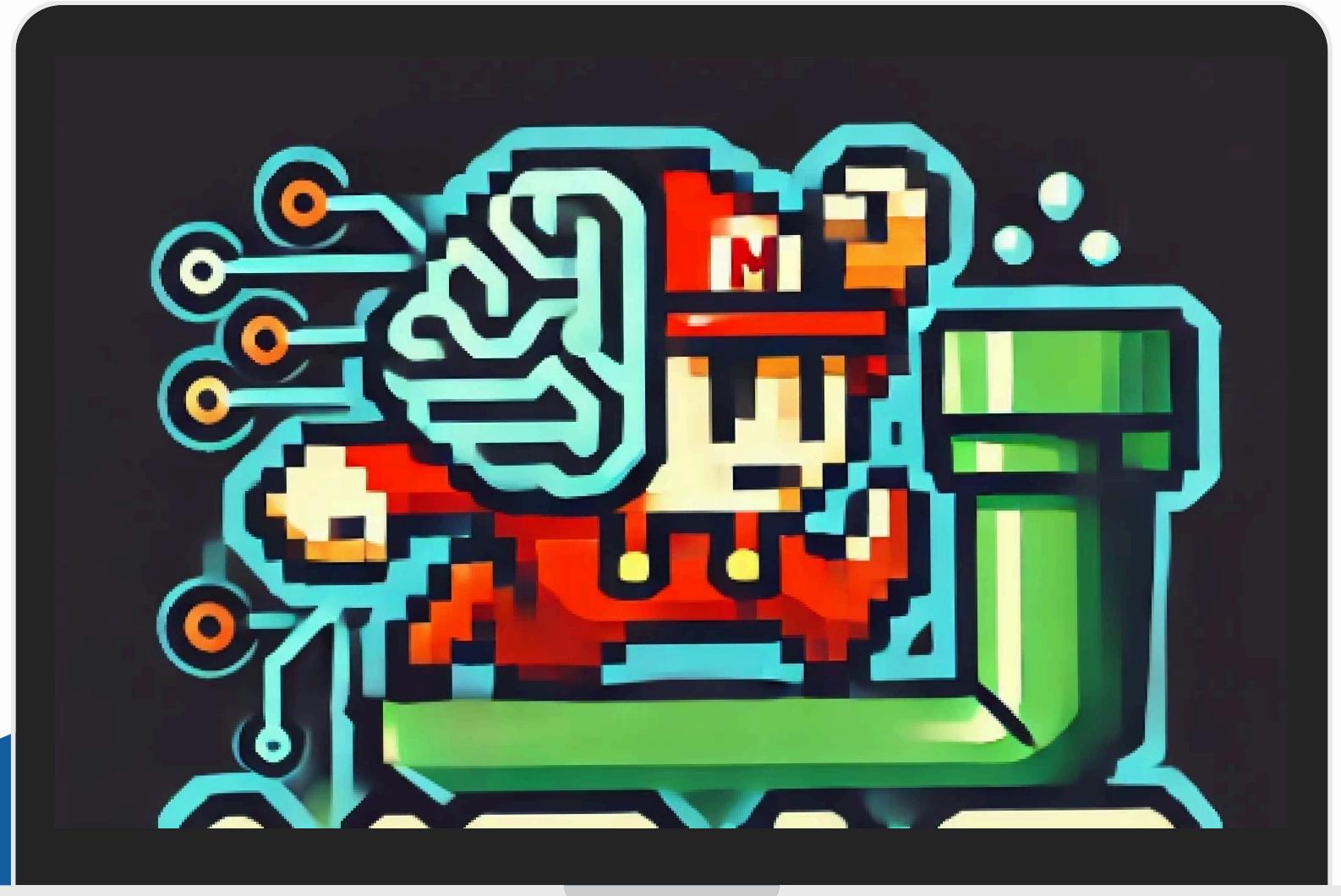
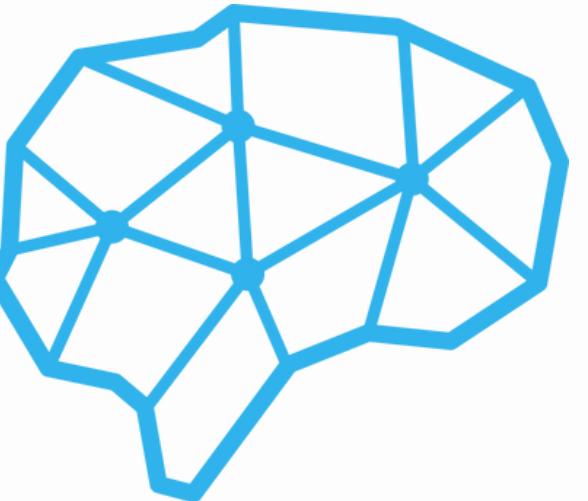


