

# Evolutionary learning

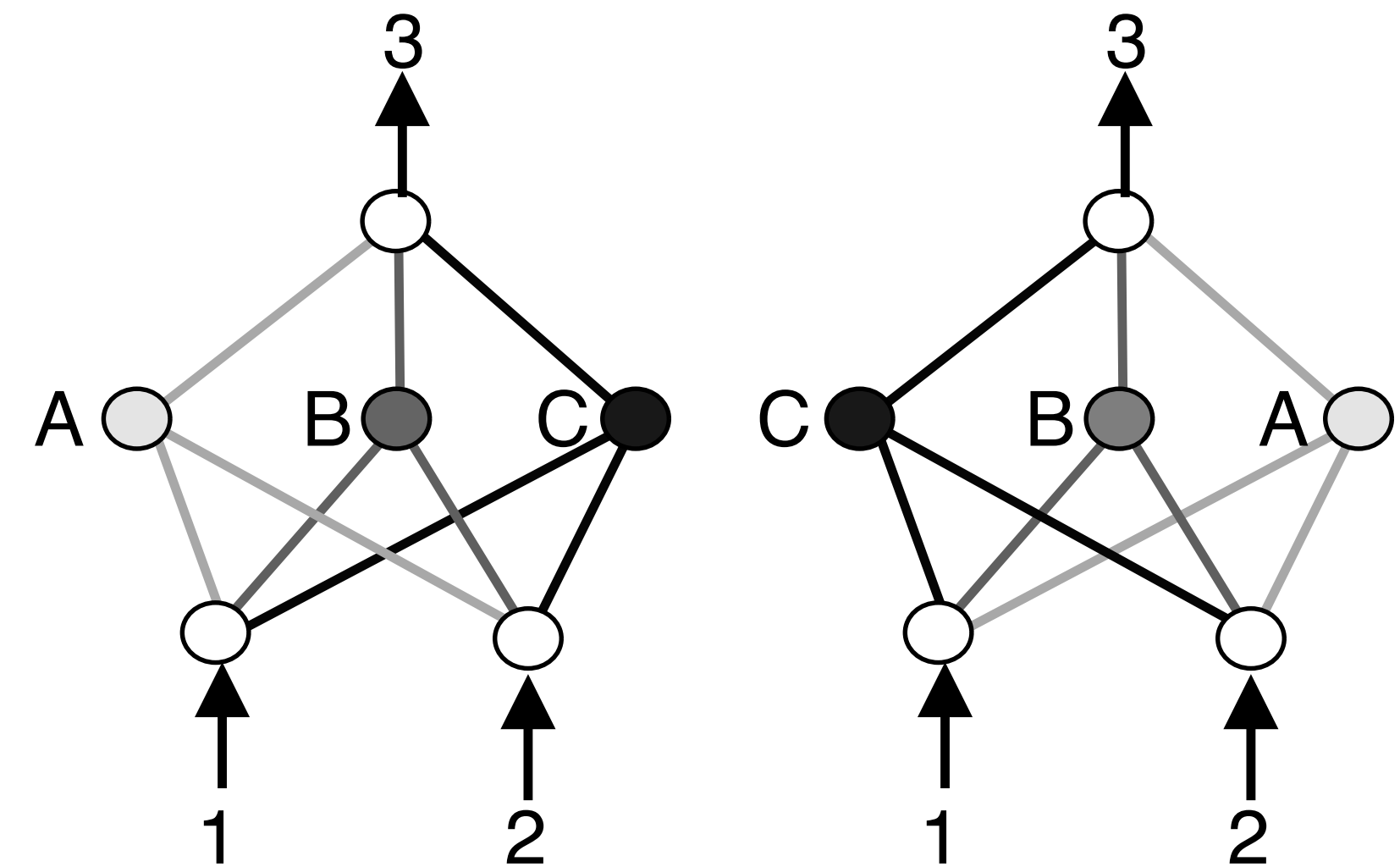
**NEAT & HyperNEAT**

**JB Mouret**

# How to evolve both the structure and the weights?

## Concept of direct encodings:

- Encode a graph as the genotype
- Labels = weights
- Mutation:
  - add a connection between two random nodes
  - remove an existing connection
  - add a neuron (usually on an existing connection)
  - change a weight (e.g., Gaussian perturbation)
- No cross-over:
  - permutation problem
  - graph matching is hard (complexity)
- Use a standard evolutionary algorithm (not an ES!)



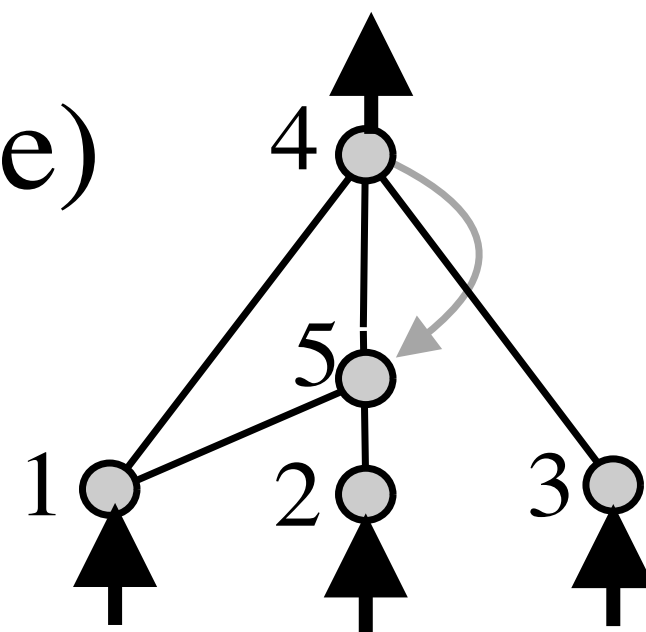
# NEAT

## Concept of NEAT:

- Grow incrementally: start with always the same topology
  - Facilitate the comparison of networks with innovation numbers
- A (global) counter for each new connection

Genome (Genotype)							
Node	Node 1	Node 2	Node 3	Node 4	Node 5		
Genes	Sensor	Sensor	Sensor	Output	Hidden		
Connect.	In 1	In 2	In 3	In 2	In 5	In 1	In 4
Genes	Out 4	Out 4	Out 4	Out 5	Out 4	Out 5	Out 5
	Weight 0.7	Weight -0.5	Weight 0.5	Weight 0.2	Weight 0.4	Weight 0.6	Weight 0.6
	Enabled	<b>DISABLED</b>	Enabled	Enabled	Enabled	Enabled	Enabled
	Innov 1	Innov 2	Innov 3	Innov 4	Innov 5	Innov 6	Innov 11

Network (Phenotype)



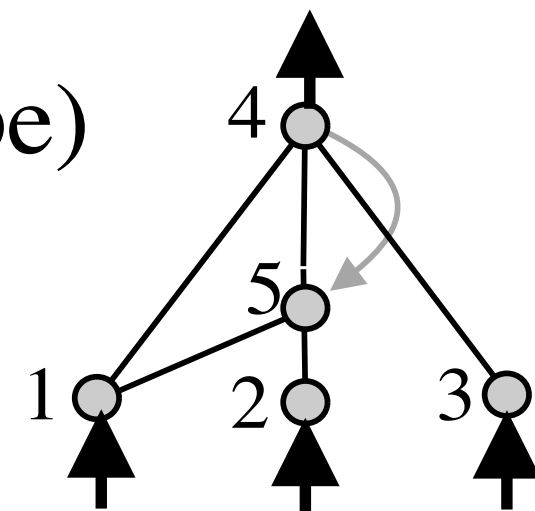
# NEAT

## Neuro-Evolution of Augmenting Topologies (NEAT)

- Facilitate the comparison of networks with innovation numbers
  - A (global) counter for each new connection

Genome (Genotype)							
Node	Node 1	Node 2	Node 3	Node 4	Node 5		
Genes	Sensor	Sensor	Sensor	Output	Hidden		
Connect.	In 1	In 2	In 3	In 2	In 5	In 1	In 4
Genes	Out 4	Out 4	Out 4	Out 5	Out 4	Out 5	Out 5
	Weight 0.7	Weight -0.5	Weight 0.5	Weight 0.2	Weight 0.4	Weight 0.6	Weight 0.6
	Enabled	<b>DISABLED</b>	Enabled	Enabled	Enabled	Enabled	Enabled
	Innov 1	Innov 2	Innov 3	Innov 4	Innov 5	Innov 6	Innov 11

