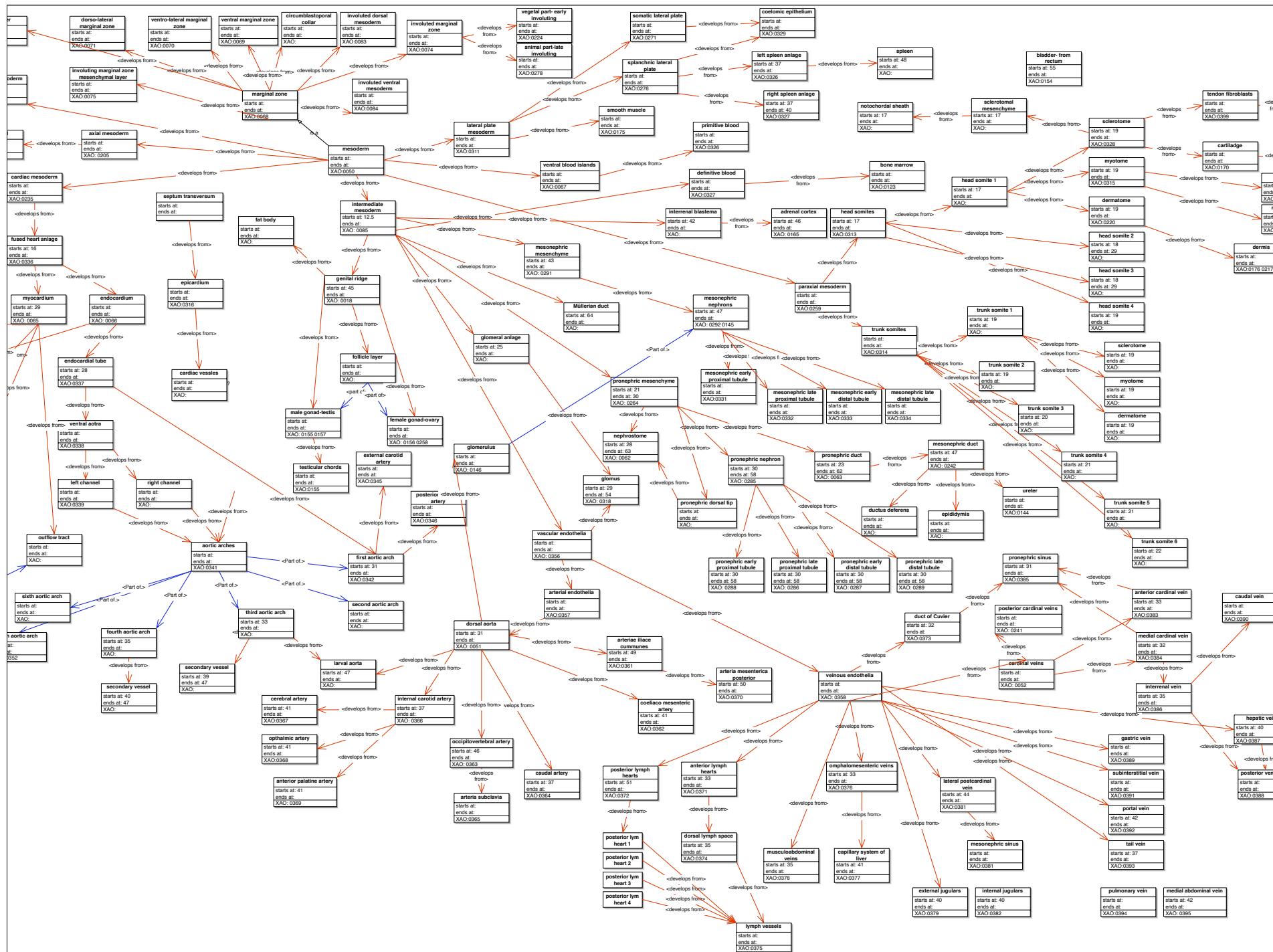
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