# Neat i Super Mario

By: Ludvig Øvrevik, Brage Kvamme, Christian Johnsen, Håkon Støren, Vetle Støren, Kristian Cornelius og Kacper Pawłowski



# AGENDA

01

Om Neat

02

Implementasjon

03

Resultater

04

Demo og  
Spørsmål

# MÅL

## Realistiske mål



- Lage en evolusjonær algoritme fra bunn
- Få Mario til å klare første bane

- Få verdensrekord i Super Mario
- Kunne spille alle typer spill, med litt trening

## Ambisiøse mål



# HVA ER NEAT?

## NeuroEvolution of Augmenting Topologies

- Etterligner evolusjonsprosessen
- Survival of the fittest
- Muterer og utvikler seg over flere generasjoner
- Hensikt: Løsning med minimalt nettverk

Fig. 1. A genotype to phenotype mapping example. The third gene is disabled, so the connection that it specifies (between nodes 2 and 5) is not expressed in the phenotype.

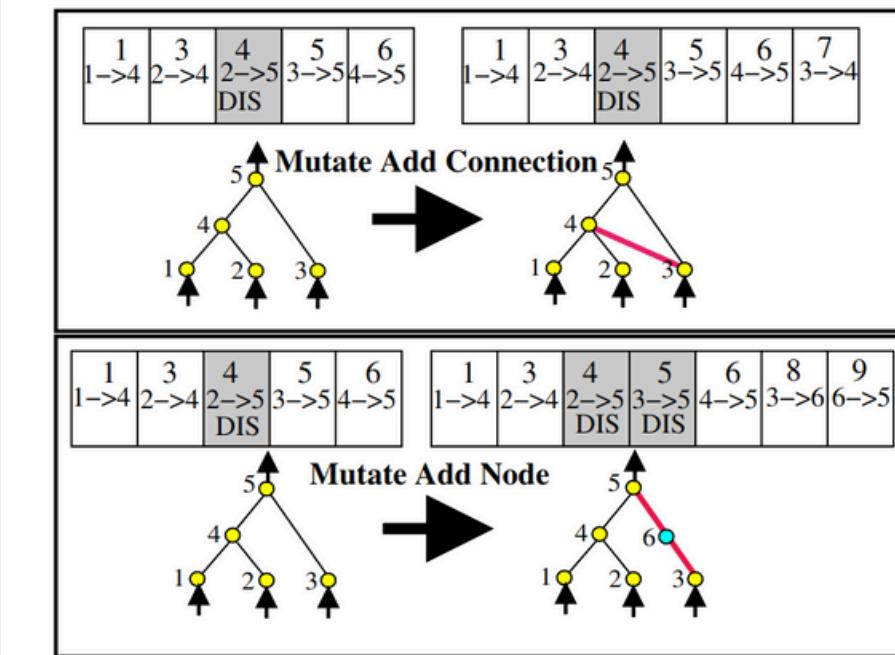
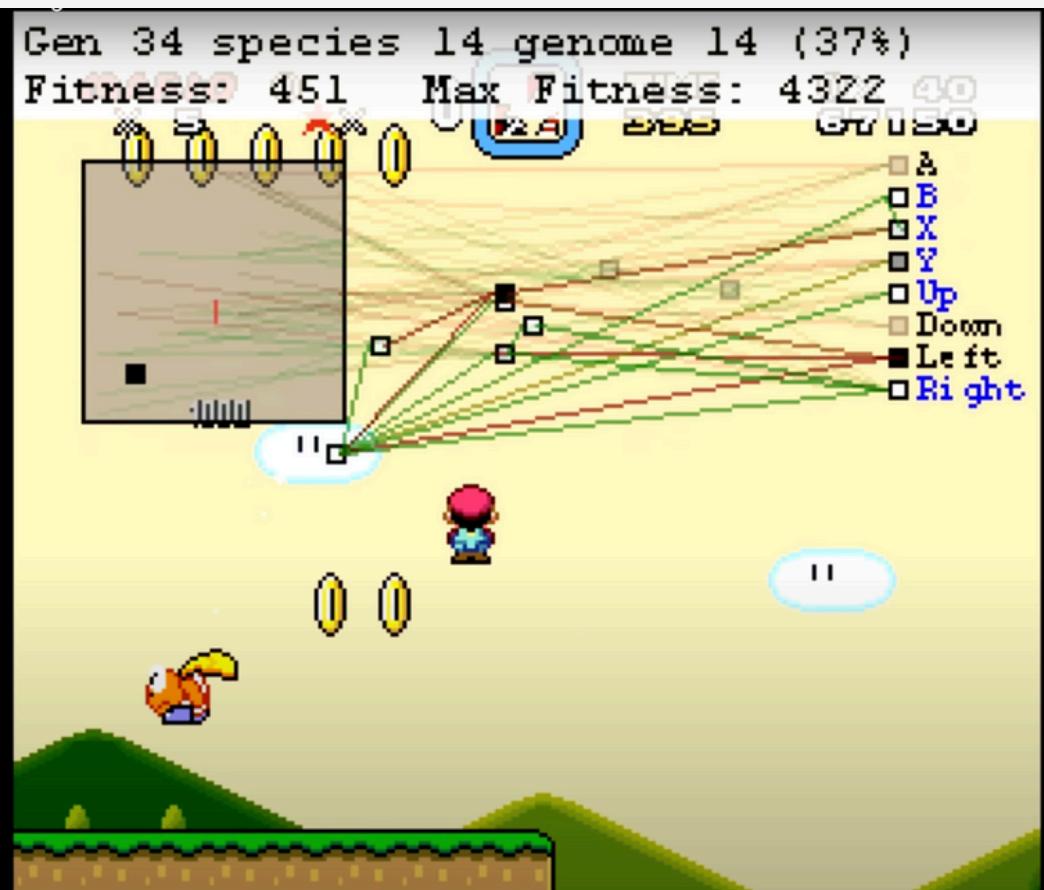


