

Monte Carlo Simulation of Somatic Twist in Ancient Marine Worms

How the First Digital Organism in a Computer Evolves *In Silico* to Become the *First Fish*



All graphics made by the student researcher unless otherwise cited. Al was not used when creating this display item.

INTRODUCTION

Background

Kinsbourne (2013) had proposed **somatic twist theory** for the evolution of **decussation** in vertebrates, as a by-product of **dorsoventral inversion** during the invertebrate-to-vertebrate transition 550 million years ago. Comparative study of select **morphological models** (Cheong, 2025) inside an aquarium suggests a plausible evolutionary pathway for how this might have occurred in ancient marine worms, leading to corticospinal tracts decussation in *Pikaia*, the first fish.

Research Question

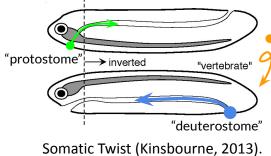
The smallest worm will turn — but how? What force(s) could possibly drive the biomechanical tissues of ancient marine worms to perform a somatic twist?

Purpose

Run Monte Carlo simulation to recreate evolutionary events inside a computer; and *see* how the first digital *C. elegans* worm evolves *in silico* to become the 'first fish' — with a dorsoventrally inverted body plan after a somatic twist.

Hypothesis

Underwater **buoyancy force** drove somatic twist of ancient marine worms.

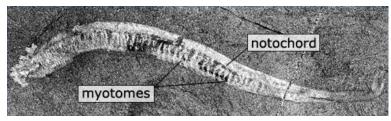




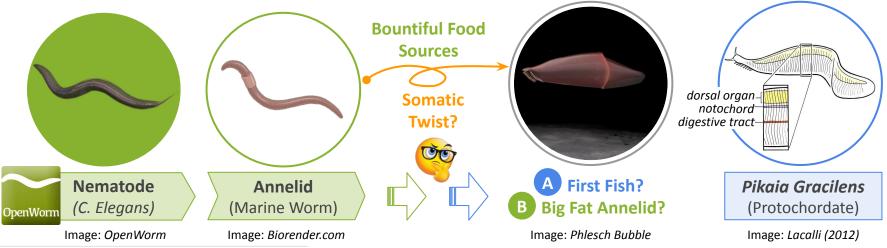
Comparative study of morphological models from ancient marine worms to the first fish inside an aquarium (Cheong, 2025).

1. Experimental Design

Starting with *C. elegans*, adjust model parameters for biomechanical matter and environment to evolve successive new generations of *marine worms* to see if any such instances, when given bountiful food sources: (a) turn into the **First Fish** after a somatic twist; or (b) grow into **Big Fat Annelids** without any somatic twist.

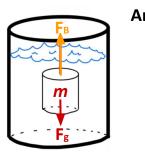


A fossil *Pikaia* (protochordate) has a visible notochord. Photo: *Chip Clark, Museum of Natural History, Smithsonian Institution.*



2. Ask: "What Happens When Food Sources are Bountiful?"

- □ Intestinal gut expands with food and becomes heavier also becomes denser.
- Excess fat plausibly stored alongside ventral nerve cord mirroring the *Pikaia* dorsal organ.
- Or would *C. elegans* evolve to become more like an annelid along the way?
- Because hydrostatic coelomic fluid allows free movement and expansion of internal organs.
- Over geologic time, aquatic environment attains salinity higher viscosity & density, too.
- Underwater **buoyancy force**, at or beyond turning point, drives somatic twist of marine worm.

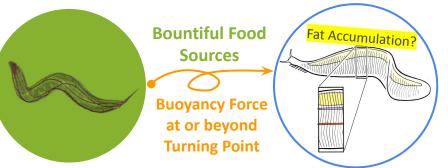


Archimedes Principle: $F_{B} = \rho_{w}gV$ $F_{g} = mg$ $W = F_{g} - F_{B}$

Underwater buoyancy force, at or beyond turning point, drives somatic twist of marine worm. Image: *James Charbonneau*.

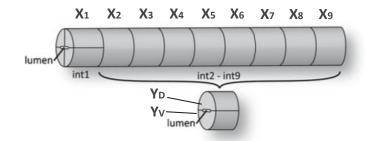
Why *C. Elegans*? Start where data is most complete. Image: *OpenWorm*.