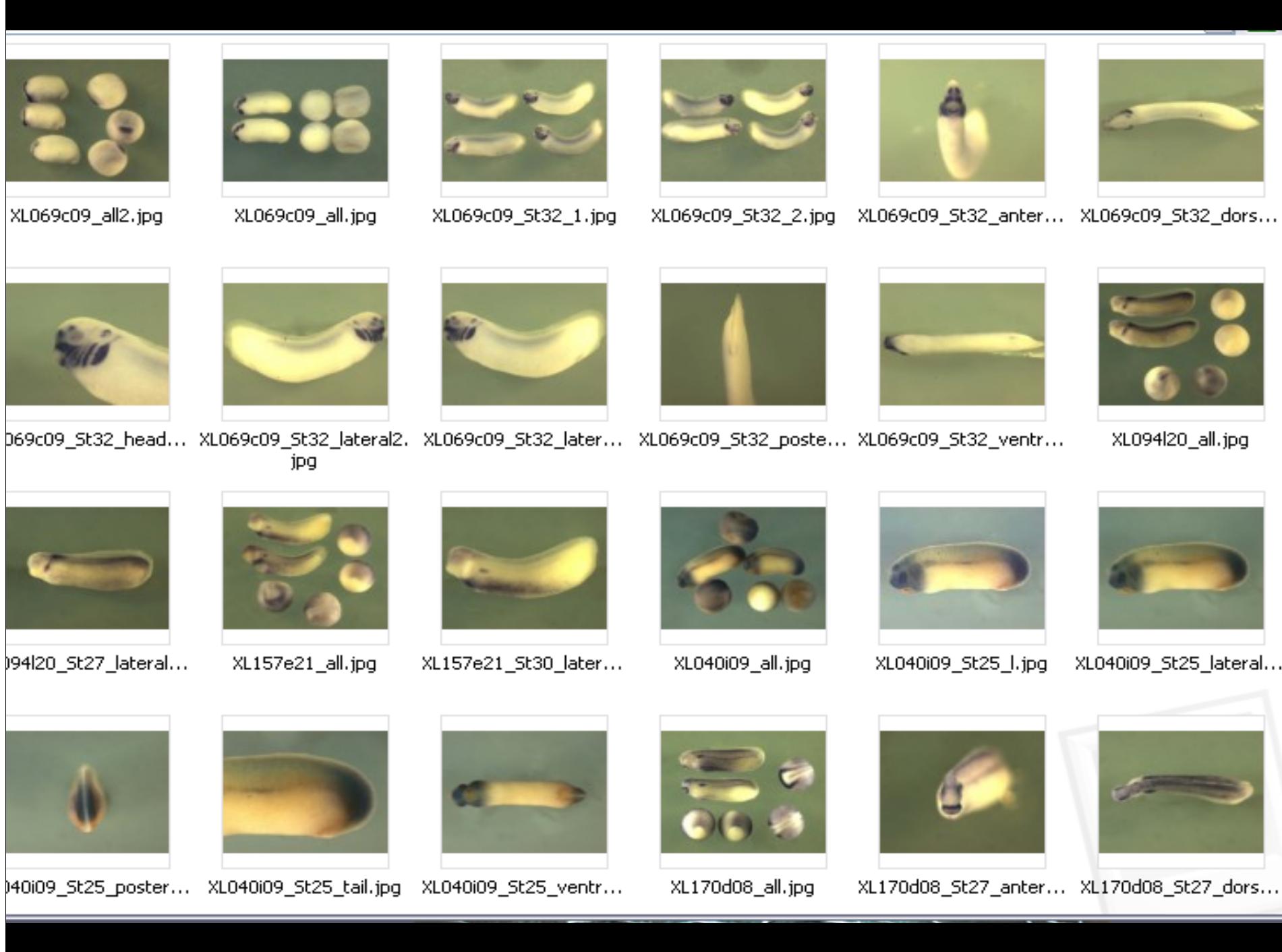
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Save Data