Fig. 2. The two types of structural mutation in NEAT. Both types, adding a connection and adding a node, are illustrated with the genes above their phenotypes. The top number in each genome is the *innovation number* of that gene. These numbers identify the original historical ancestor of each gene, making it possible to find matching genes during crossover. New genes are assigned new increasingly higher numbers.

# Hva vi tok inspirasjon fra



## Efficient Evolution of Neural Network Topologies

Kenneth O. Stanley and Risto Miikkulainen  
Department of Computer Sciences  
The University of Texas at Austin  
Austin, TX 78712  
[kstanley, risto@cs.utexas.edu](mailto:kstanley, risto@cs.utexas.edu)

### Abstract

Neuroevolution, i.e. evolving artificial neural networks with genetic algorithms, has been highly effective in reinforcement learning tasks, particularly those with hidden state information. An important question neuroevolution is how to gain an advantage from evolving neural network topologies along with weights. We present a method, NeuroEvolution of Augmenting Topologies (NEAT) that outperforms the best fixed-topology methods on a challenging benchmark reinforcement learning task. We claim that the increased efficiency is due to (1) employing a principled method of crossover of different topologies, (2) protecting structural innovation using speciation, and (3) incrementally growing from minimal structure. We test this claim through a series of ablation studies that demonstrate that each component is necessary to the system as a whole and to each other. What results is significantly faster learning. NEAT is also an important contribution to GAs because it shows how it is possible for evolution to both optimize and complexify solutions simultaneously, making it possible to evolve increasingly complex solutions over time, thereby strengthening the analogy with biological evolution.

### I. INTRODUCTION

Neuroevolution (NE), the artificial evolution of neural networks using genetic algorithms, has shown great promise in reinforcement learning tasks. NE outperforms standard reinforcement learning methods in many benchmark tasks [6, 10, 11]. Neural networks are a good class of decision making systems to evolve because they are capable of representing solutions to many different kinds of problems, and the mapping from genotype to phenotype is generally efficient. NE is particularly well suited to reinforcement learning tasks because NE does not require supervision.

