



WormGUIDES an Interactive Informatic Developmental
Atlas at Subcellular Resolution

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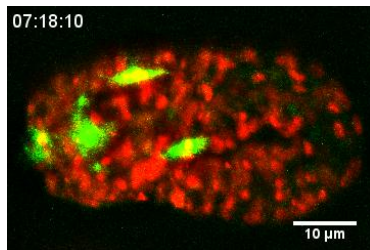
² Dept. of Cell Biology, Yale University, New Haven, CT

³ NIBIB, NIH, Bethesda, MD.

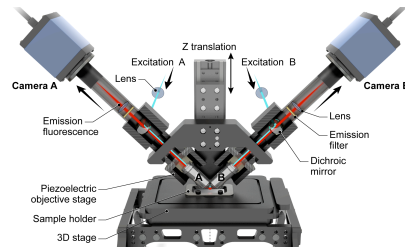
⁴ Dept. of Genetics & Dev Biology & Ctr. For Cell Analysis and Modeling, UConn Health Ctr., Farmington, CT

WormGUIDES Team

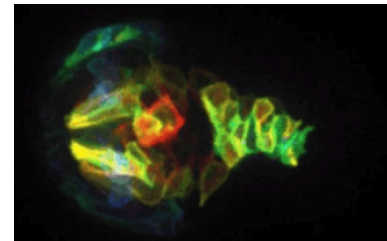
**Neuro
Development**



Microscopy



**Dev. Bio
Computational
Bio.
Computer Vision**



**Dev. Bio
Computational Bio.**



**Daniel
Colón Ramos**
Yale Medical School

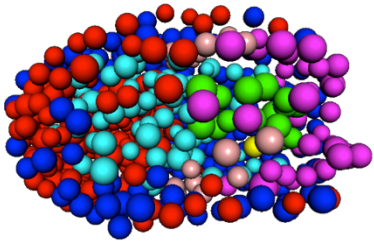
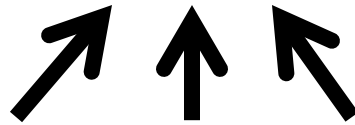
**Hari
Shroff**
NIBIB

**Zhirong
Bao**
Sloan Kettering Inst.

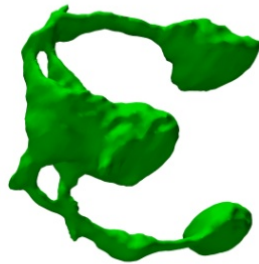
**Bill
Mohler**
UConn Health Ctr.



Atlas



Nuclear positions



neural morphology

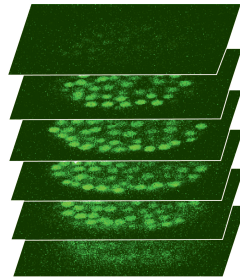
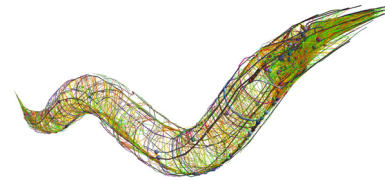


image data



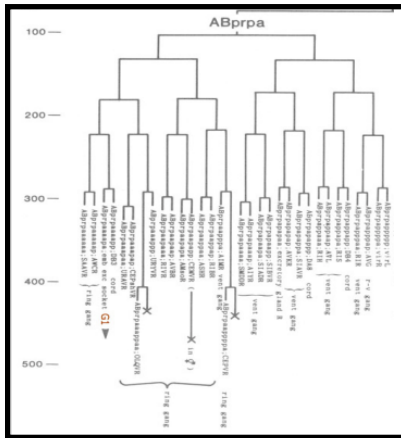
wiring diagram

AIA
RMED
CANL
CANR
...

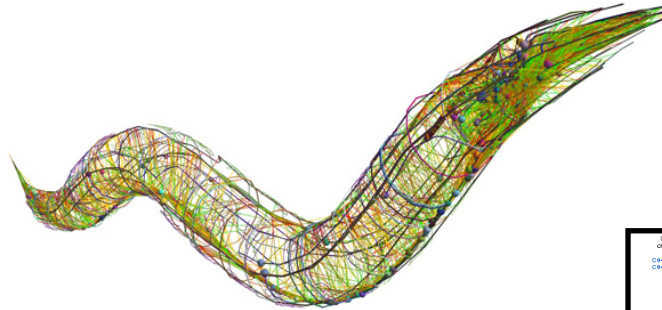
annotation

Motivation

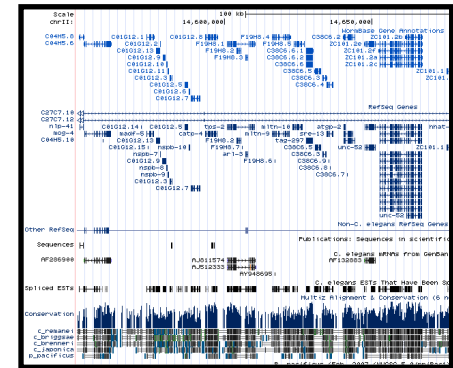
Cornerstones of
C. elegans Research:



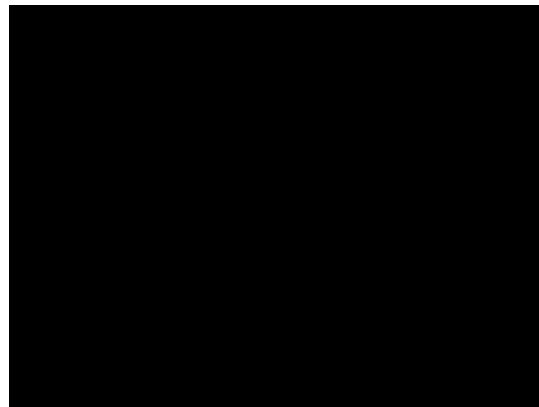
LINEAGE



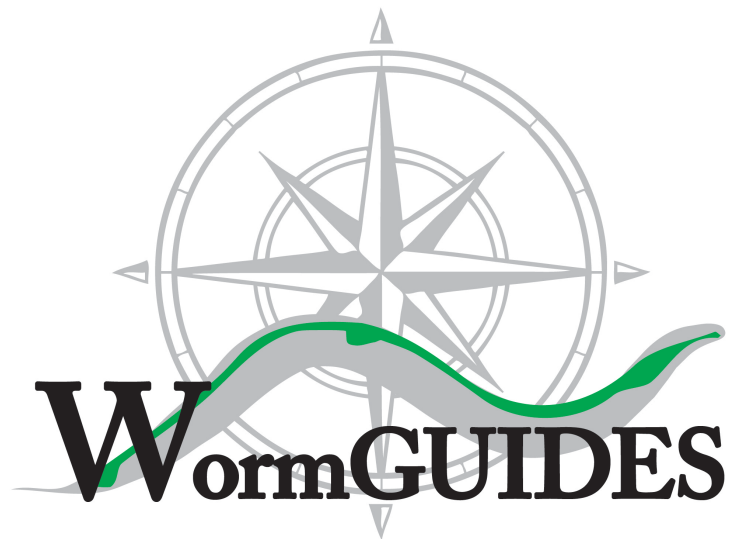
ANATOMY/
WIRING



GENOME



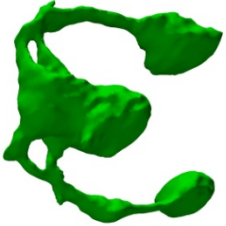
OBSERVABLE DEVELOPMENT



WormGUIDES



Starrynite
Cell tracking



Neural segmentation

Computer Vision

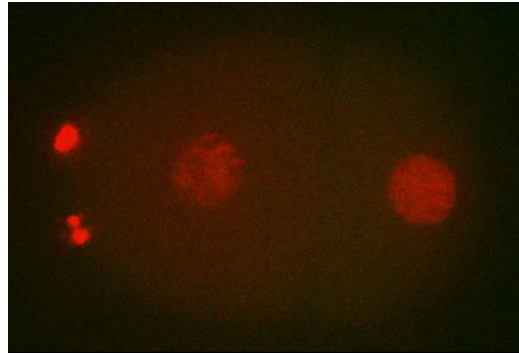


CytoSHOW

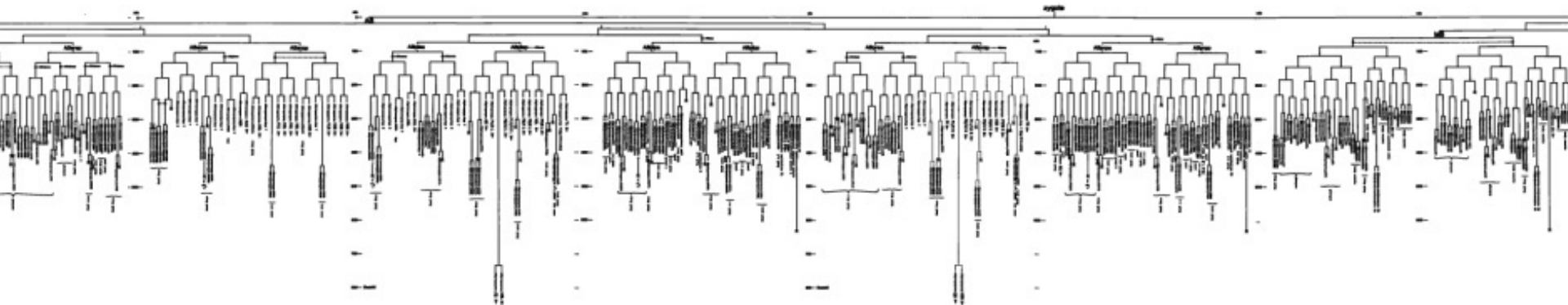
AceTree

Visualization

Cell Tracking



Bao Lab

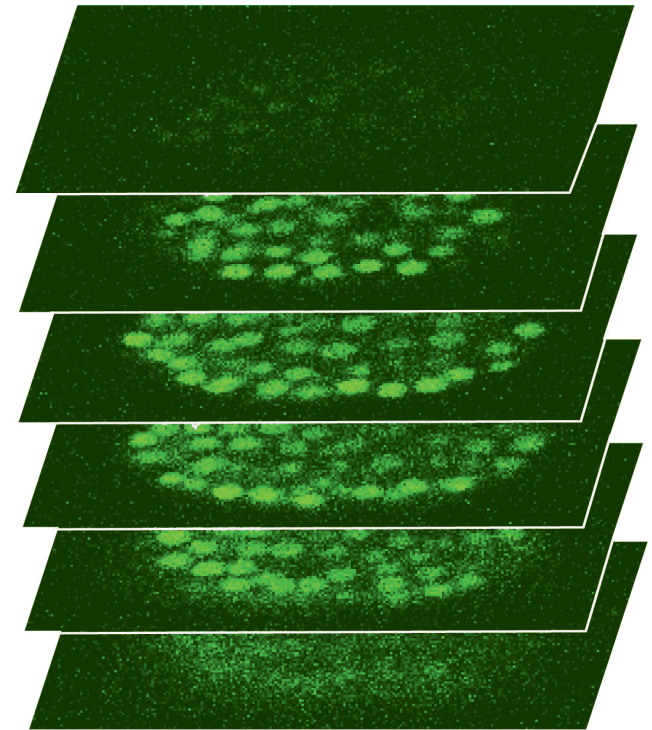