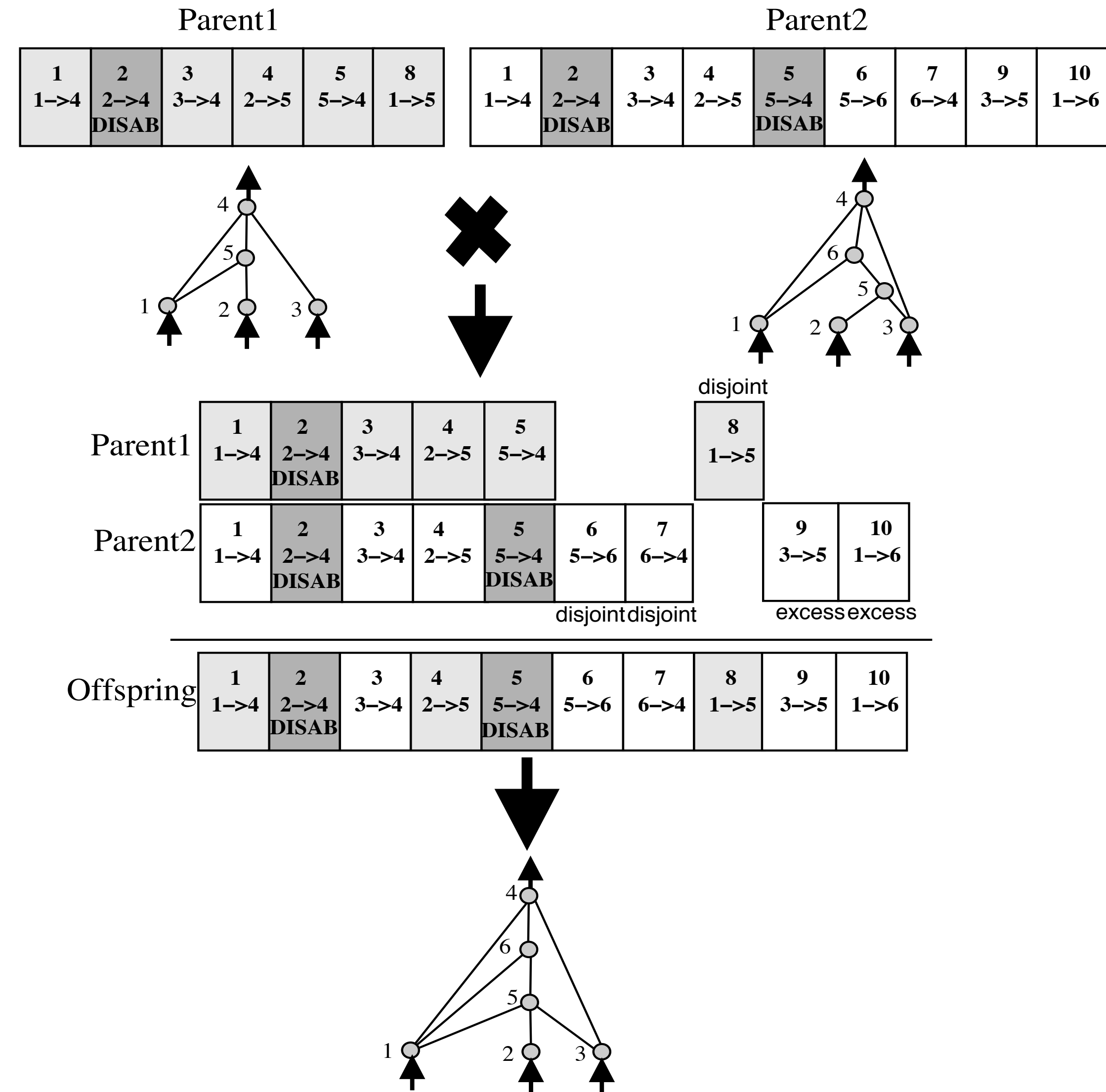
Network (Phenotype)



# NEAT: cross-over

## Neuro-Evolution of Augmenting Topologies (NEAT)

- Align the genes with the same innovation number
- Matching genes are inherited randomly,
- disjoint genes and excess genes are inherited from the more fit parent



# NEAT: speciation

## Neuro-Evolution of Augmenting Topologies (NEAT)

**Objective:** define niches by network topology

(so that a novel topology is "protected")

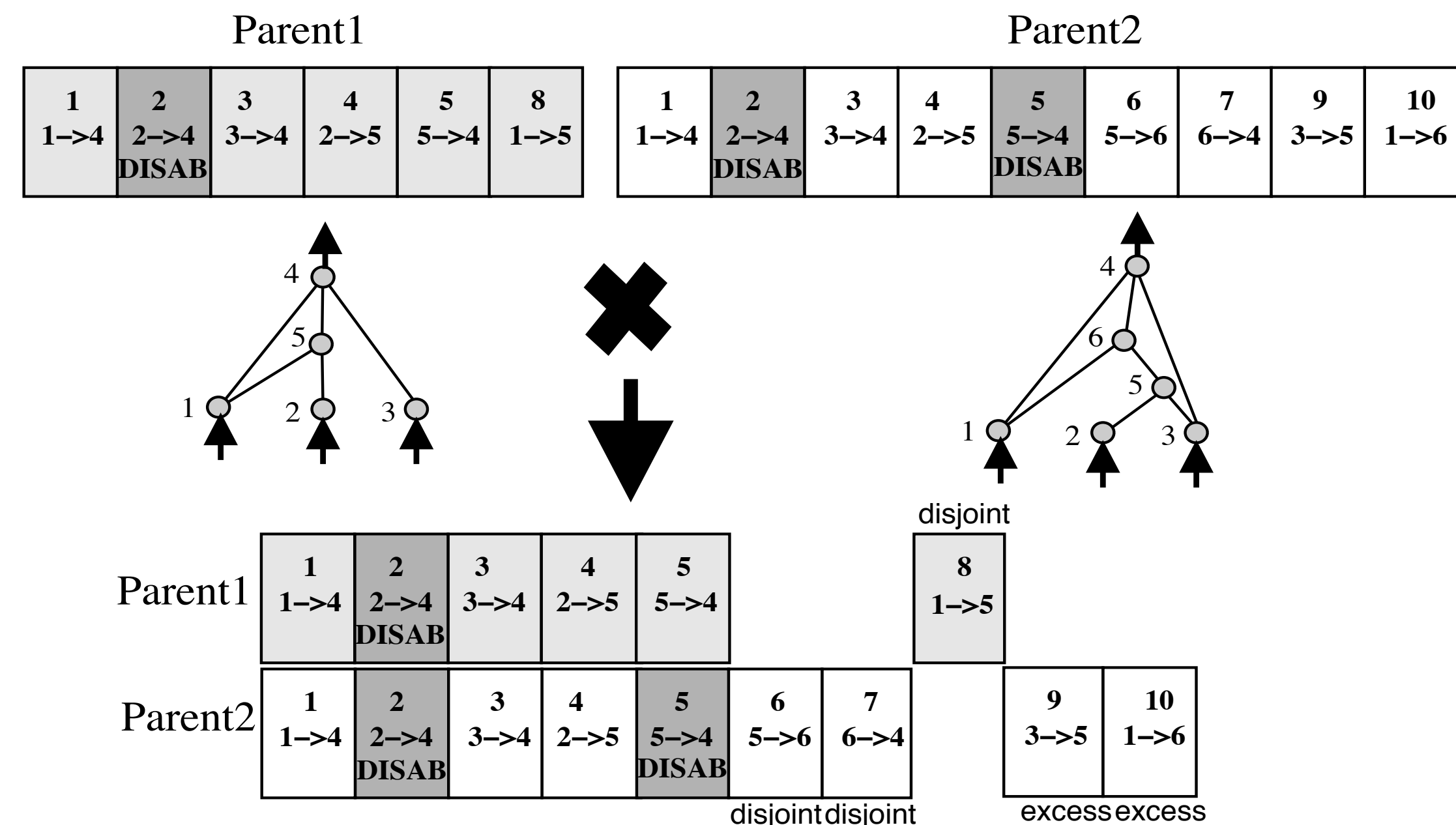
- Align the genes with the same innovation number
- Compute a distance between individuals:

$$\delta = \frac{c_1 E}{N} + \frac{c_2 D}{N} + c_3 \cdot \overline{W}.$$

Excess genes

Disjoint genes

Weight difference (joint genes)



Use fitness sharing:

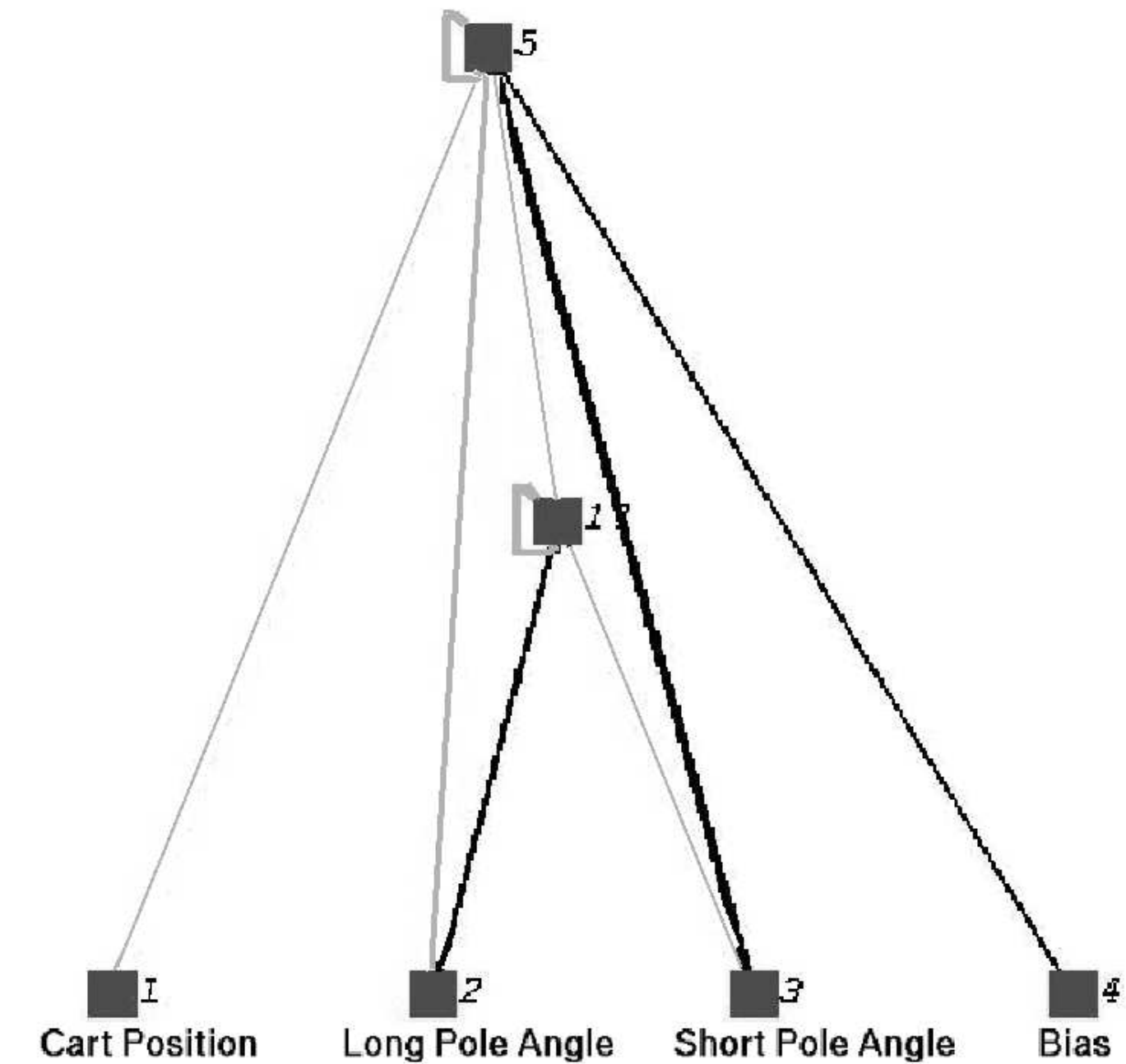
$$f'_i = \frac{f_i}{\sum_{j=1}^n \text{sh}(\delta(i, j))}.$$

1 if  $\delta(i, j) > \delta_{tr}$ , 0 otherwise

# NEAT: results

Method	Evaluations	Failure Rate
No-Growth NEAT (Fixed-Topologies)	30,239	80%
Nonspeciated NEAT	25,600	25%
Initial Random NEAT	23,033	5%
Nonmating NEAT	5,557	0
Full NEAT	3,600	0

double pole balancing with velocities



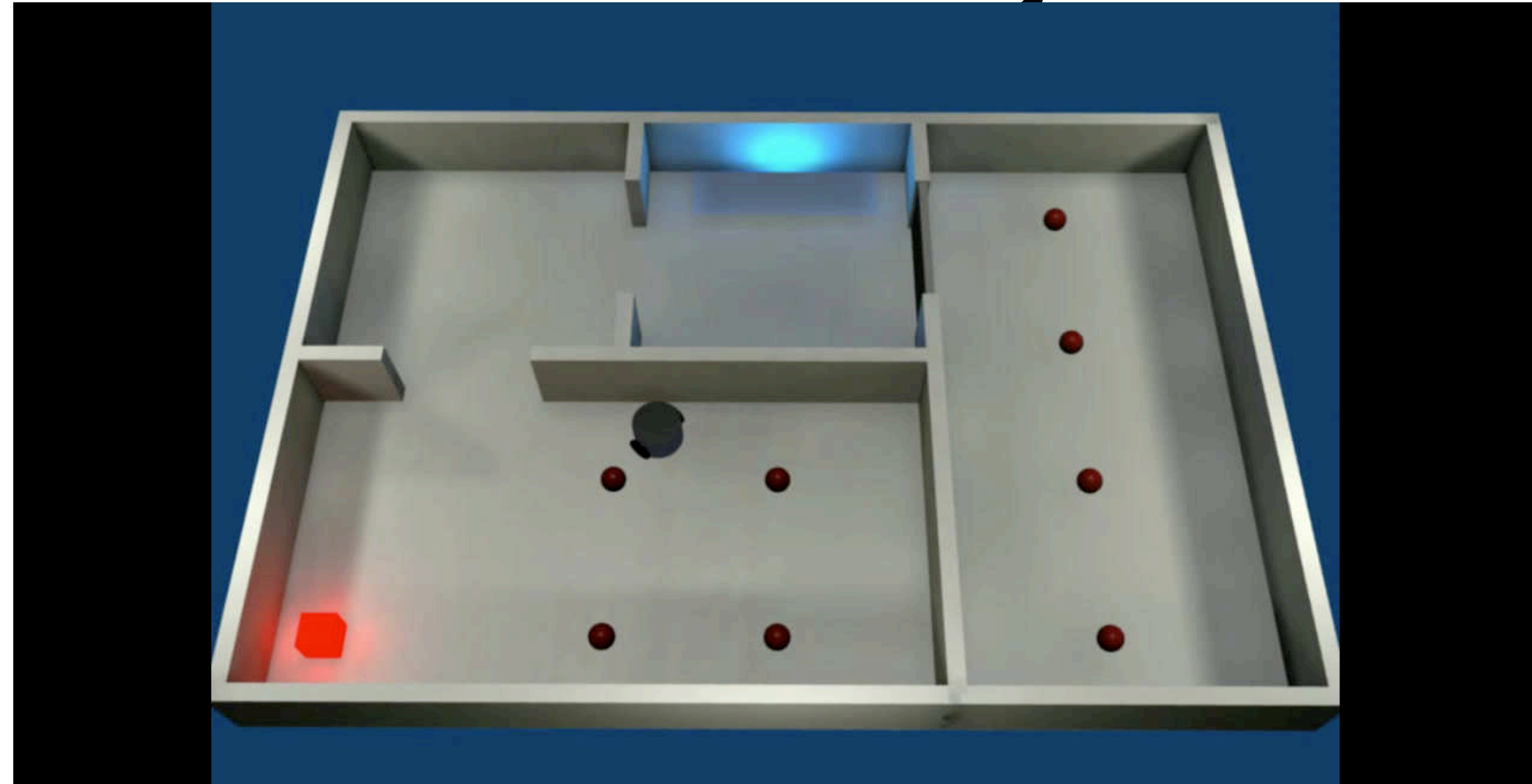
double inverse pendulum  
(note the recurrences!)