Pikaia as target reference and end point of simulation. Image: *Lacalli (2012)*.



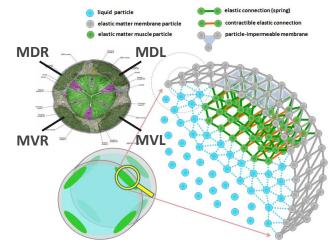
3. Gather Parameters for "Biomechanical Matter + Environment" Simulation

- C. elegans digestive tract: pharynx (57 + 38 cells), intestine (20 cells), and rectum (11 cells).
- Intestinal cells: simple tube runs along **80% body length** and **1/3 of somatic mass** of organism.
- Model built with biomechanical matter: contractile matter, elastic matter, and membranes.



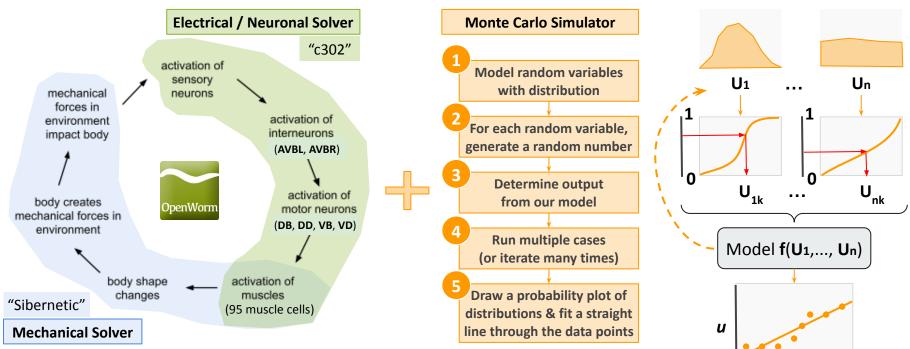
Intestine with 20 cells modeled as 9 rings

Basic structure of the intestine as a set of rings: four cells in the anterior-most X_1 ring and two cells (Y_D and Y_V) in each of the ring from X_2 through X_9 . Dorsoventral inversion is detected from relative positions of Y_D and Y_V . Image: *Dimov & Maduro (2019)*.



Types of particles used in the Sibernetic simulation framework for creating a model of the worm body with cuticle, **muscle quadrants** (MDR, MVR, MVL, and MDL) and internal liquid. Image: *Gleeson et al. (2023)*.

4. Extend Sibernetic & OpenWorm Software with Monte Carlo Simulator

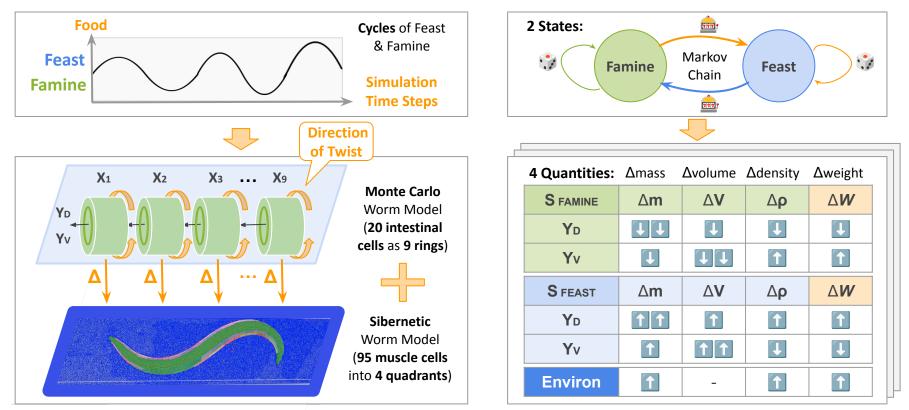


Feedback control loop of simulation engine. Solvers for two systems of equations: smoothed particle hydrodynamics and Hodgkin-Huxley, overlap at the activation of muscle cells. Image: *OpenWorm*.

Adding Monte Carlo method so as to predict probability of various outcomes (e.g., somatic twist?) when random variables (e.g., feast or famine?) are present.

