

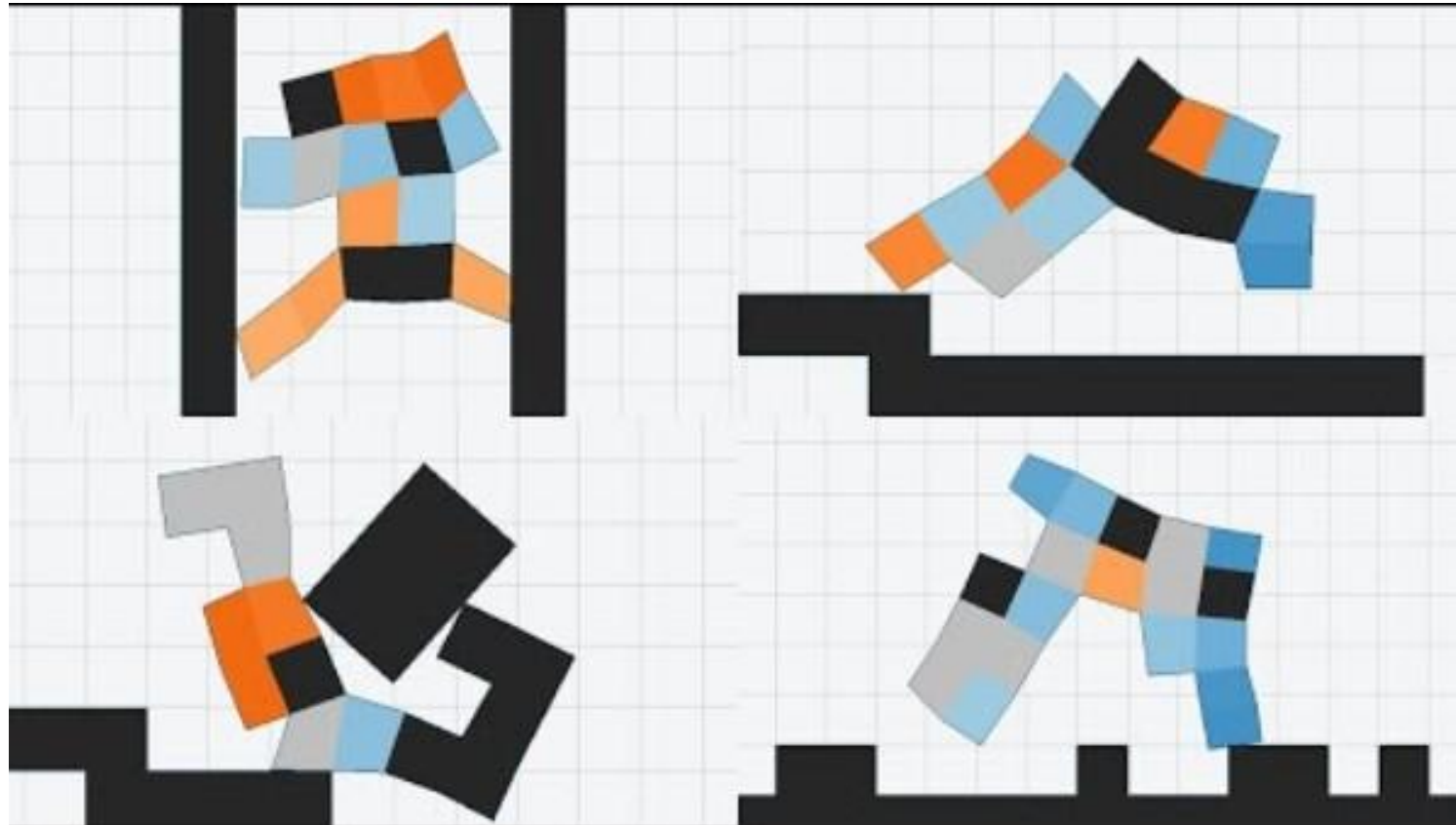
Diploidy and Indirect-Encoding, new ways to explore Body-Brain co-optimization in 2D single genome soft-robot