A major question in NE is how to gain an advantage from evolving topology in addition to connection weights. On one hand, evolving topology might overcomplicate the search. On the other, it can also save time by finding the right number of hidden neurons for a particular problem automatically [7].

A previous study showed that fixed-topology NE can outperform a topology-evolving system on the benchmark double pole balancing task [6]. This finding is important because pole balancing has been a benchmark task in NE and reinforcement learning for over 30 years [1, 6, 7, 9], and double pole balancing is challenging to even the best of modern

methods. Doing well at this important benchmark suggests that a method will do well in other tasks as well. Whether Topology and Weight Evolving Artificial Neural Networks (TWEANNs) can enhance the performance of NE remains an open question.

In this article, we aim to show that evolving topology can indeed increase performance. We present a new TWEANN, NeuroEvolution of Augmenting Topologies (NEAT), that significantly outperforms the fixed-topology NE method that currently takes the fewest evaluations on the double pole balancing task. We identify three major challenges for TWEANNs and present solutions to each of them: (1) Is there a genetic representation that allows disparate topologies to crossover in a meaningful way? Our solution is to use historical markings to line up genes with the same origin. (2) How can topological innovation that needs a few generations to optimize be protected so that it does not disappear from the population prematurely? Our solution is to separate each innovation into a different species. (3) How can topologies be minimized *throughout evolution* without the need for a specially contrived fitness function that measures complexity? Our solution is to start from a minimal structure and grow only when necessary. This paper establishes that each of our solutions is necessary by showing that NE performance significantly declines with the ablation of any of the major solution components. Working together in NEAT these components constitute a promising new approach to difficult reinforcement learning tasks.

We begin by describing the NEAT method, including results showing that NEAT is significantly faster than other NE methods on the hardest pole balancing benchmark. We then present ablation studies designed to explain NEAT's performance in terms of its components.

### II. NEUROEVOLUTION OF AUGMENTING TOPOLOGIES (NEAT)

#### A. Genetic Encoding

NEAT is designed specifically to address the three challenges raised in the introduction. Each genome includes a list of *connection genes*, each of which refers to two *node genes* being connected (figure 1). Each connection gene specifies the in-node, the out-node, the weight of the connection, whether

To appear in *Proceedings of the 2002 Congress on Evolutionary Computation (CEC '02)*. Piscataway, NJ: IEEE

Sethbling

Originalt paper fra 2002 om NEAT

# NEAT Paper

---

DÅRLIG!

- Var ikke lett å forstå hvordan de selv hadde implementert algoritmen
- Måtte gjøre mange antagelser og finne egne løsninger på problemer

# Implementasjon

- Input: Piksel fra spillet - 10x20 grayscale | 20x40 RGB
- Output: Action (Jump, move left, move right...)
- Ikke brukt pytorch eller tensorflow.
- Alt fra genomes, til noder og connections er klasser