Filter:









XAO:1000345 = NF stage 33

XAO:0000567 = liver

XAO:0000213 = pronephric duct

XAO:0000332 = cranial ganglia

human anatomy ← human disease ← human gene exp

CARO

fish anatomy

PATO

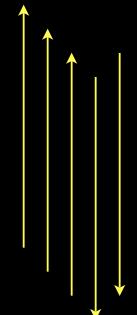
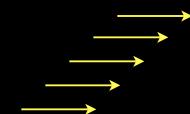
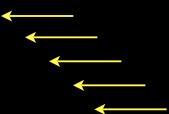
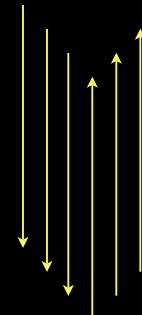
fish mutants

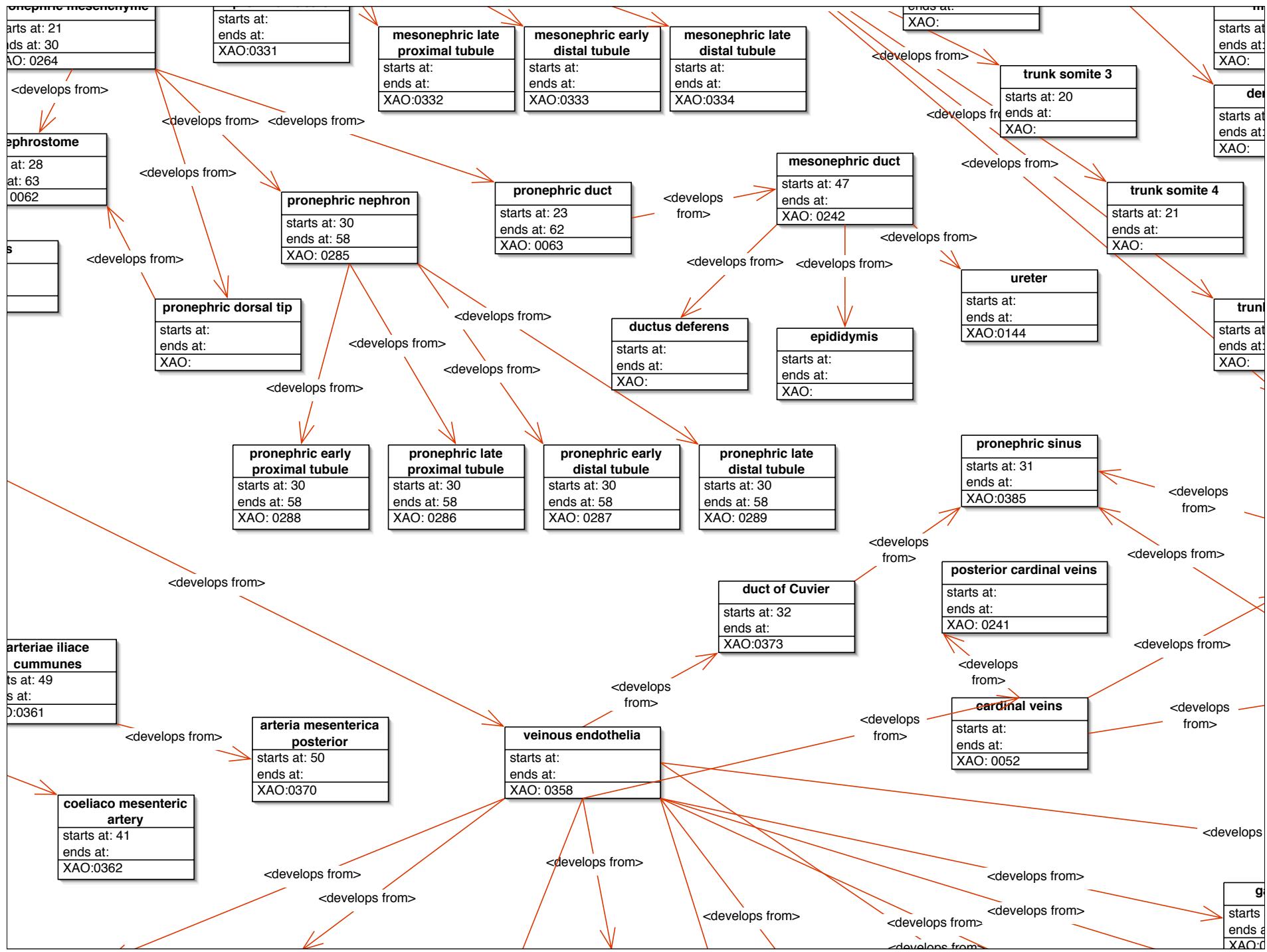
fish gene exp

frog anatomy

frog morphants

frog gene exp





- ontologies accurately represent gene expression patterns in space and time
- ontologies can use other ontologies for predicting gene function

acknowledgments...

3D; victor gerth; nato ueno (nibb)

EST analysis; kevin synder; jeff bowes

XAO; erik segerdell; monte westerfield (U of Oregon)

funding; national institutes of health, NICHD