MATTEO VACHER

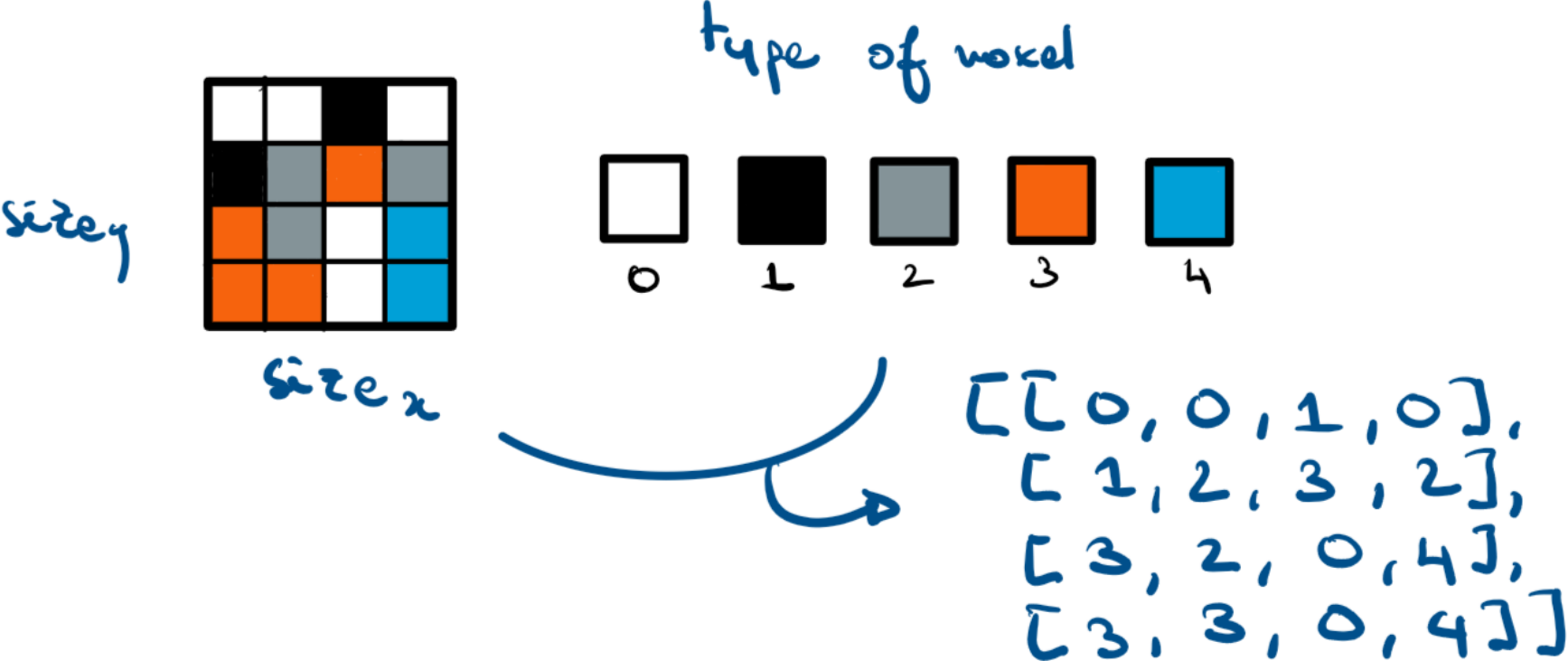
ENGINEERING STUDENT AT ISAE-SUPAERO

EXCHANGE RESEARCH STUDENT AT TSUKUBA UNIVERSITY

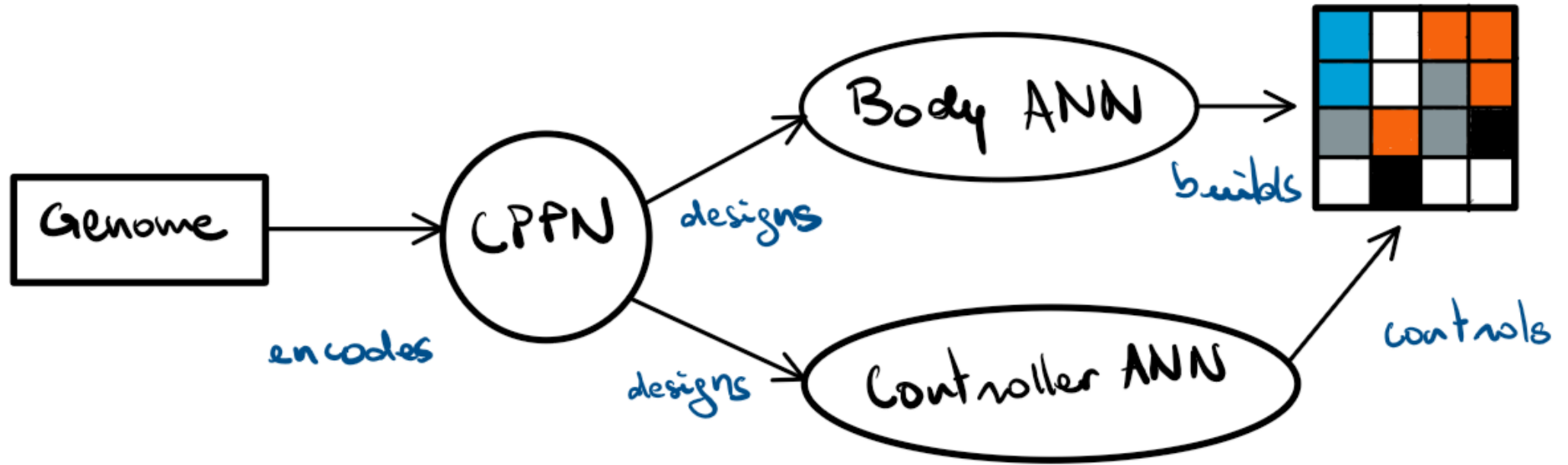
Evogym



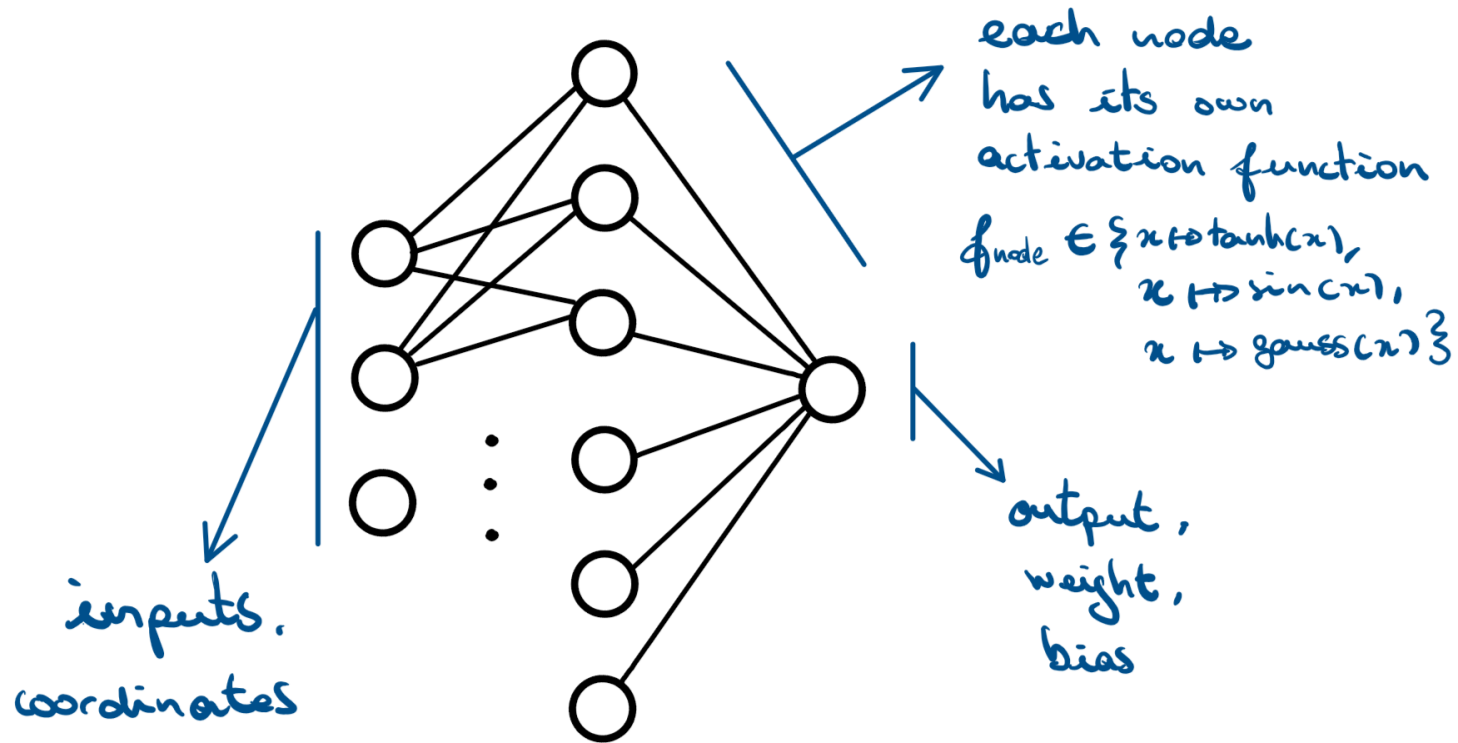
What is a Soft-Robot



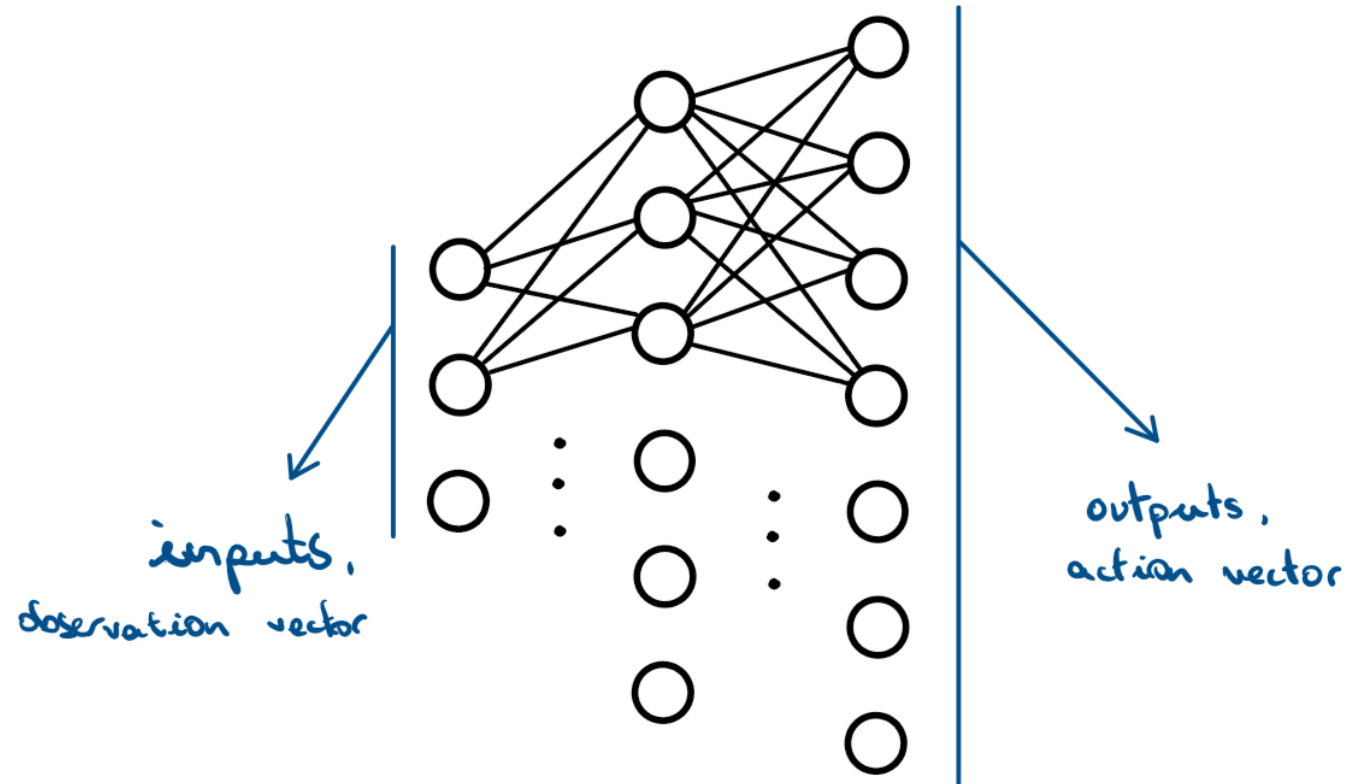
Indirect Encoding



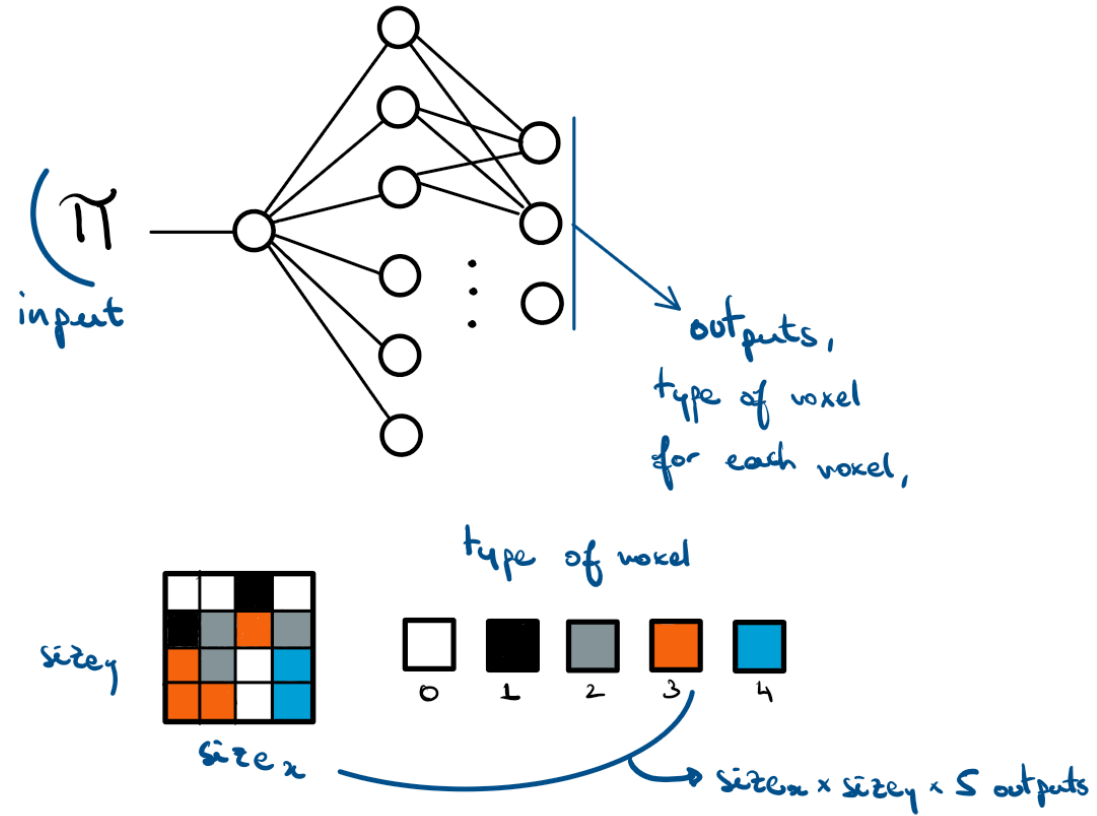