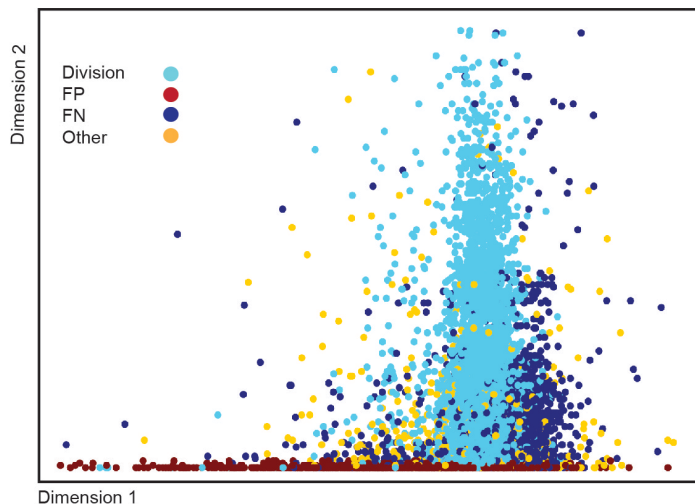


Starrynite: Cell Tracking

Algorithmically

- Supervised learning approach to segmentation and tracking.
- Used in a dozen+ labs worldwide
- ~35k lines Matlab

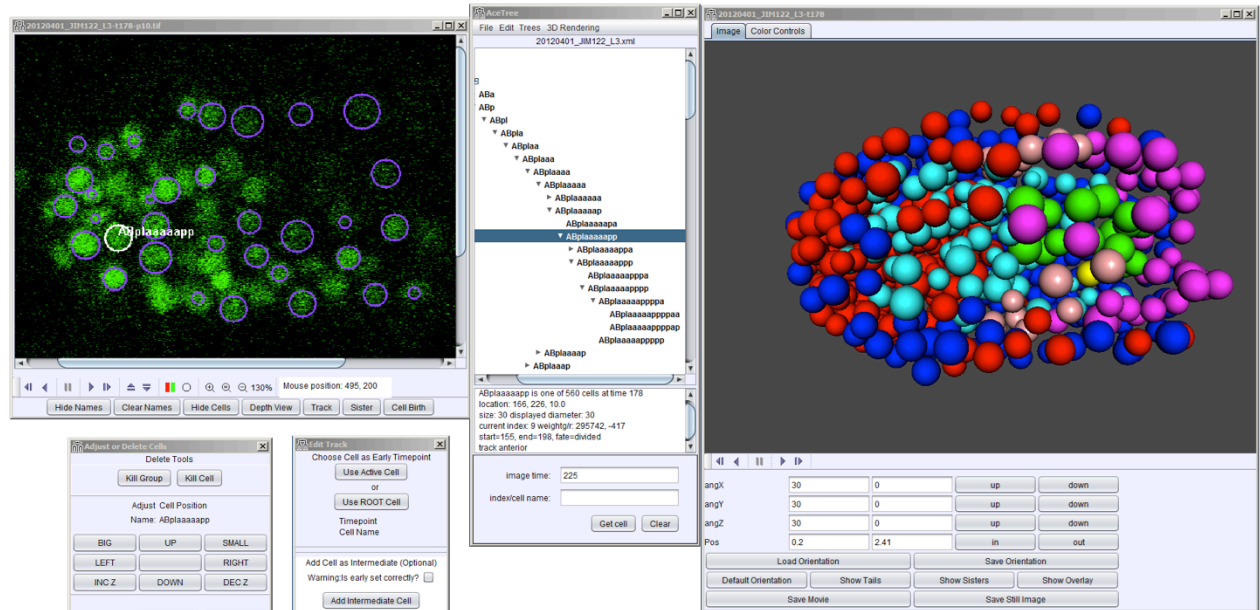
<http://starrynite.sourceforge.net/>



A semi-local neighborhood-based framework for probabilistic cell lineage tracing
Anthony Santella, Zhuo Du and Zhirong Bao
BMC Bioinformatics 2014, 15:217