# Instead of speciation: behavioral diversity

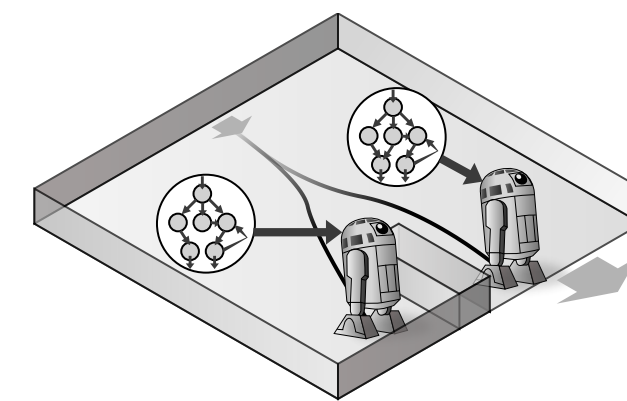
- Define a distance between behaviors  
e.g., diff. in trajectory or sensor streams
- Use NSGA-II to maximize:

$$\begin{cases} \textit{Fitness}(x) \\ \sum_i d(x, P_i) \end{cases}$$

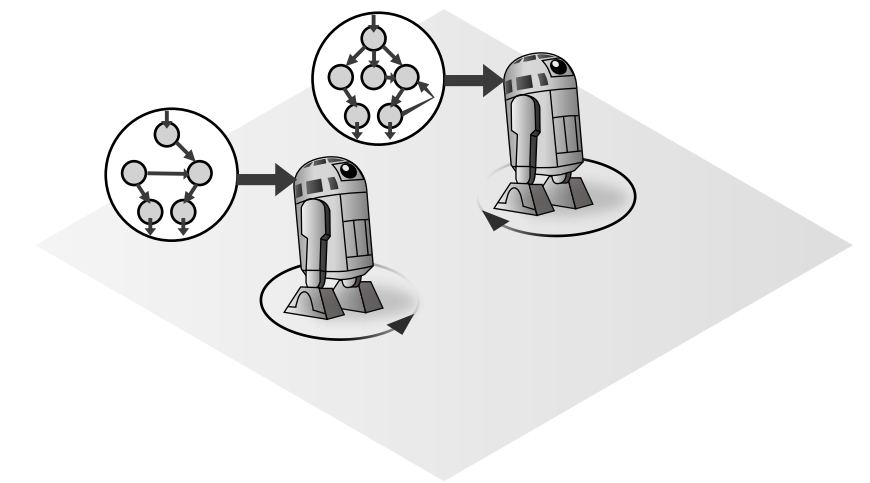
Diversity with regards  
to the population



**The key for evolving topologies is diversity**



Close genotype / different behavior



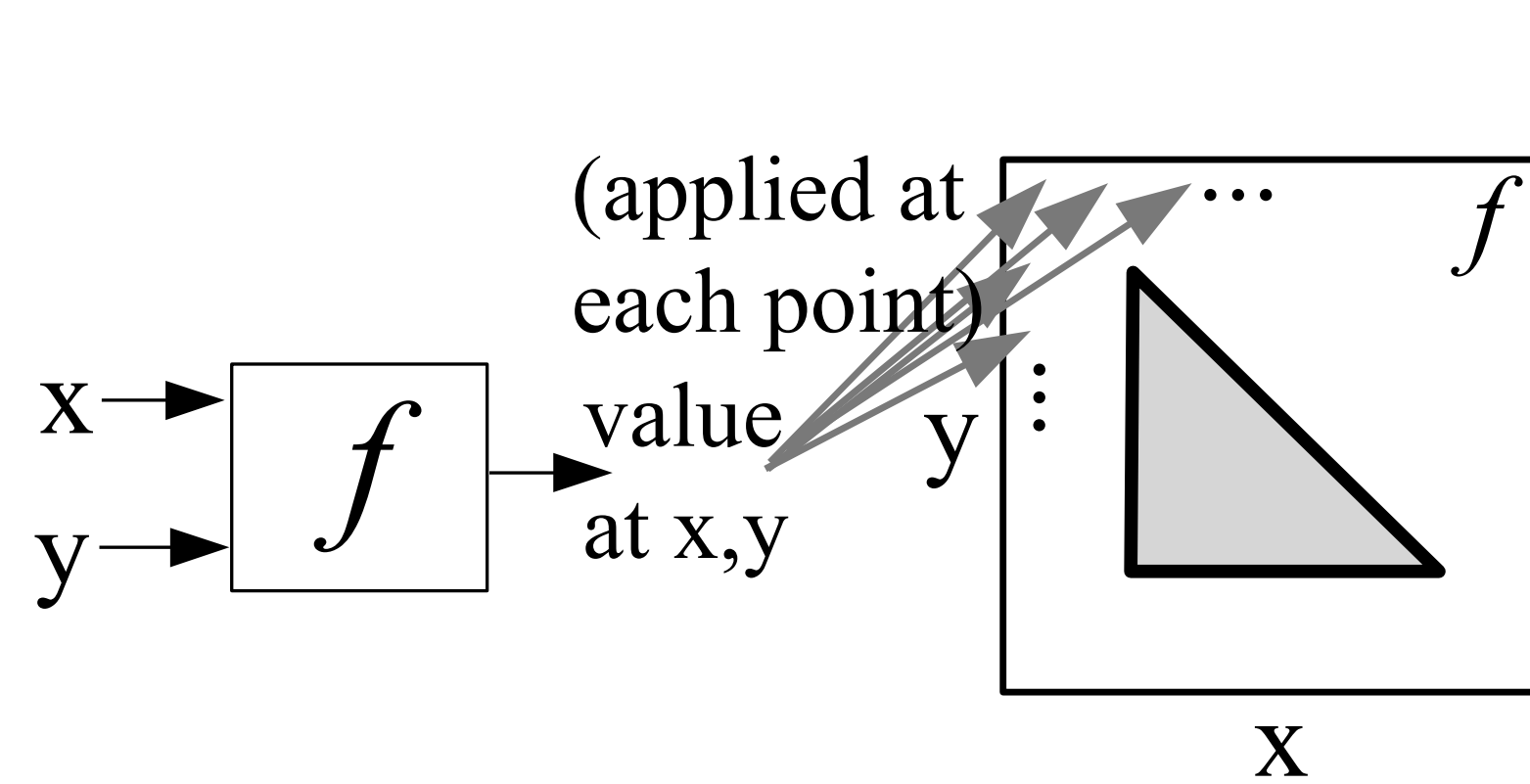
Different genotype / same behavior

**Mouret, J. B., & Doncieux, S. (2012).** Encouraging behavioral diversity in evolutionary robotics: An empirical study. *Evolutionary computation*, 20(1), 91-133.

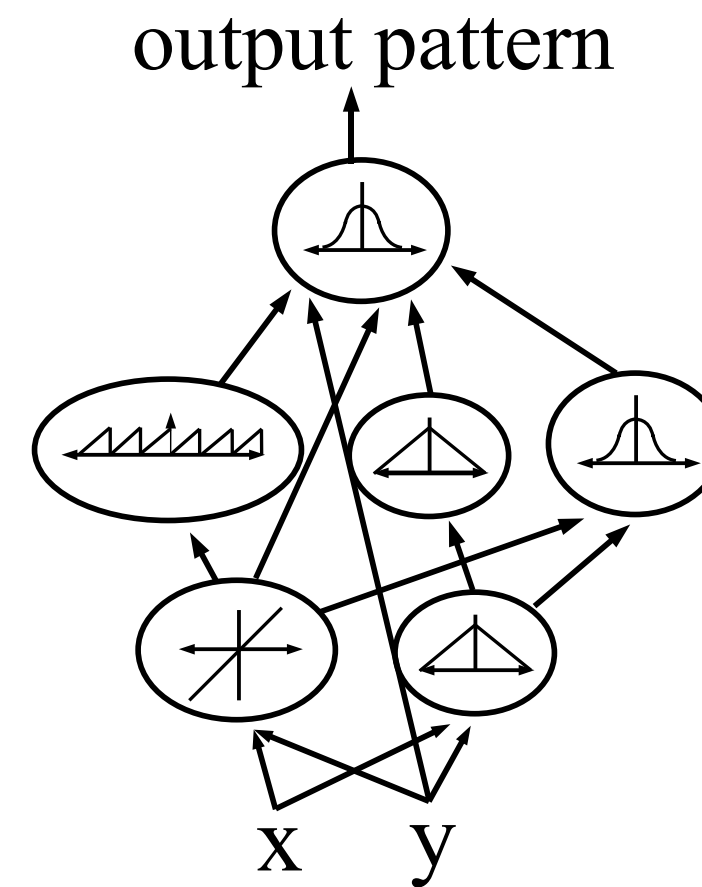
**Doncieux, S., & Mouret, J. B. (2014).** Beyond black-box optimization: a review of selective pressures for evolutionary robotics. *Evolutionary Intelligence*, 7(2), 71-93.