Compositional Pattern Producing Network



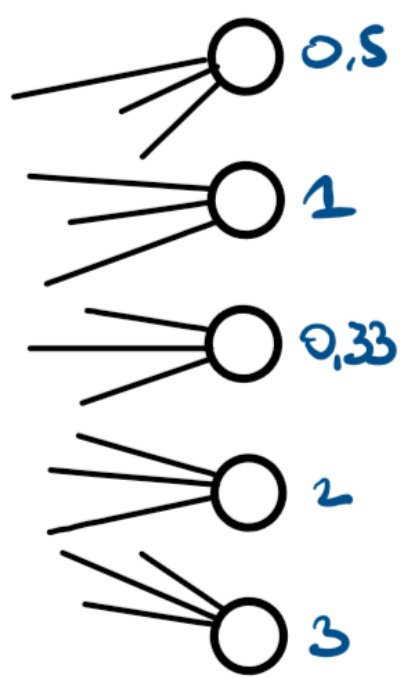
Controller Artificial Neural Network



Body Producer Neural Network



Example



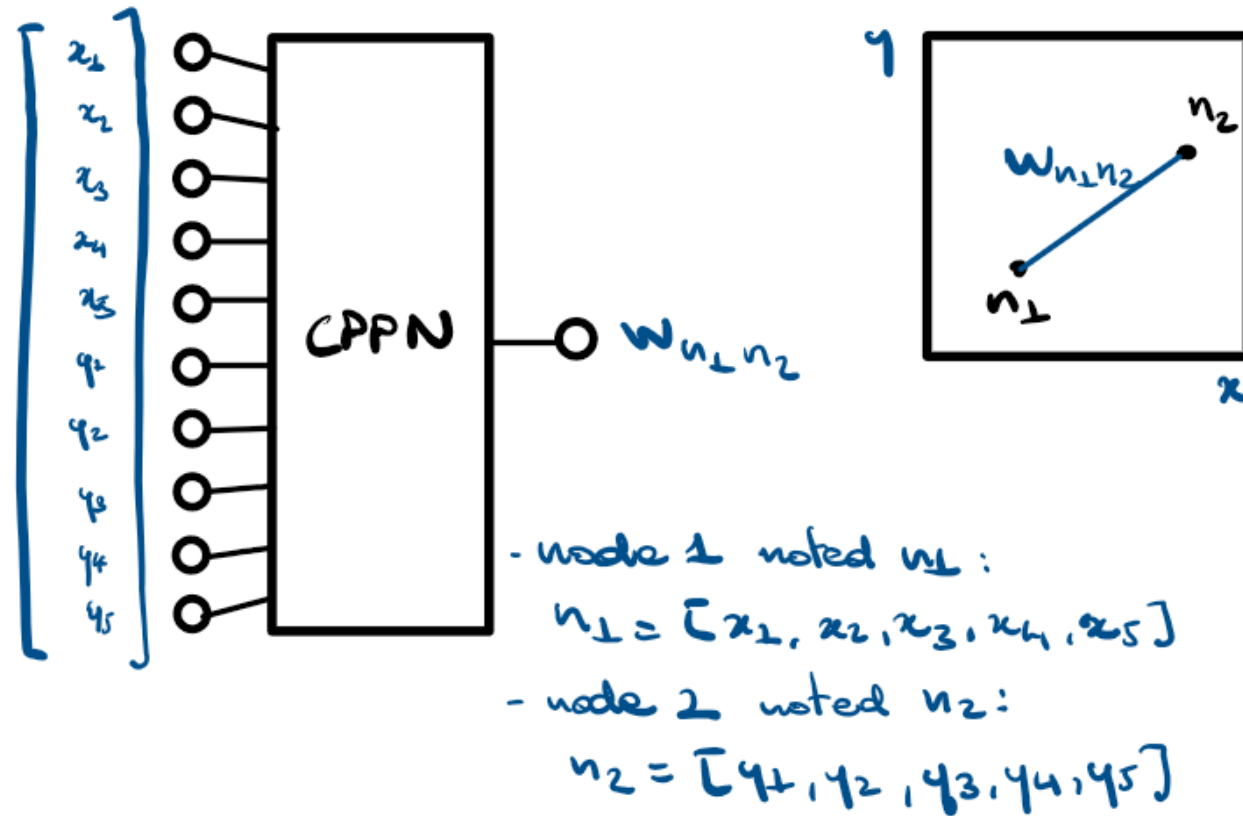
output for voxel (i, j)

output = [0,5; 1; 0,33; 2, 3]

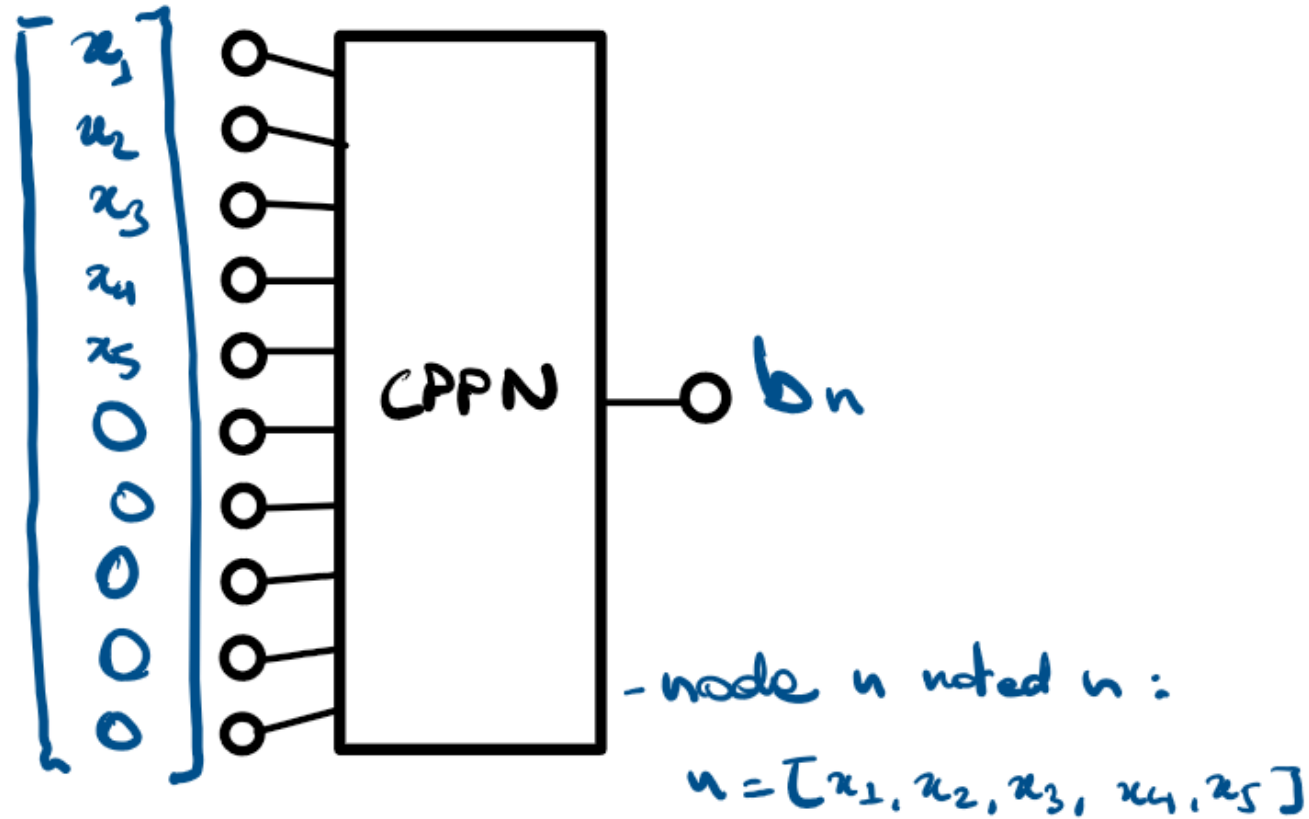
type = $\text{argmax}(\text{output}) = 4$



CPPN and Weights



CPPN and Biases



The Substrate

Cartesian product:

E_1, E_2, \dots, E_n set of elements:

$$E_1 \times E_2 \times \dots \times E_n = (x_1, x_2, \dots, x_n)$$

for $x_1 \in E_1, x_2 \in E_2, \dots, x_n \in E_n$

Body ANN substrate:

for n nodes in neural network

$$n = [x_1, x_2, x_3, x_4] \in E_1 \times \dots \times E_n$$

How can every point represent
the reality of the problem?