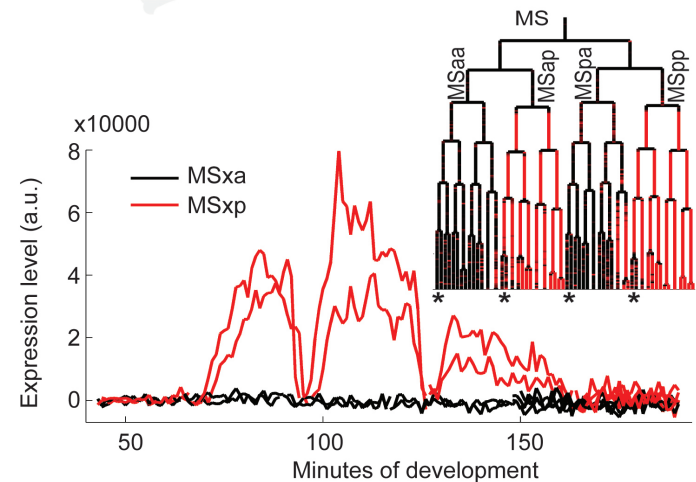
AceTree: Single Cell Analysis

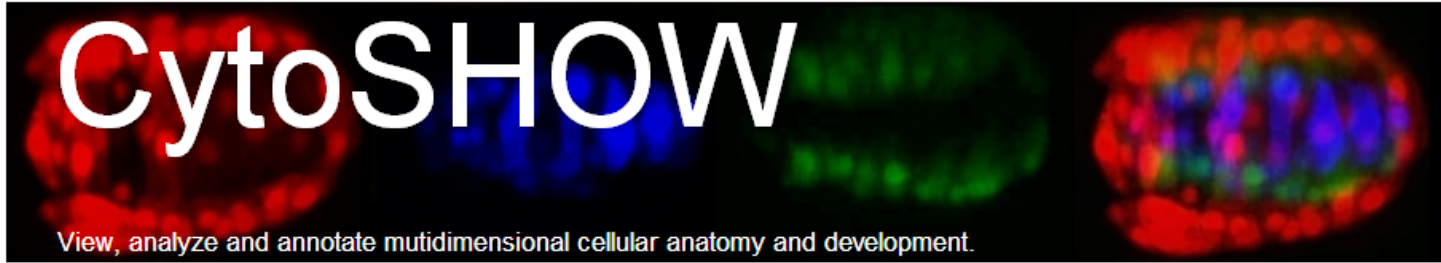
- Edit
- View
 - Tracks
 - image data
 - Lineage view
 - Gene expression
- Data analysis
- Stand alone app
 - ~60k lines Java
 - Built on ImageJ, Java3D



