



## Tufts University: The Levin Lab, Department of Biology: Research

data grow exponentially, true insight into shape generation and repair is significantly impaired because bioinformatics is focused on gene sequences but not applicable to analyses of <u>shape</u>. Thus, several fields are stymied by a lack of conceptual and computerized tools to link mechanistic understanding of molecular signals with behavior of the patterning systems they encode. The field is missing (1) convenient symbolic mathematical tools with which to formalize shape and changes in shape, such that the outcomes of patterning experiments can be stored in a searchable database (like Entrez at NCBI, but for morphogenesis instead of gene expression), (2) generally-accessible agent-based virtual environments within which mechanistic models of patterning can be simulated *in silico* and integrated with existing data for testing and derivation of key regulatory properties, and (3) accessible artificial intelligence tools to help discover models consistent with experimental results in fields where the data are so abundant and complex that scientists cannot invent models consistent with empirical data.

We are using the data on genetic and bioelectrical mechanisms of regeneration in planarian flatworms (a very popular model system for molecular genetics work) as a proof-of-principle to 1) create a prototype for a *symbolic* mathematical formalism for encoding knowledge about shape, 2) implement a computing platform (expert system on planarian regeneration) so that anyone can query the existing literature for information about functional experiments that modify morphology, 3) produce a flexible and easy-to-use system for modeling the patterning consequences of control networks including both biochemical and physiological mechanisms, 4) create an Artificial Intelligence tool to assist users to discover mechanistic, constructivist models of signaling among components that match sets of functional data on patterning pathways, 5) use this system to identify a model explaining some of the remarkable regenerative abilities of planarian worms, which can regenerate any part of their body regardless of how they are cut, and 6) experimentally test new predictions of the models we identify in this way. Our work is yielding conceptual modeling and automated mining tools to revolutionize the building of algorithmic, understandable models directly from functional data that are too difficult to discover manually, thus impacting many fields of biology and engineering.

- Using techniques from artificial intelligence, computational neuroscience, and cognitive science to make models of morphogenesis - treating patterning systems as primitive cognitive <u>agents</u>
  - Modeling pattern formation and cell regulation as neural-like circuits with plasticity, memory, and goal satisfaction circuits; using modulation of global neurotransmitter and electrical synapse properties to write pattern memories and behavioral repertoires into living tissue
  - Constructing quantitative models of patterning using <u>extremal (least-action) principles</u>

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