Building E_1, E_2, \dots, E_4

$$E_1 = \{0,33; 0,66; 1\}$$

$$E_2 = \{-1; -0,5; 0; 0,5; 1\}$$

$$E_3 = \{-1; -0,5; 0; 0,5; 1\}$$

$$E_4 = \{(\cos(k\frac{2\pi}{5}); \sin(k\frac{4\pi}{5})); k \in \mathbb{I}[0; 4]\}$$

\hookrightarrow $n = [0,66; -0,5; 1; \cos(2\frac{2\pi}{5}); \sin(2\frac{4\pi}{5})]$

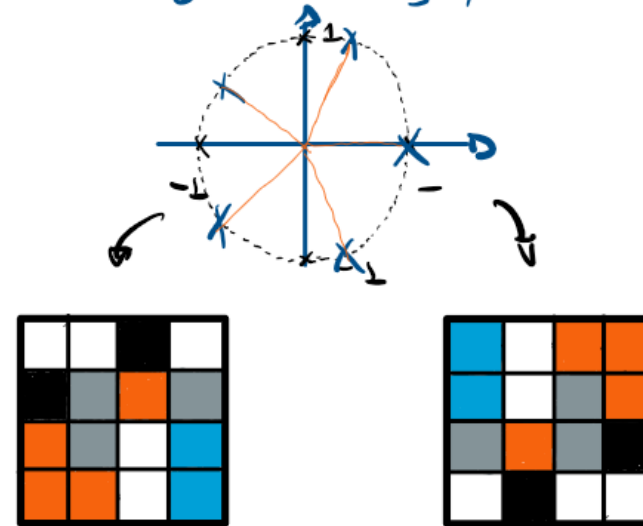
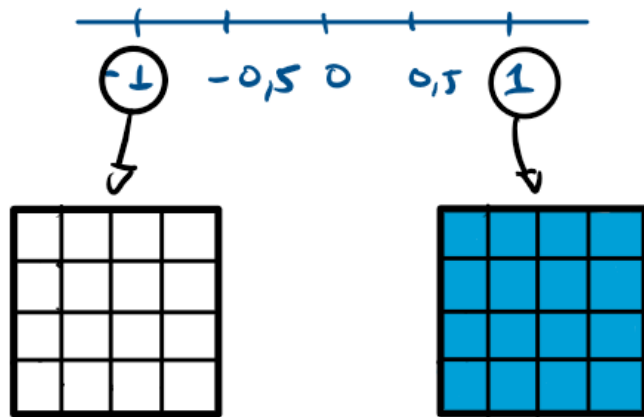
The problem

Side effect

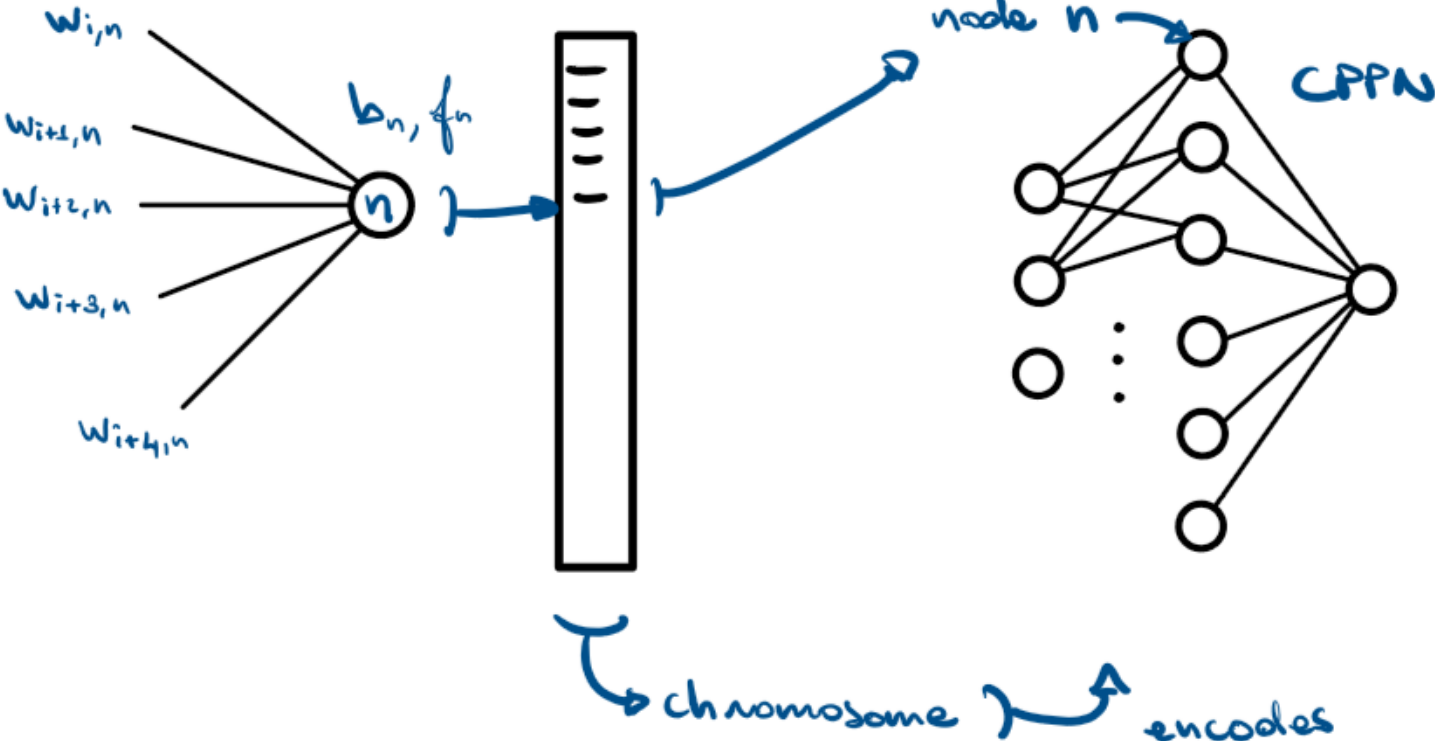
$E_4 = \text{type of vowel}$

$$E_4 = \{-1; -0.5; 0; 0.5; 1\}$$

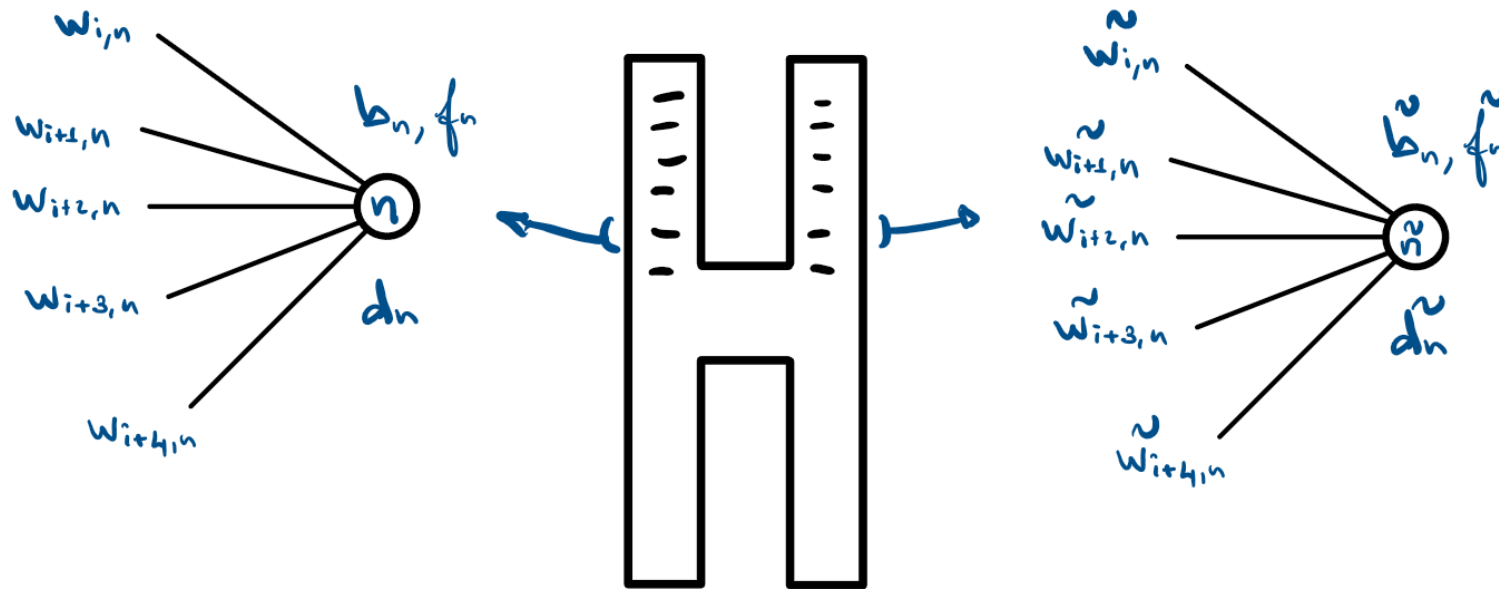
$$E_4 = \{(\cos(k\frac{2\pi}{5}); \sin(k\frac{4\pi}{5})) ; k \in \{0; 1; 2; 3; 4\}\}$$