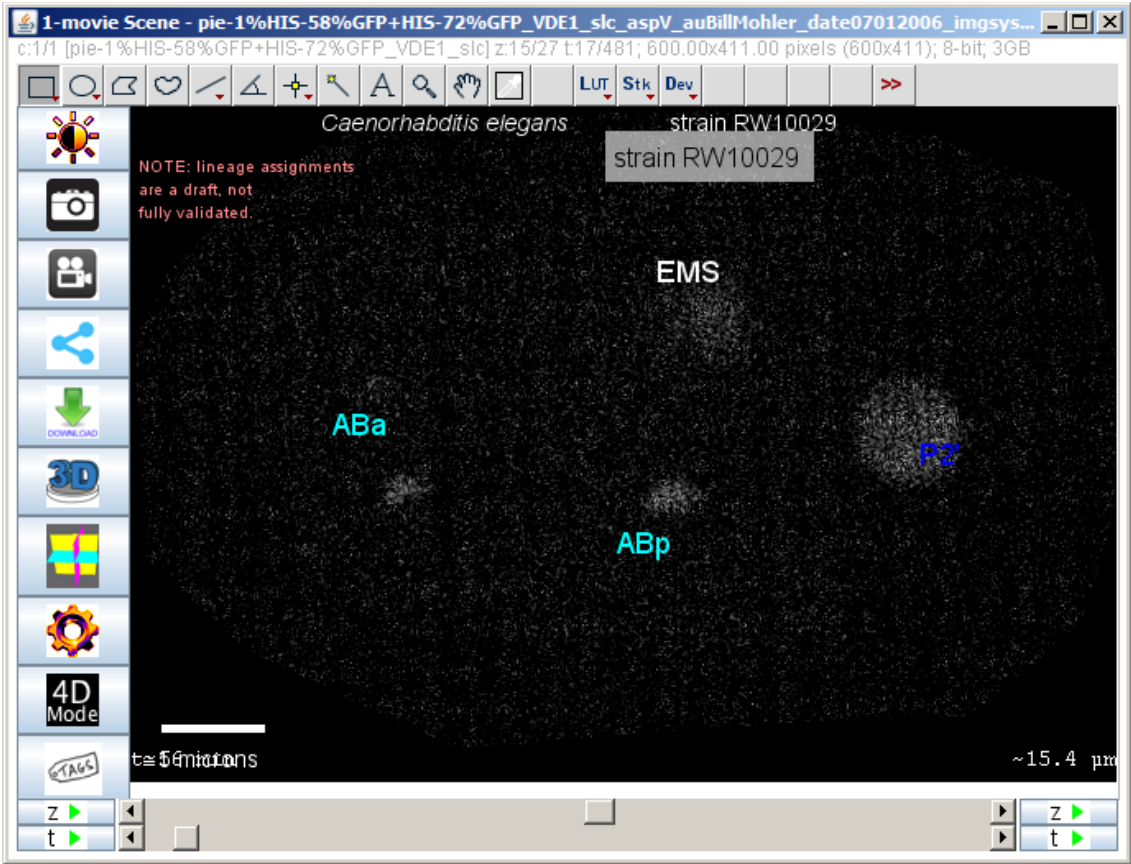
<http://starrynite.sourceforge.net/>

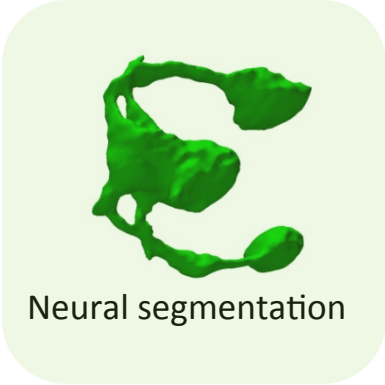
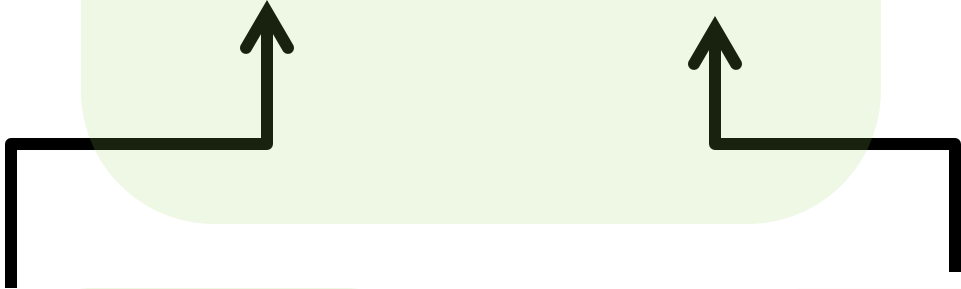
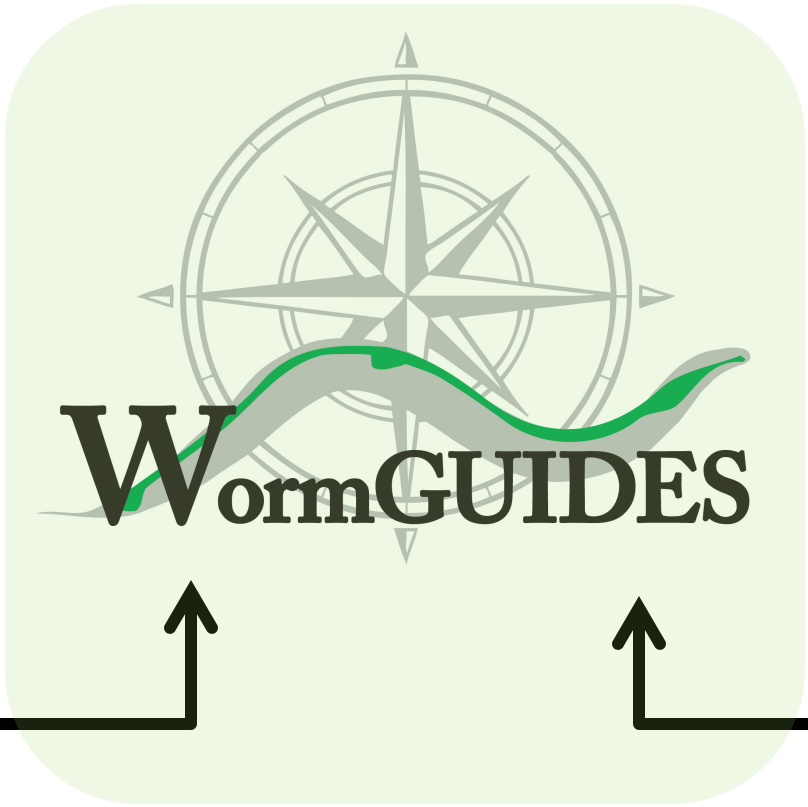
AceTree: a tool for visual analysis of *Caenorhabditis elegans* embryogenesis
 Thomas J Boyle, Zhirong Bao, John I Murray, Carlos L Araya and Robert H Waterston*
 BMC Bioinformatics 2006, 7:275





- Web launch Java app, built on top of ImageJ www.cytoshow.org





Computer Vision



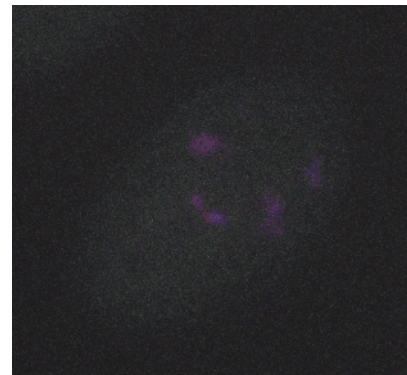
AceTree

Visualization

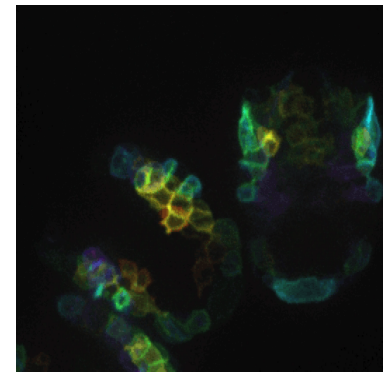
Long-Term Project Goal

Alignment and integration of multiple data sets in unified multi-layer virtual environment.