# Finding natural patterns



(a) Mapping



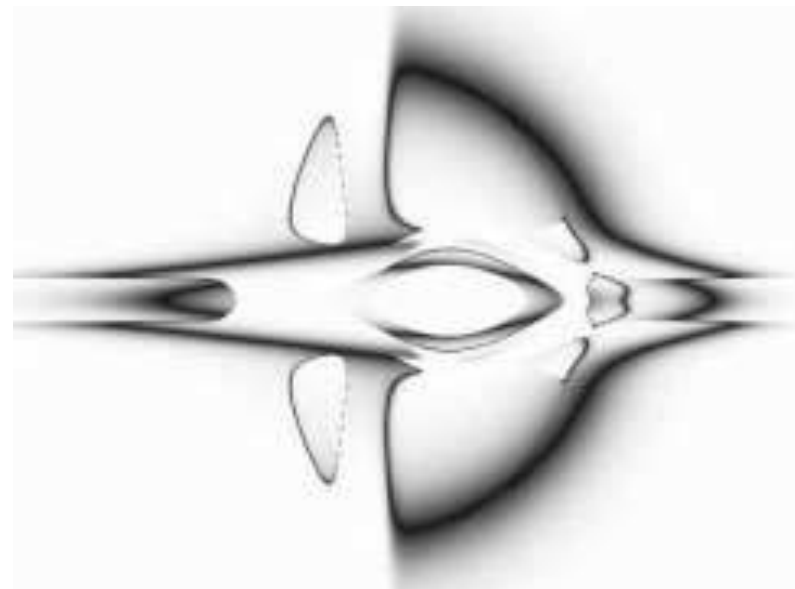
(b) Composition

Evolved with NEAT

Additional mutation: change the activation function

Functions:

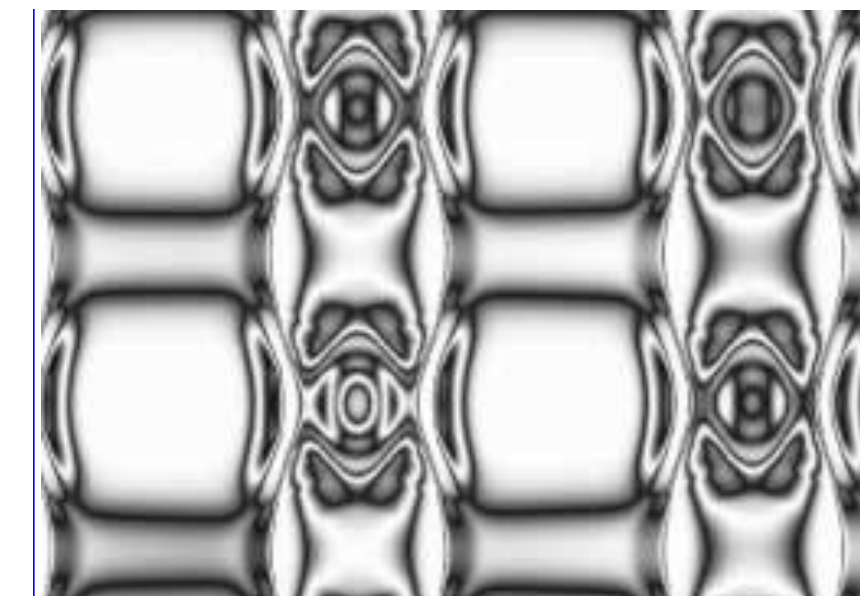
- sine wave (repetition)
- Gaussian (symmetry)
- sigmoid
- ...



Symmetry



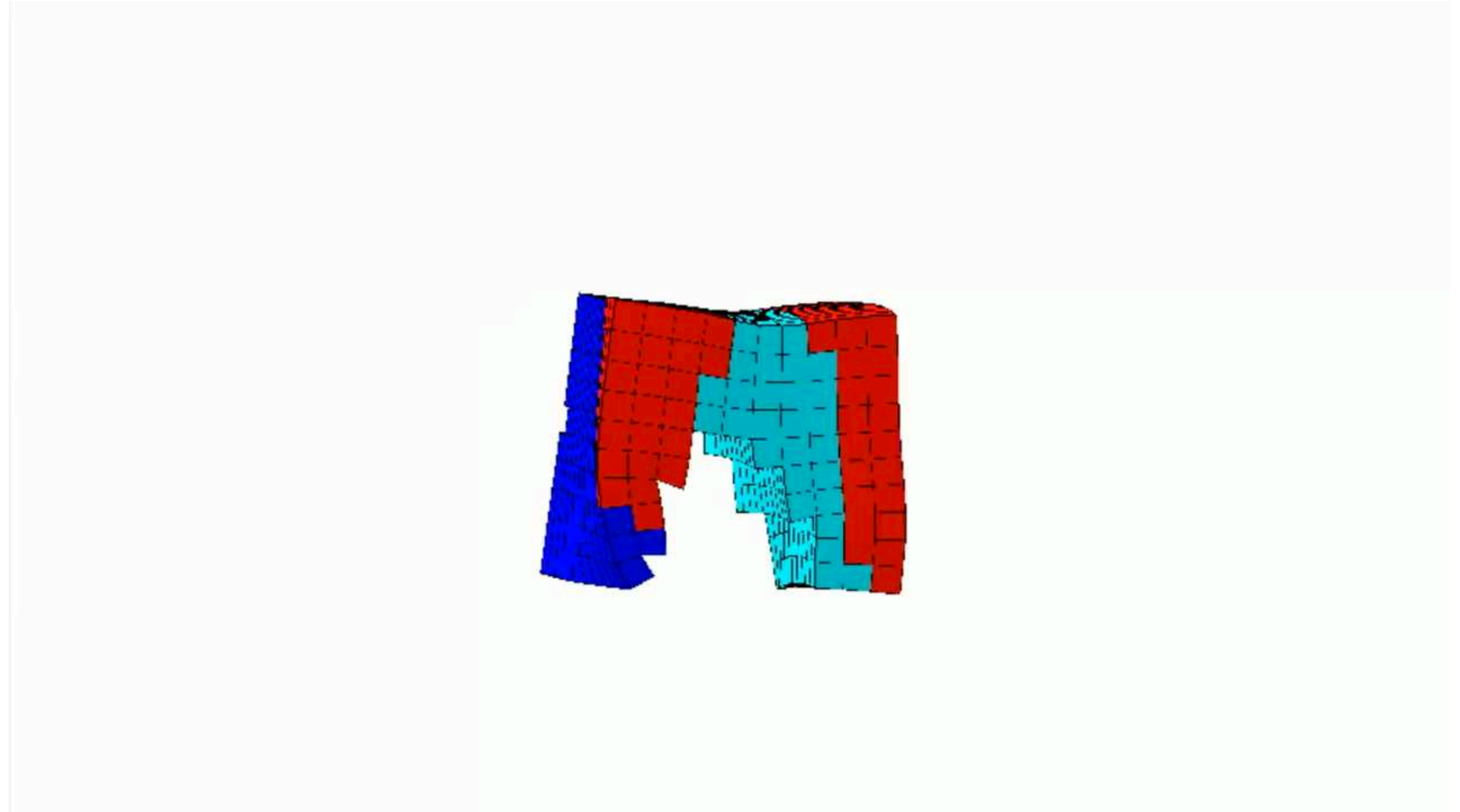
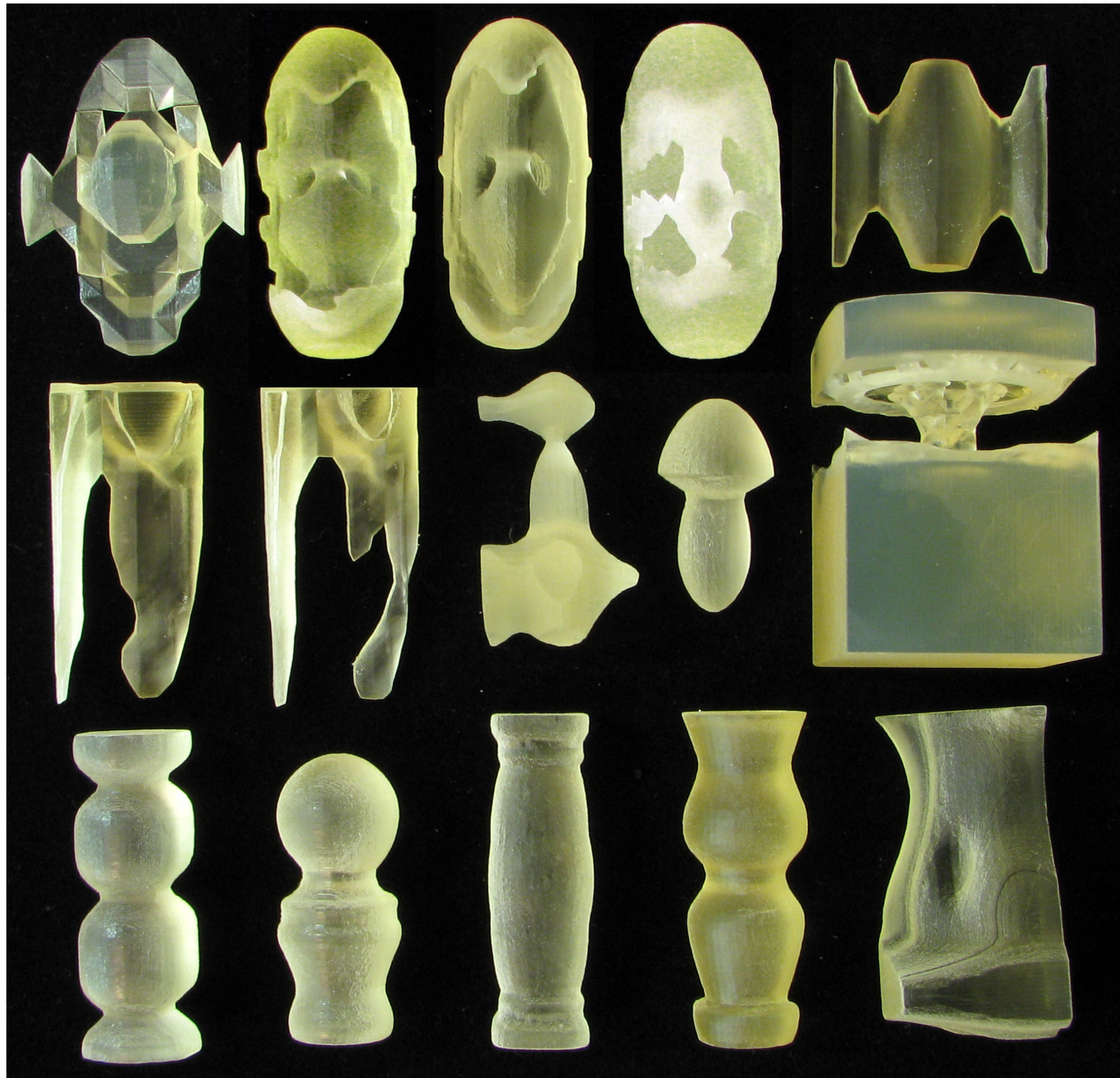
Imperfect symmetry