Haploidy



Diploidy

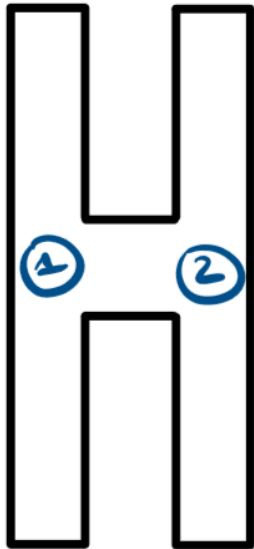
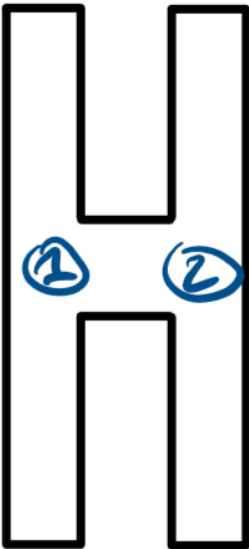


$d_n, \tilde{d}_n \in [0, SD]$, compare (d_n, \tilde{d}_n)

Crossover

Parent 1

Parent 2



→
random
selection

Parent 1

Parent 2

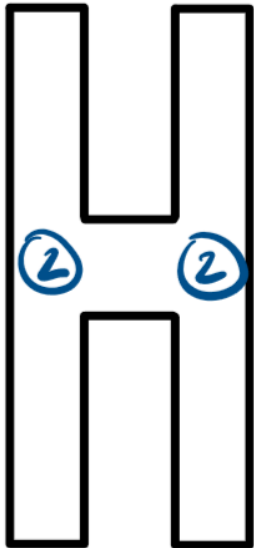


+

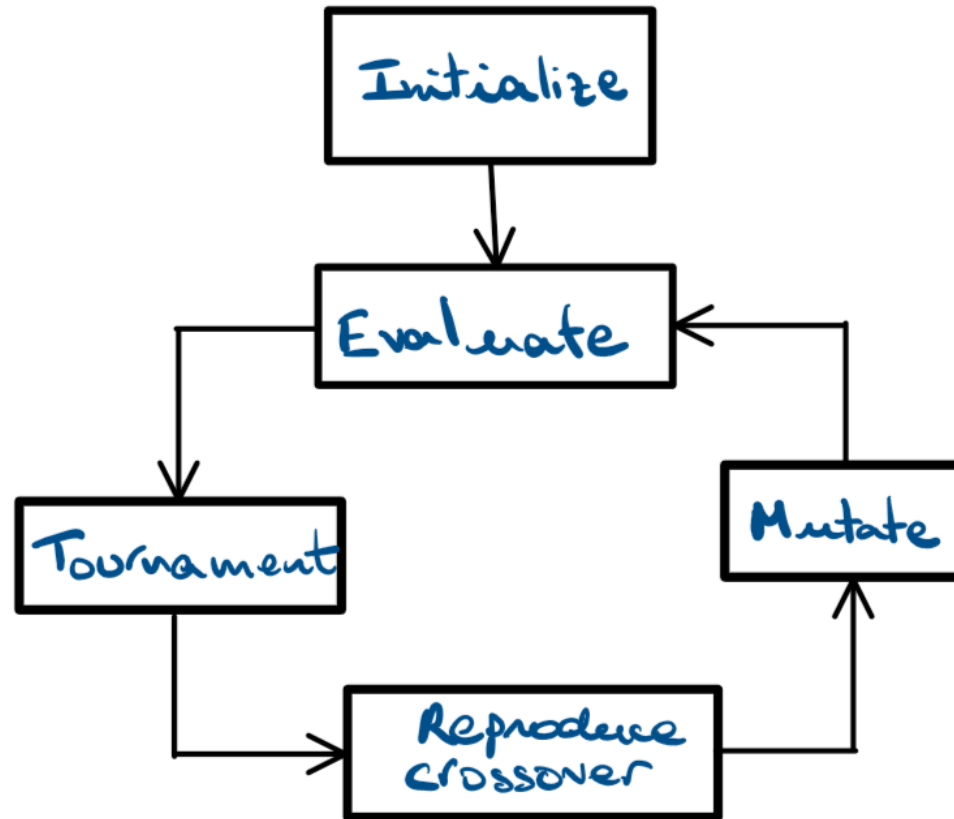


→
fusion

New individual



The Algorithm



The benefits

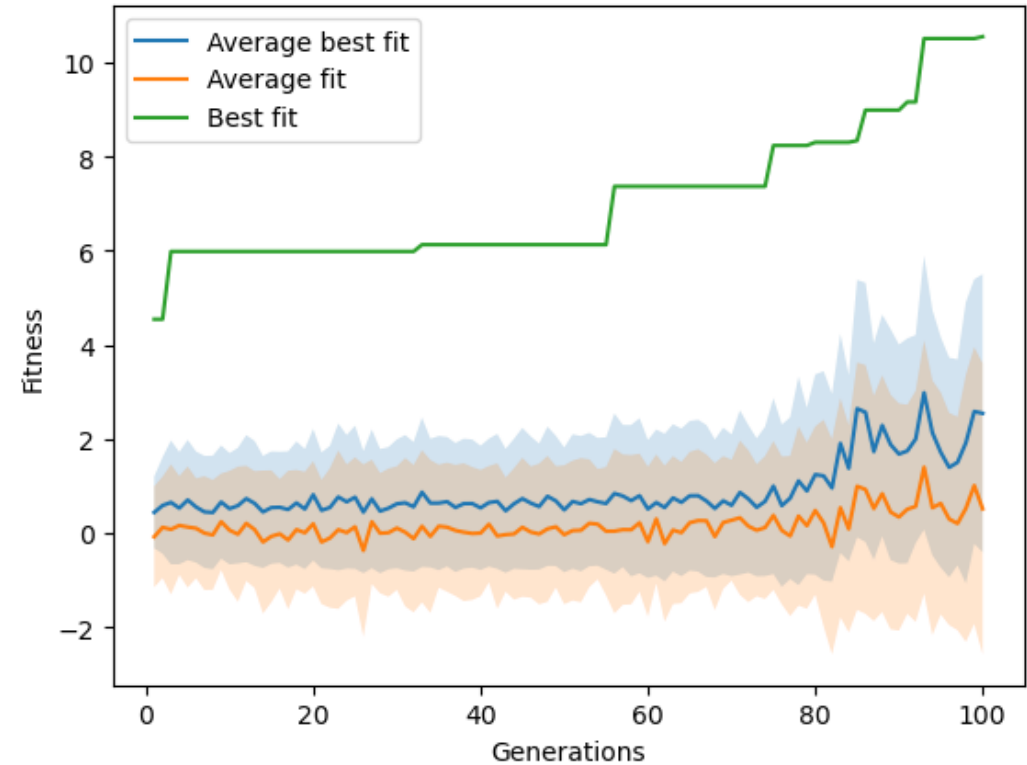
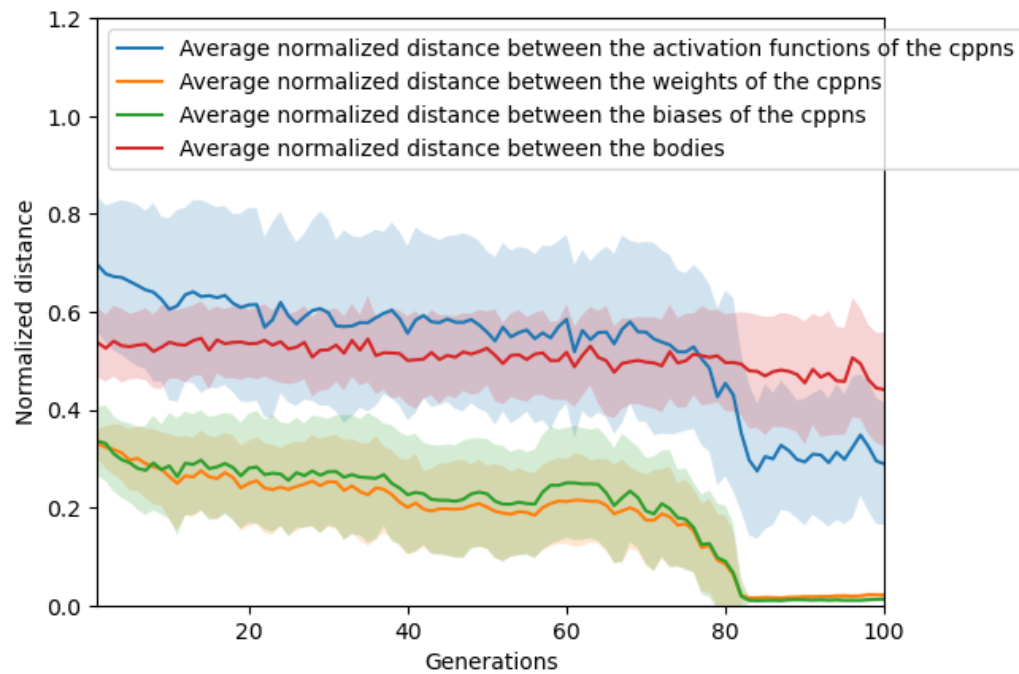