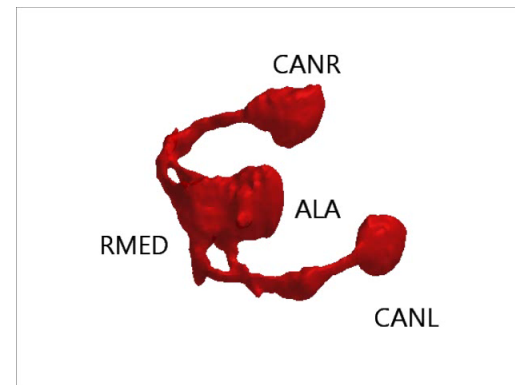
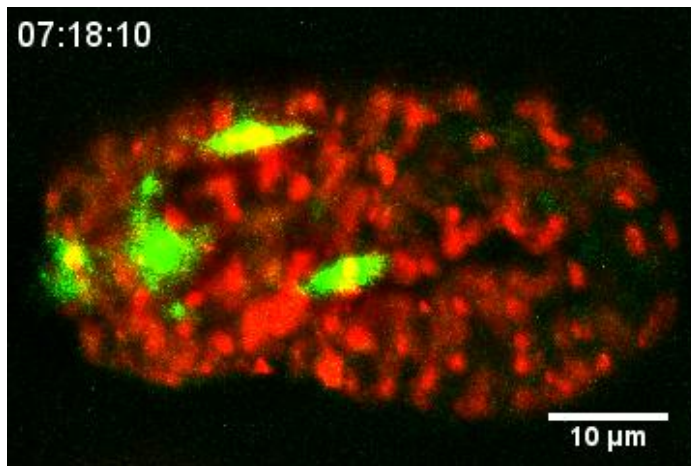
- Neurons from different strains
- Aligned using nuclei