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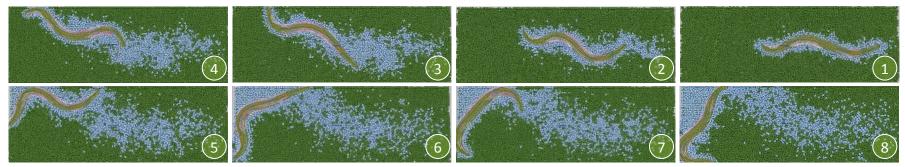
5. Run Monte Carlo Simulation — Until Somatic Twist Event Detected



RESULTS

1. Calibration: "The Smallest Worm Will Turn." (Shakespeare, "Henry VI", 1591)

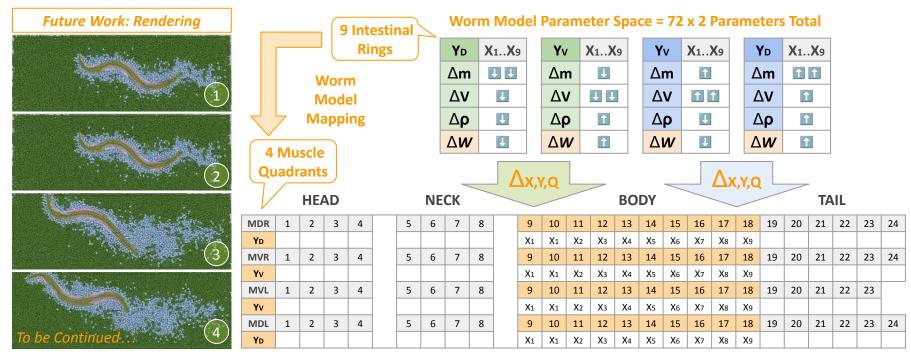
Trial	Platform	Processor	GPU	Memory	Duration	Total Steps	Run Time	OK?
1A	MacBook Air	8-Core Apple M2	10-Core GPU	16 GB	15 ms	3000	2466 sec	X
1B	MacBook Pro	8-Core Intel i9 2.4 GHz	Radeon Pro 8 GB	32 GB	15 ms	3000	199 sec	\checkmark
2B	MacBook Pro	8-Core Intel i9 2.4 GHz	Radeon Pro 8 GB	32 GB	5000 ms	1000000	34.65 hrs	
2C	MacBook Pro	16-Core M4 Max	40-Core GPU	64 GB	5000 ms	1000000	TBD	
2D	AWS Cloud	TBD	-	TBD	5000 ms	1000000	TBD	



A simulated *C. elegans* makes three turns in five seconds inside a bounded aquatic environment. Different processor and memory configurations were benchmarked to estimate computing resources required for running repeated trials on OpenWorm + Sibernetic. Original Docker image and Python source code: *https://hub.docker.com/r/openworm/openworm*

RESULTS

2. Visualization: "The Worm Did Turn. It Turned on Itself." (Kinsbourne, 2013)



Worm behavior emerges from simulation of empirical data, with tiny proportional changes: $\Delta x_{,Y,Q}$ applied to the worm body: **MDR** 9..18, **MVR** 9..18, **MVL** 9..18, and **MDL** 9..18 at each time step, driven by famine and feast cycles which successive worm generations must adapt to.

DISCUSSION

Validation In Silico: Monte Carlo Simulation of Somatic Twist

Theories Matching Simulation Results

- □ Simulation results provide *in silico* validation for **somatic twist theory** of Kinsbourne (2013).
- □ That means **dorsoventral inversion** (St. Hilaire, 1882), a long-held hypothesis in the field, has also been validated.
- □ **"From Worm to Man"**: underwater **buoyancy** is the force that drives somatic twist, which in turn cause dorsoventral inversion that ultimately led to **decussation** in vertebrates.

Review of Known Facts About C. Elegans



- □ Four bands of muscles run the length of the body enable locomotion.
- □ Head moves freely all four muscle quadrants independently wired.
- Dorsal or ventral bending for body movements never left or right.
- Locomotion: lie on *left or right side* when crossing horizontal surface.

Possible Errors

- □ All models are *wrong*, but some are *useful*.
- □ More than one way to map 9 intestinal rings onto 95 muscle cells for Sibernetic simulation.
- Geometric progression from tiny proportional changes to model parameters at each time step.