Repetition with variation



# Finding natural patterns



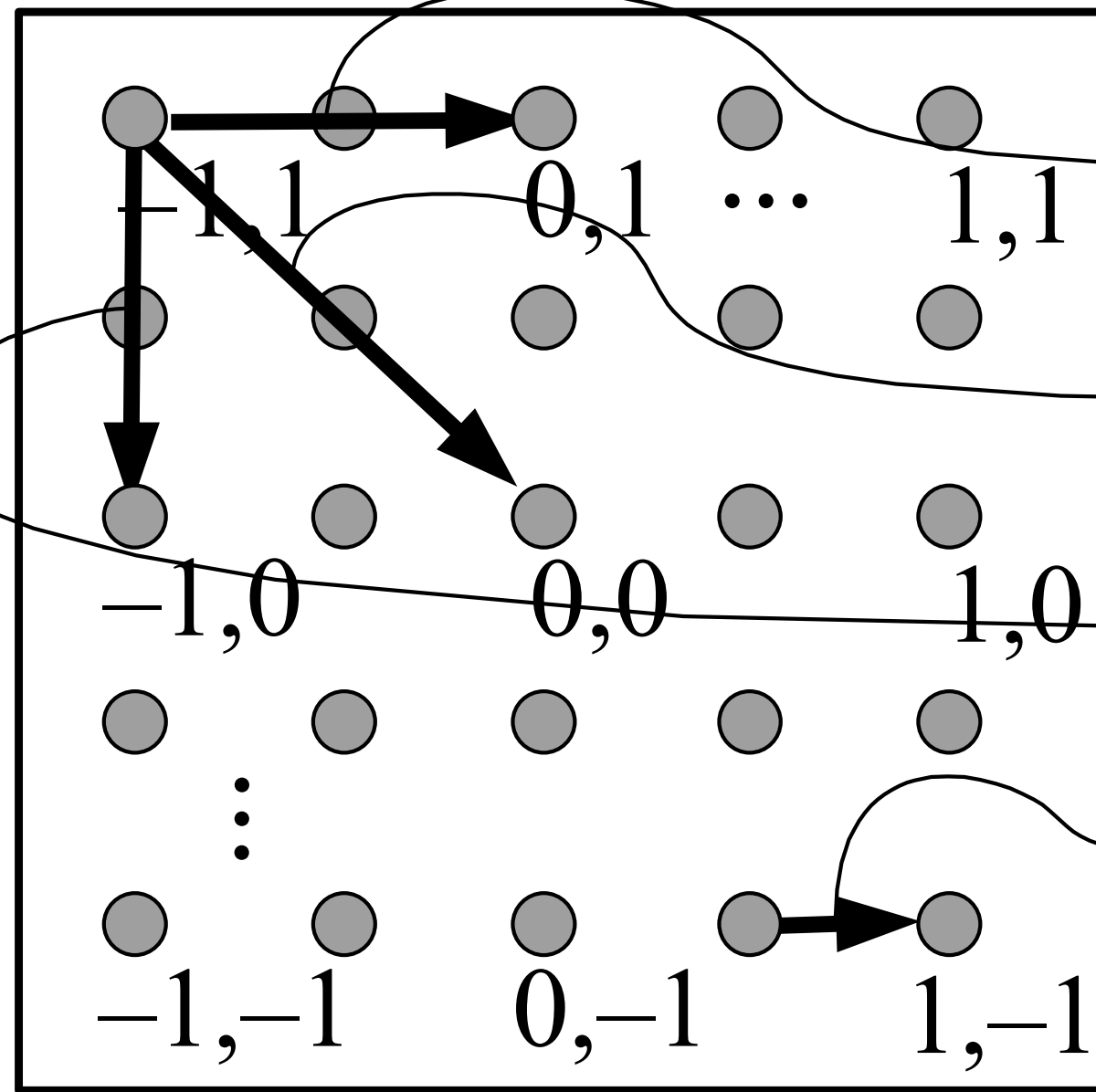
**Clune, Jeff, and Hod Lipson. (2011)** "Evolving three-dimensional objects with a generative encoding inspired by developmental biology." Proc. of *ECAL*. 2011.

**Cheney N, MacCurdy R, Clune J, Lipson H. (2013)** Unshackling evolution: evolving soft robots with multiple materials and a powerful generative encoding. In. *Proc of GECCO* 2013