- Greater ability to search complex spaces and find better optima.
- Better adaptation to changing environments
- Less sensitive to mutation rate, i.e. to destructive mutations
- Depending on the crossover, can match the results of haploid systems on stationary problems

The Parameters

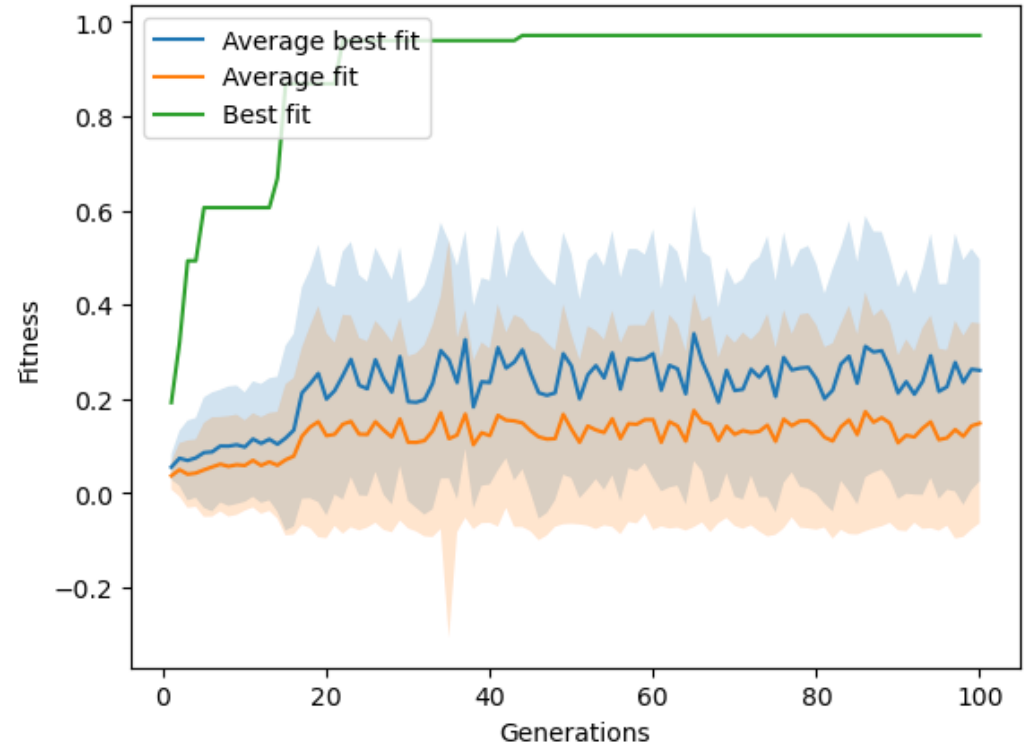
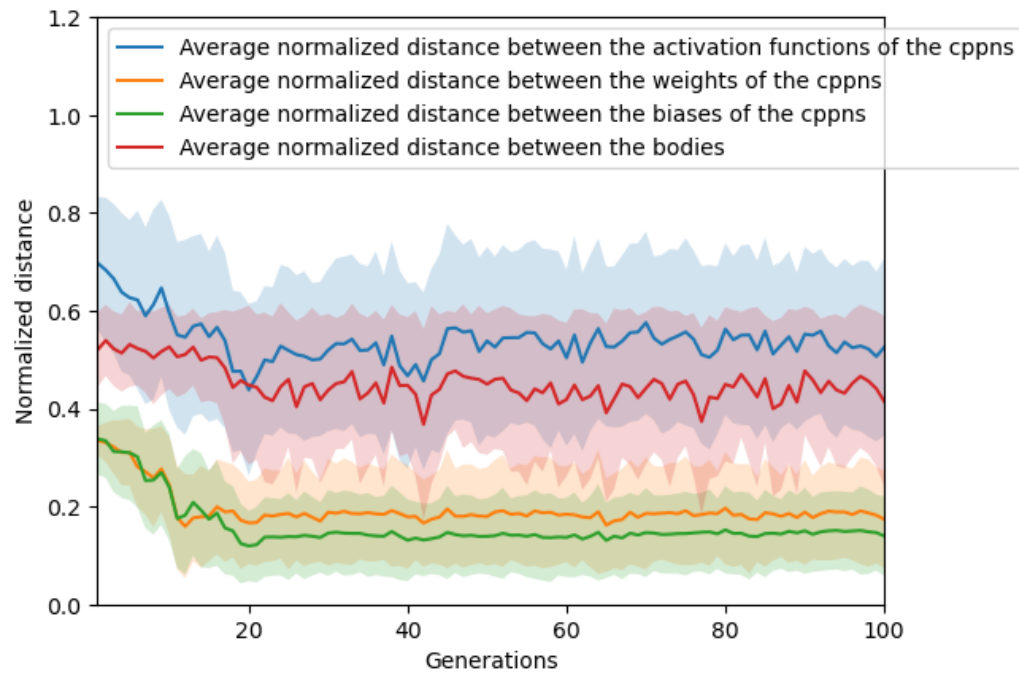
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  "max_weight" : 20,  
  "max_bias" : 20,  
  "response" : 1,  
  