Unexpected Challenges

OpenWorm simulation very compute intensive.
 2 ½ days on MacBook simulates 5-sec duration.
 Run future repeated simulations on AWS cloud?
 Highlighting worm body for visual confirmation
 Somatic twist is bottom-up emergent behavior
 Design considerations of somatic twist detector

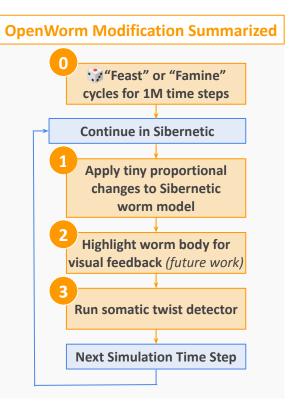
Repeated Trials

□ Repeated Monte Carlo trials expensive to run!

CONCLUSIONS

Underwater Buoyancy Drives Somatic Twist of Marine Worm

- Somatic twist theory of Kinsbourne (2013) offers a testable hypothesis for how corticospinal tracts became decussated as by-product of dorsoventral inversion in ancient marine worms.
- My results validated somatic twist theory by simulating underwater buoyancy force that drives somatic twist in marine worms during times when food was bountiful.
- My contribution: novel design of a somatic twist detector that can track relative cell positions during Sibernetic simulation to identify when dorsoventral inversion has just occurred.
- A biomechanical worm model has greater reproducibility *in silico* compared with mechanical 'twistable body plan structures' submerged inside an aquarium (Cheong, 2025).
- Monte Carlo simulation shows great promise as a practical new approach to conducting evolutionary biology experiments *in silico* when combined with Sibernetic and OpenWorm.



REFERENCES

- 1. Allentoft-Larsen, M. C., Gonzalez, B. C., Daniels, J., Katija, K., Osborn, K., & Worsaae, K. (2021). Muscular adaptations in swimming scale worms (Polynoidae, Annelida). *Royal Society Open Science*, *8*(10). https://doi.org/10.1098/rsos.210541
- 2. Cheong, A. (2025). Why Do Vertebrates Have Decussated Corticospinal Tracts?
- 3. Dimov, I., & Maduro, M. F. (2019). The C. elegans intestine: organogenesis, digestion, and physiology. *Cell and Tissue Research*, 377(3), 383–396. https://doi.org/10.1007/s00441-019-03036-4
- 4. Gleeson, P., Alicea, B., Cantarelli, M., Idili, G., Lee,C.W., Palyanov, A., Khayrulin, S., Portegys, T., Ghayoomie, V. & Larson, S. (2023, June 24–28). Updates from the OpenWorm project: incorporating NeuroPAL data and ASH neuron electrophysiological recordings [Poster session]. 24th International C. elegans Conference, Glasgow, Scotland.
- 5. Gleeson, P., Cantarelli, M., Currie, M., Hokanson, J., Idili, G., Khayrulin, S., Palyanov, A., Szigeti, B., & Larson, S. (2015). The OpenWorm Project: currently available resources and future plans. *BMC Neuroscience*, *16*(S1). https://doi.org/10.1186/1471-2202-16-s1-p141
- 6. Kinsbourne, M. (2013). Somatic twist: A model for the evolution of decussation. *Neuropsychology*, 27(5), 511–515. https://doi.org/10.1037/a0033662
- 7. Lacalli, T. (2012). The Middle Cambrian fossil Pikaia and the evolution of chordate swimming. *EvoDevo*, *3*(1), 12. https://doi.org/10.1186/2041-9139-3-12
- 8. OpenWorm. (n.d.). Openworm Website. https://openworm.org & https://github.com/openworm/OpenWorm
- 9. Palyanov, A., Khayrulin, S., & Larson, S. D. (2016). Application of smoothed particle hydrodynamics to modeling mechanisms of biological tissue. *Advances in Engineering Software, 98*, 1–11. https://doi.org/10.1016/j.advengsoft.2016.03.002
- 10. Pikaia gracilens. (n.d.). The Burgess Shale. https://burgess-shale.rom.on.ca/fossils/pikaia-gracilens/
- Szigeti, B., Gleeson, P., Vella, M., Khayrulin, S., Palyanov, A., Hokanson, J., Currie, M., Cantarelli, M., Idili, G., & Larson, S. (2014).
 OpenWorm: an open-science approach to modeling Caenorhabditis elegans. *Frontiers in Computational Neuroscience*, 8. https://doi.org/10.3389/fncom.2014.00137