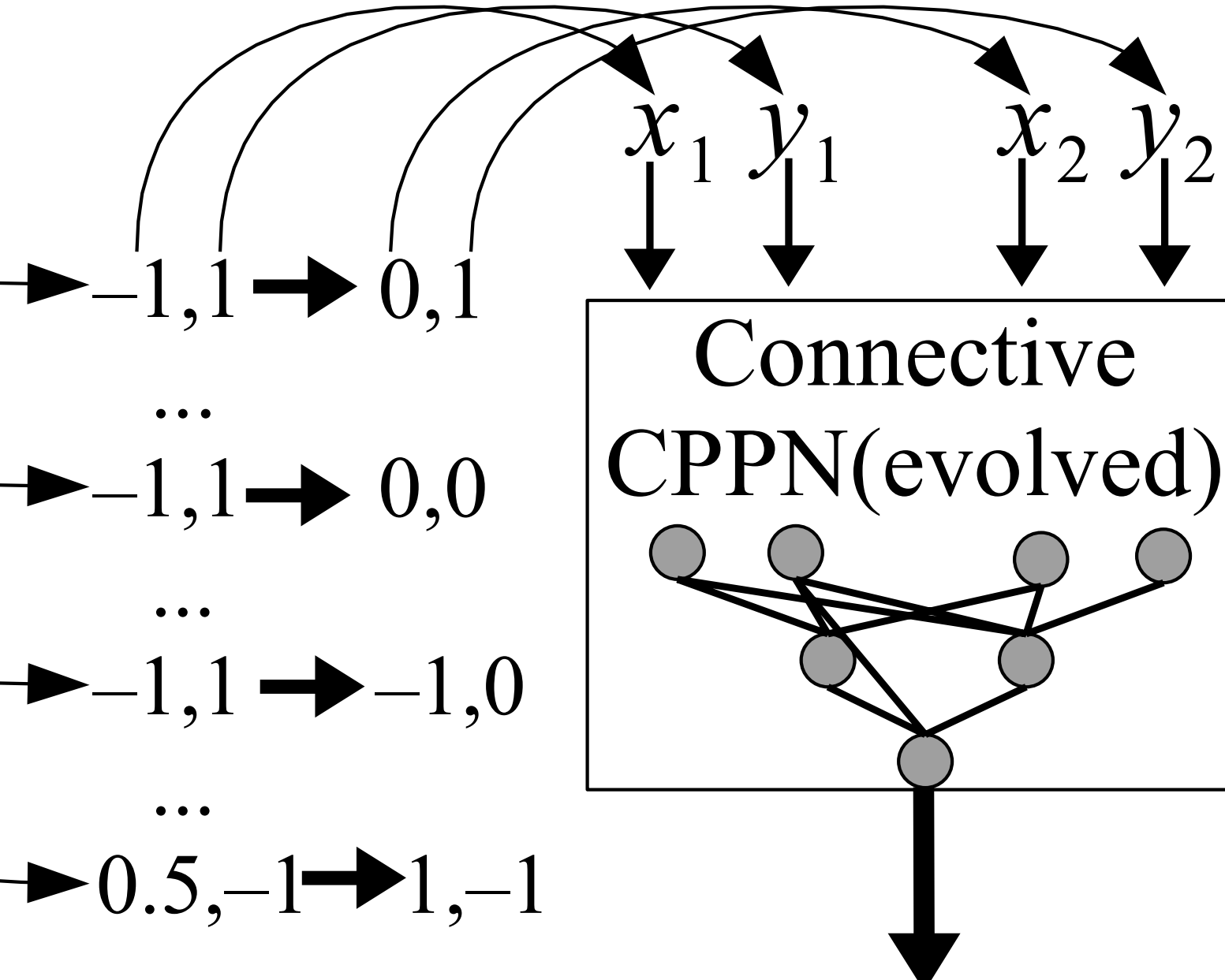
# Connection patterns

1) Query each potential connection on substrate



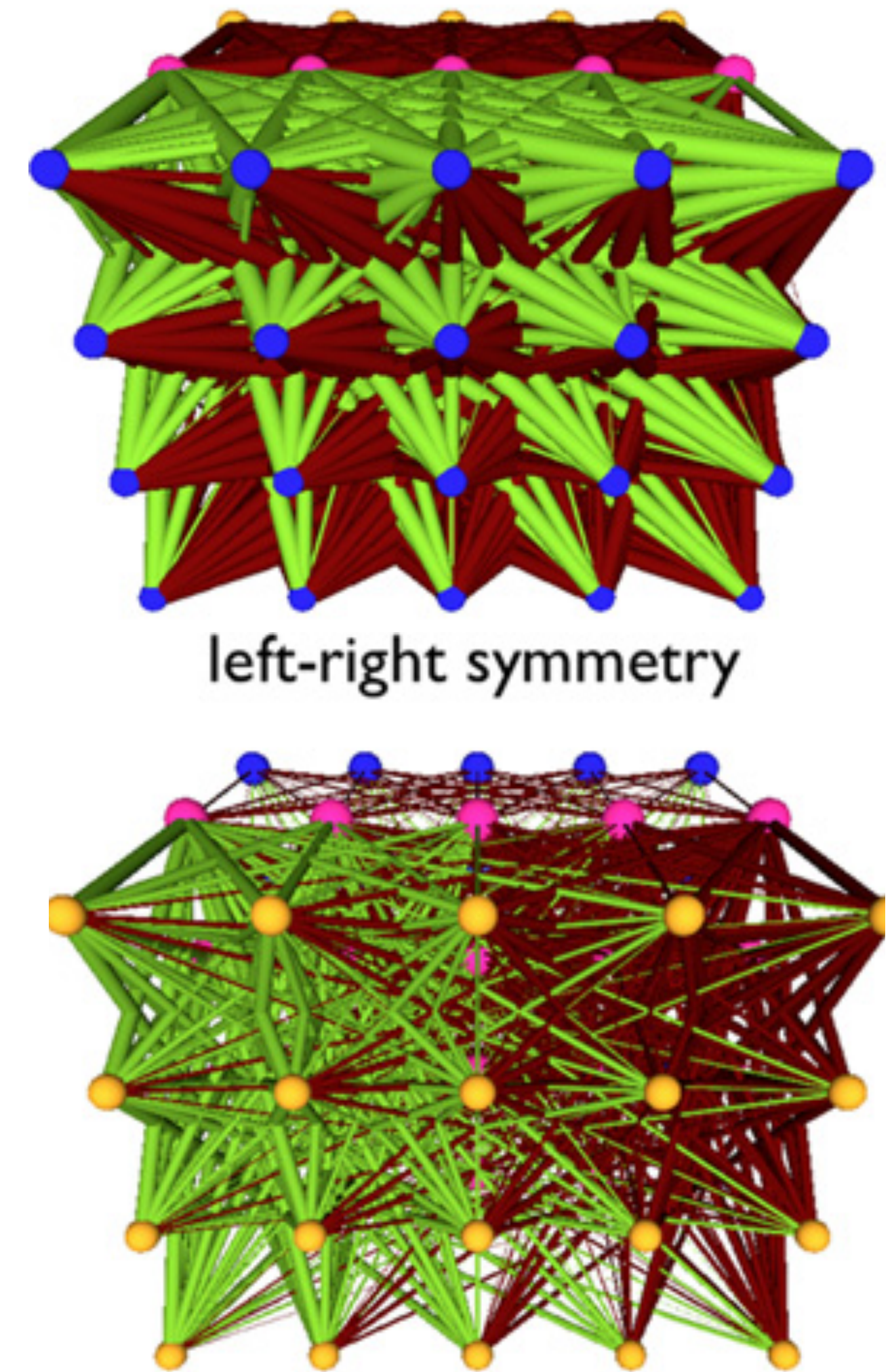
Substrate

2) Feed each coordinate pair into CPPN



3) Output is weight between  $(x_1, y_1)$  and  $(x_2, y_2)$

repeat for each node



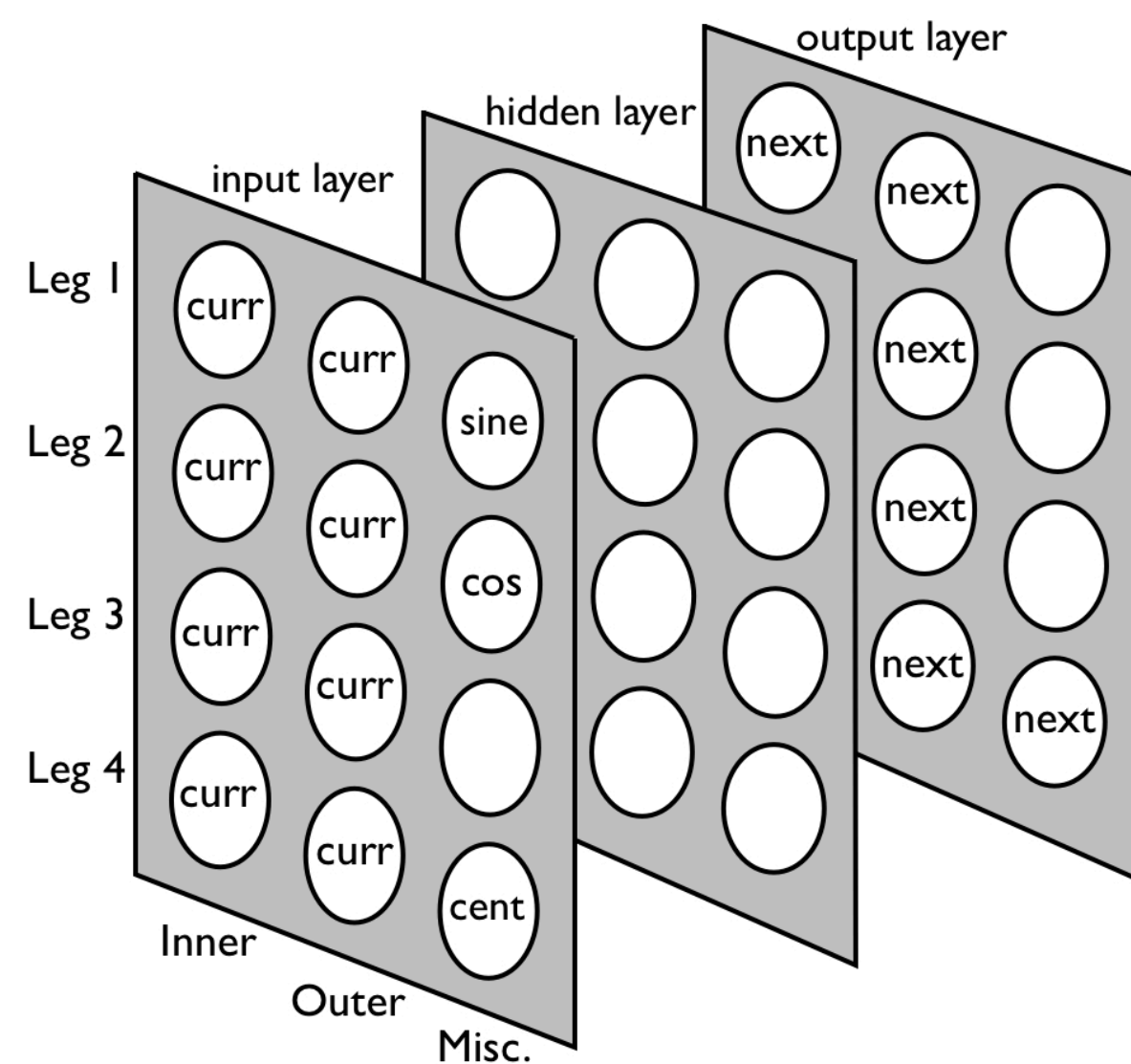
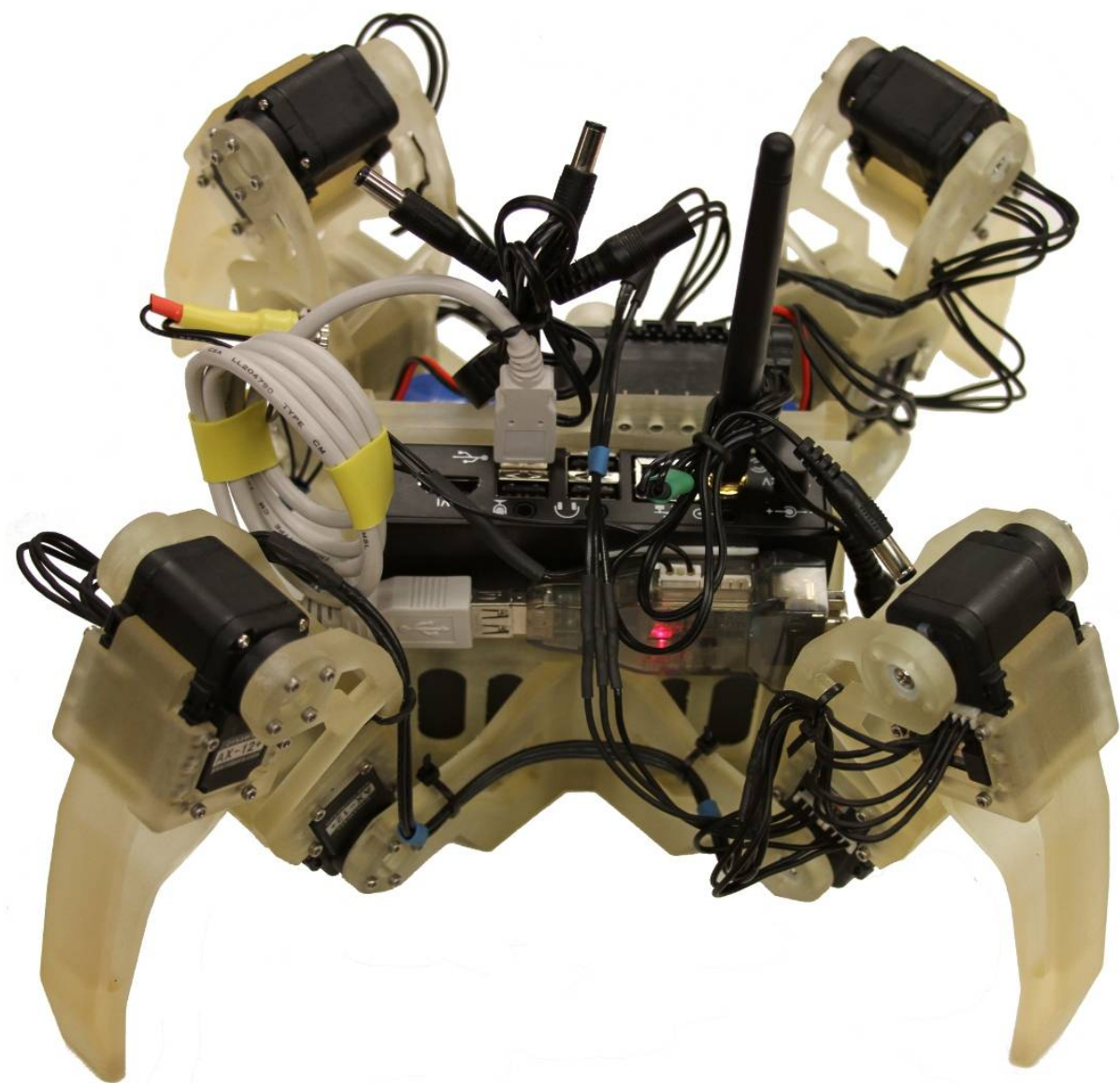
**Scales to million of neurons!**