  "range_weight" : 12,  
  "range_bias" : 12,  
  "sigma_weight" : 0.5,  
  "sigma_bias" : 0.3,  
  "threshold_weight" : 0.1,  
  "threshold_bias" : 0.1,  
  "threshold_function" : 0.05,  
  "threshold_dominance" : 0.1,  
  
  "number_of_dominances" : 5,  
}
```

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  "number_of_dominances" : 5,  
  
  "generations" : 100,  
  "population" : 100,  
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  "number_in_tournament" : 5,  
  "number_of_elites" : 2,  
  
  "shape_of_cppn" : [10, 4, 4, 2, 1],  
  
  "function_pool" : ["gaussian", "sin", "tanh"],  
  
  "n_steps" : 800,  
  "number_of_reported_individuals" : 40,  
  
  "cpus" : 7,  
  
  "env_name" : "Walker-v0"  
}
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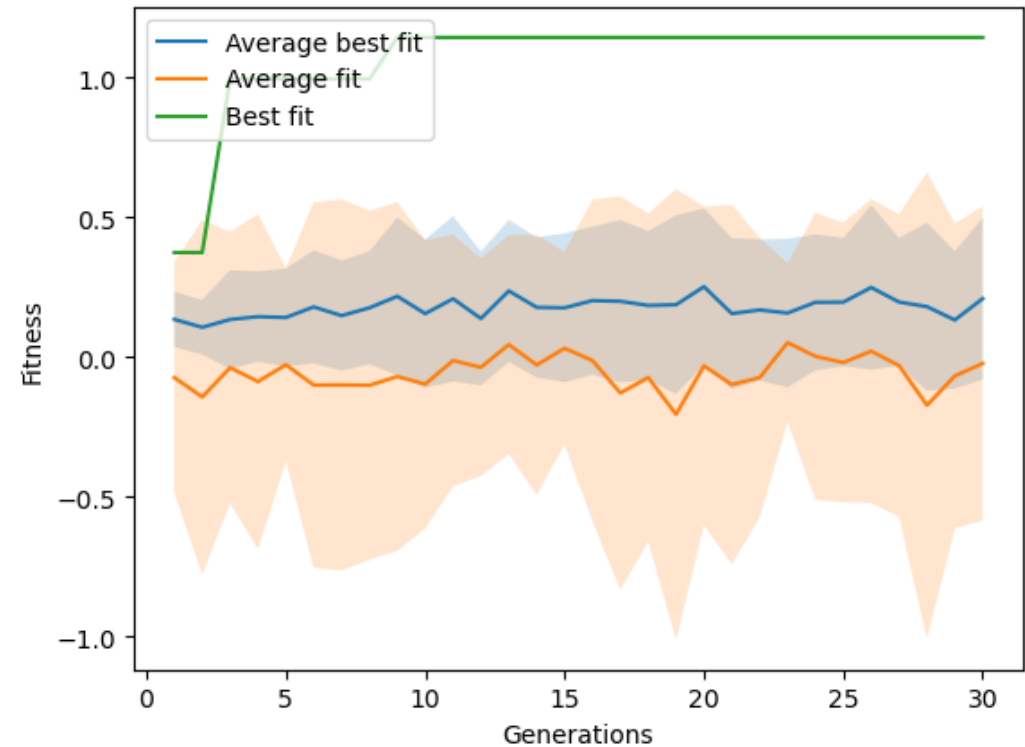
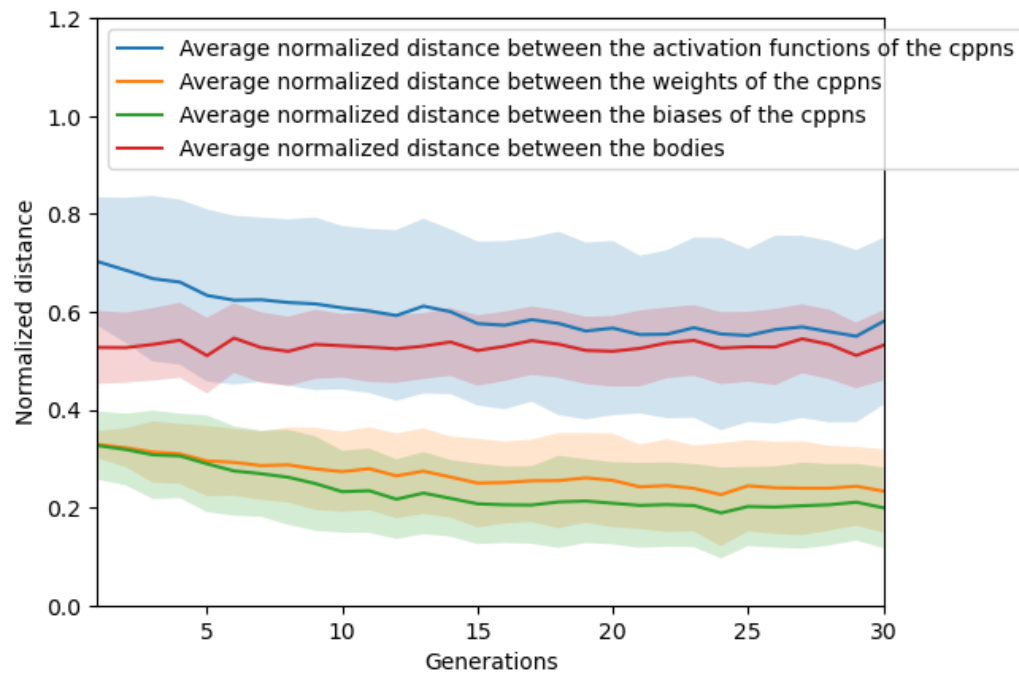
Initial Results (Stationary, Diploid, Walker-v0)



Initial Results (Stationary, Diploid, Climber-v0)



Initial Results (Stationary, Diploid, BridgeWalker-v0)



Videos

- Walker-v0 : <https://www.youtube.com/playlist?list=PL-fKXFhkBjTNbn6cMiTNqUD7EEFH63u7w>

- Climber-v0 : <https://www.youtube.com/playlist?list=PL-fKXFhkBjTPiS3WBmSGBMzGQ9fQO4ZoW>

- BridgeWalker-v0 : <https://www.youtube.com/playlist?list=PL-fKXFhkBjTM000iY-H2yw-BHk5M6r9cK>

Next

- Experiment with haploid individual and compare on stationary / changing environment
- Build and compare other forms of substrate at generation 0.

Thank you

ありがとうございます