lim-4



irx-1



WormGUIDES Mobile app

C. elegans anatomy
nuclear positions

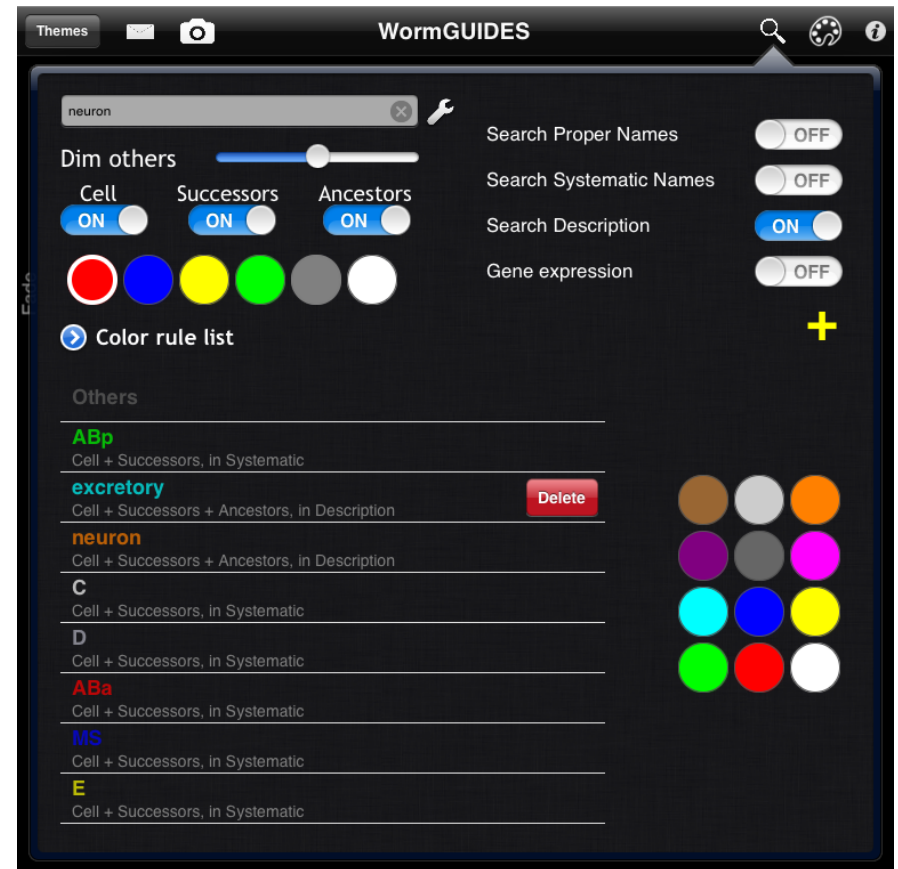
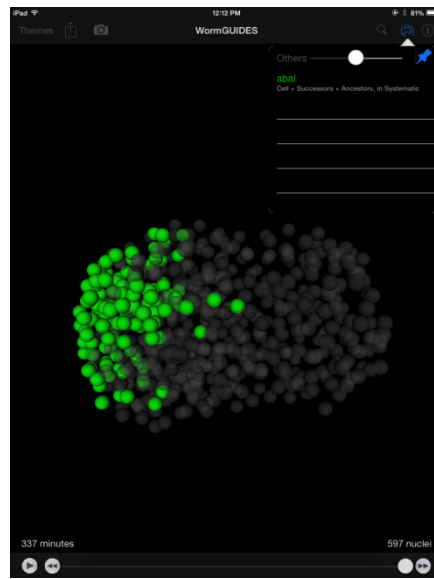
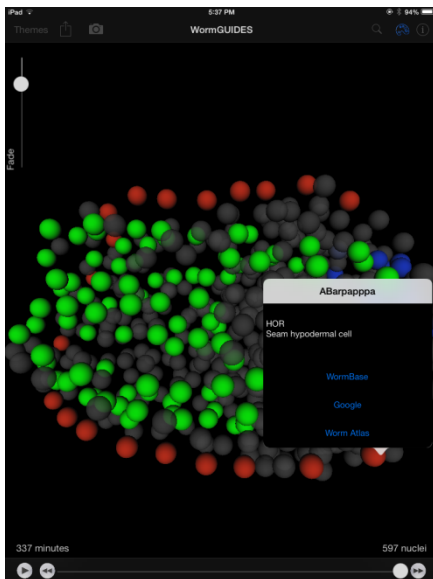
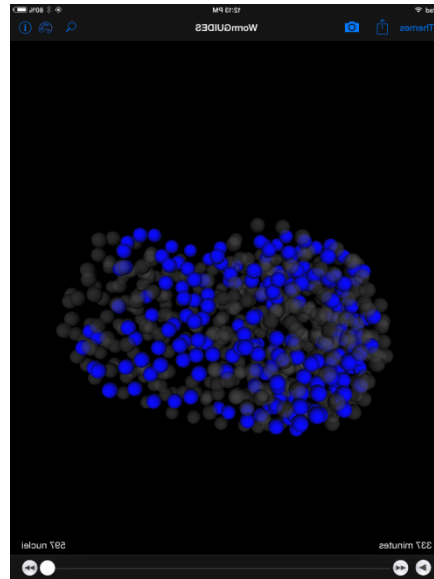
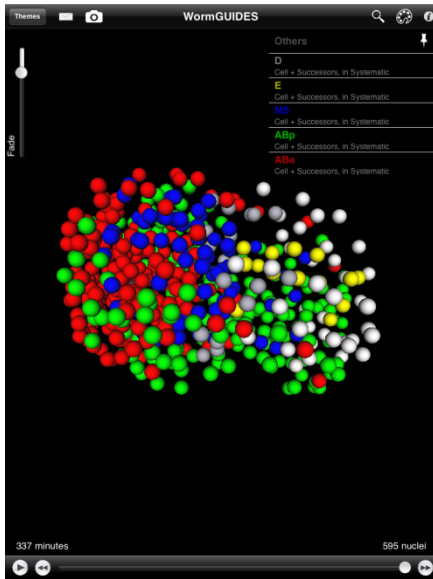
Available for