Stanley, Kenneth O., David B. D'Ambrosio, and Jason Gauci. (2009) "A hypercube-based encoding for evolving large-scale neural networks." *Artificial life* 15.2 (2009): 185-212.

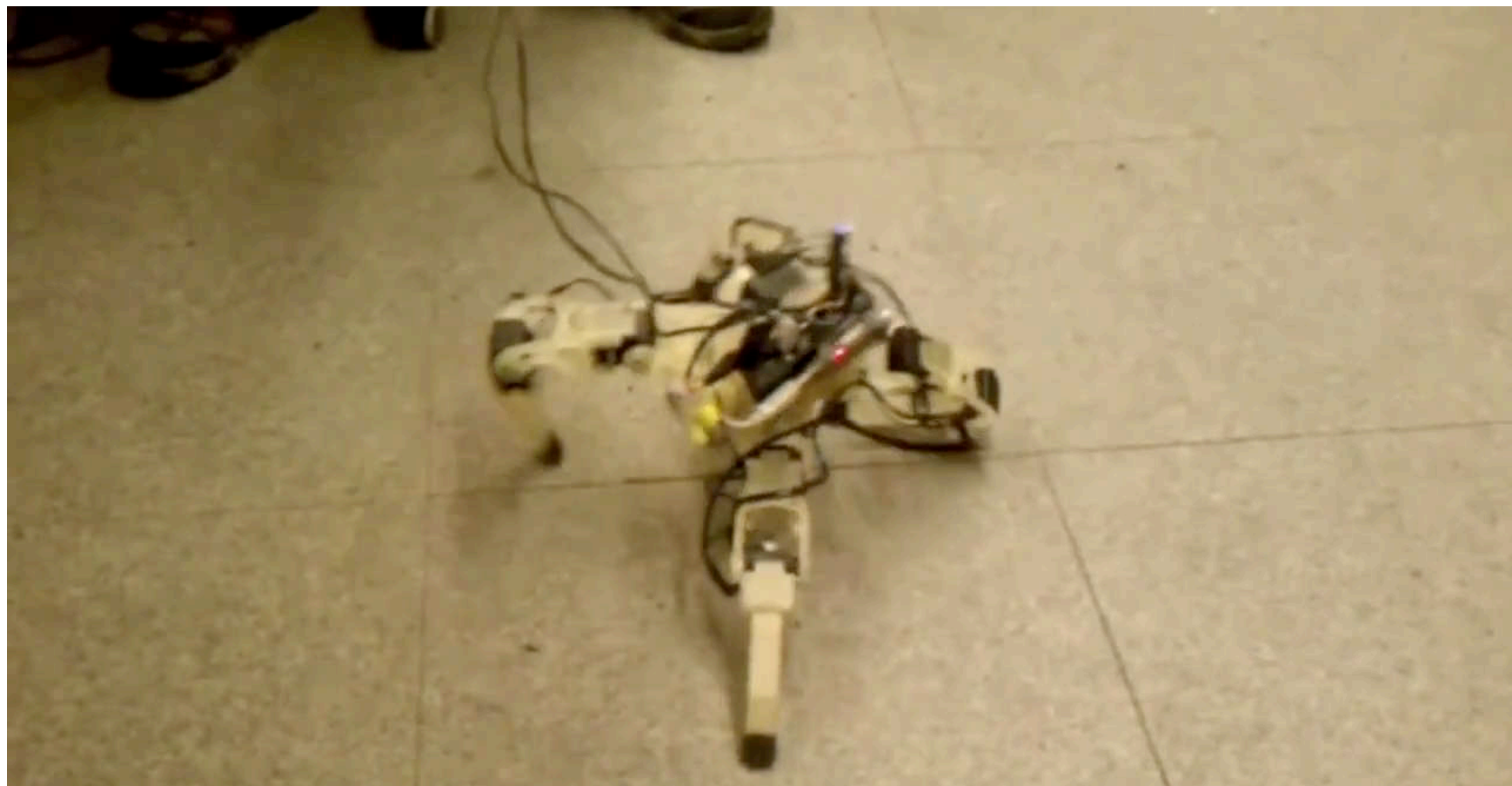
Clune J, Stanley KO, Pennock RT, Ofria C. (2011) On the performance of indirect encoding across the continuum of regularity. *IEEE Transactions on Evolutionary Computation*. 2011 Jun;15(3):346-67.



# HyperNEAT: learning gaits



	Average	Std. Dev.
Previous hand-coded gait	5.16	–
Random search	9.40	6.83
Uniform Random Hill Climbing	7.83	4.56
Gaussian Random Hill Climbing	10.03	6.00
Policy Gradient Descent	6.32	7.39
Nelder-Mead simplex	12.32	3.35
Linear Regression	14.01	12.88
Evolved Neural Network (HyperNEAT)	29.26	6.37



Yosinski, J., Clune, J., Hidalgo, D., Nguyen, S., Zagal, J. C., & Lipson, H. (2011). Evolving robot gaits in hardware: the HyperNEAT generative encoding vs. parameter optimization. In ECAL (pp. 890-897).