- Android
- iOS

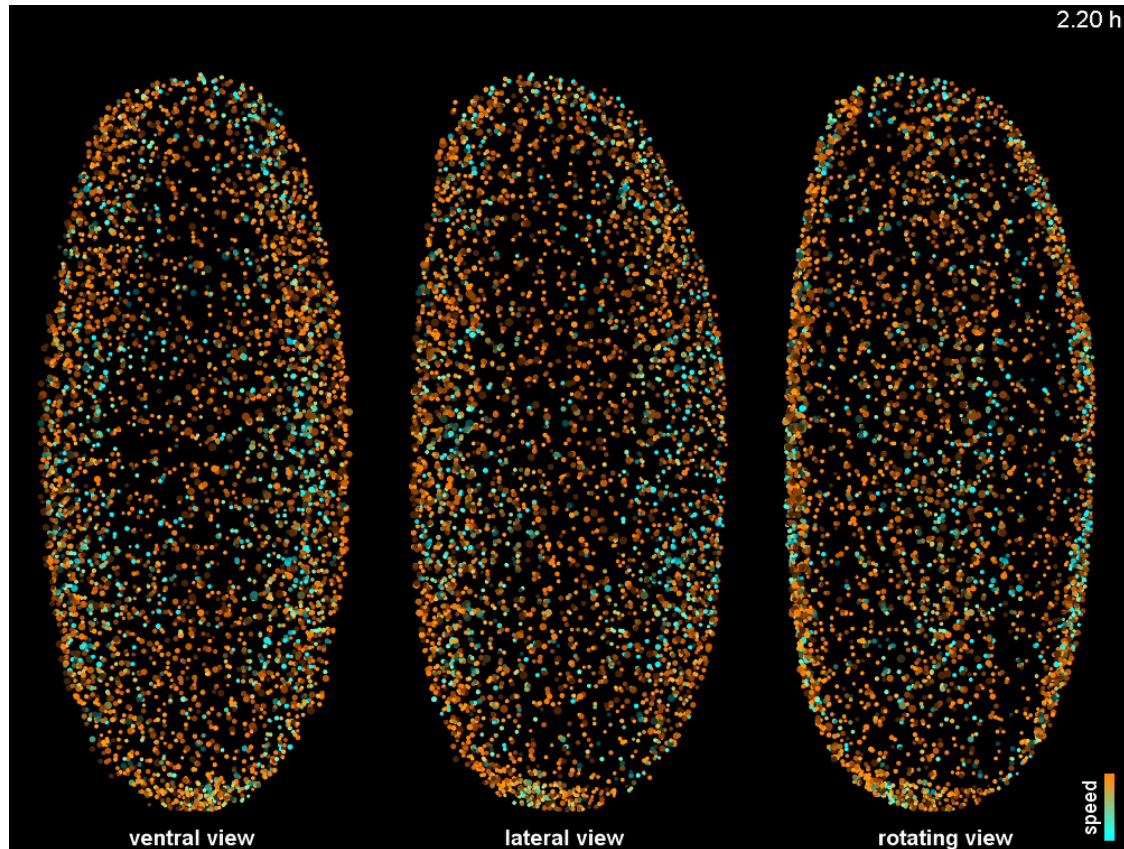
www.wormguides.org



WormGUIDES app



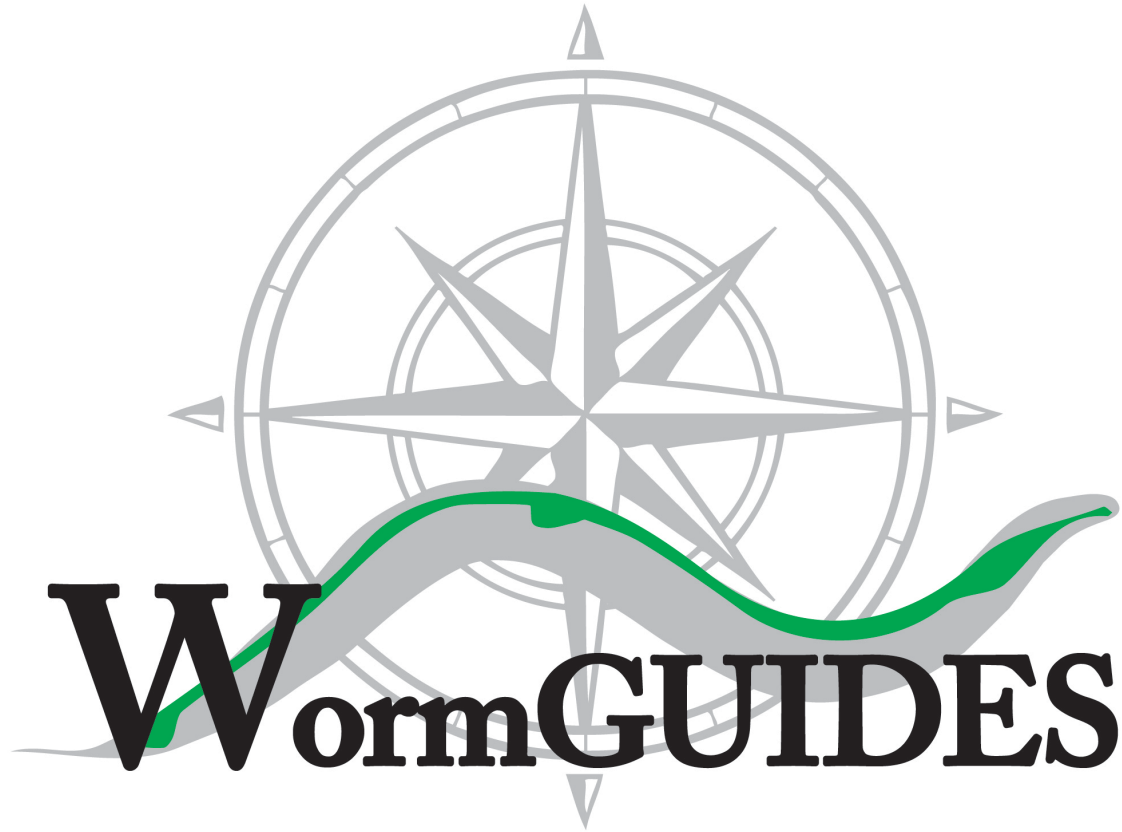
Futures



Starrynite reconstruction of drosophila development

P. J. Keller, A. D. Schmidt, A. Santella, K. Khairy, Z. Bao, J. Wittbrodt and E. H. K. Stelzer, "Fast, high-contrast imaging of animal development with scanned light sheet-based structured-illumination microscopy" *Nature Methods* 2010, 7, pp. 637-642

Input, collaboration, contributions welcome!



www.wormguides.org

<http://starrynite.sourceforge.net/>

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