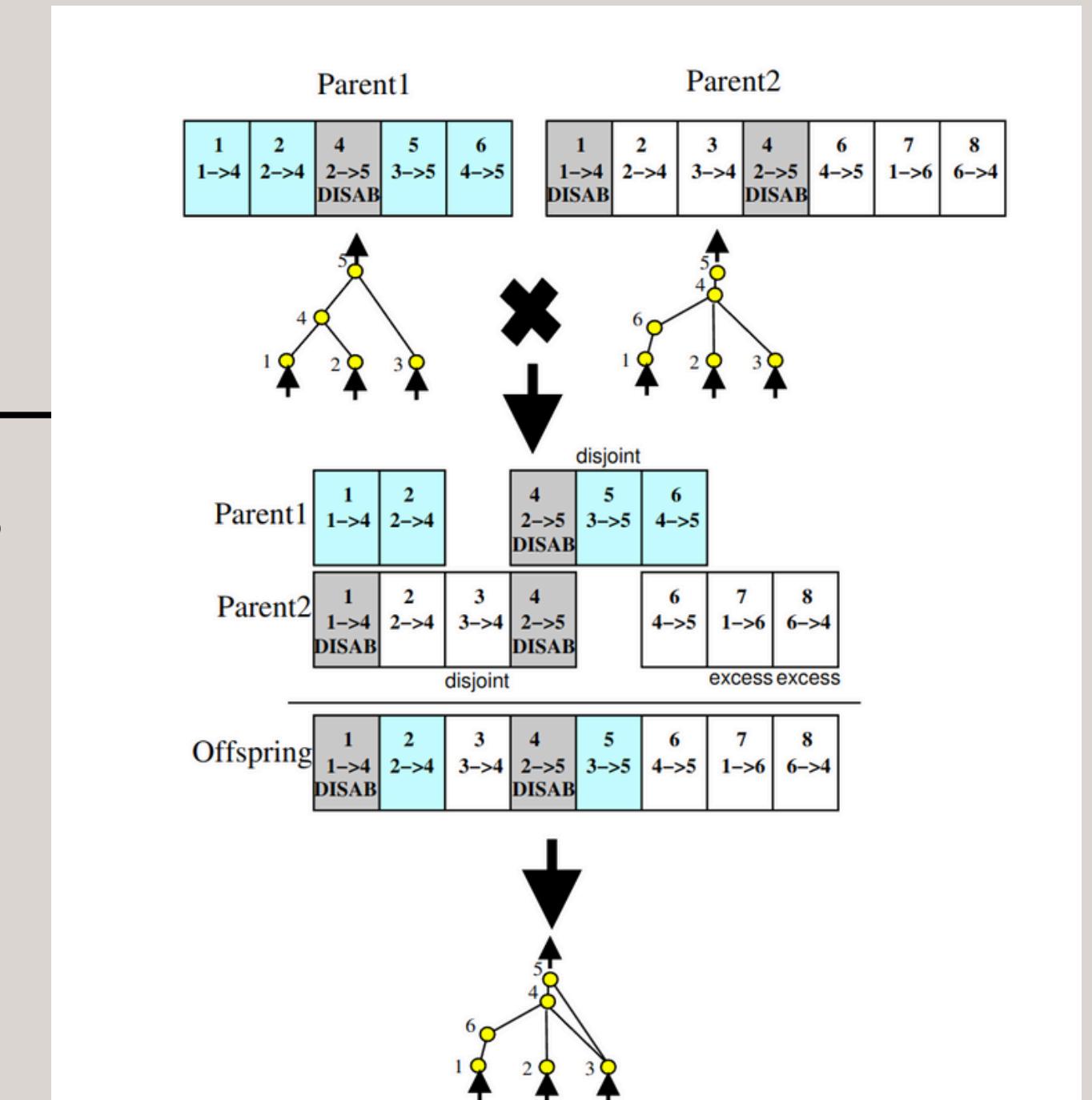
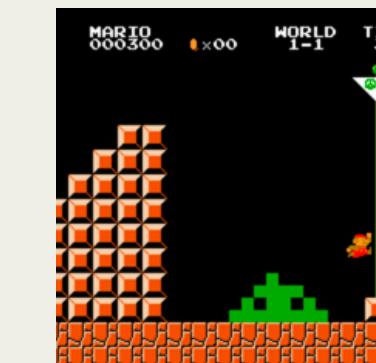
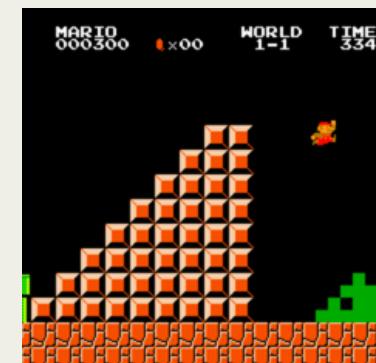
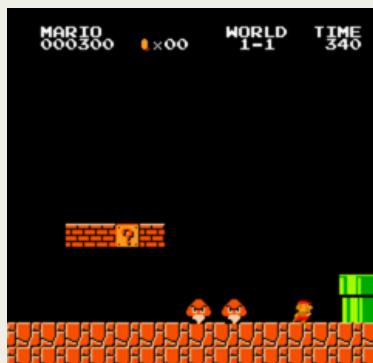
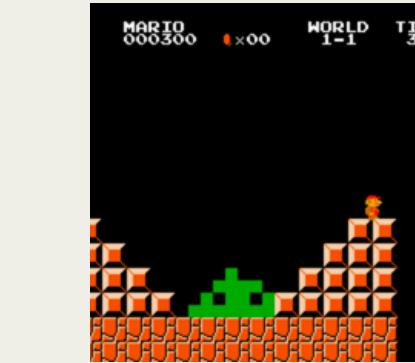
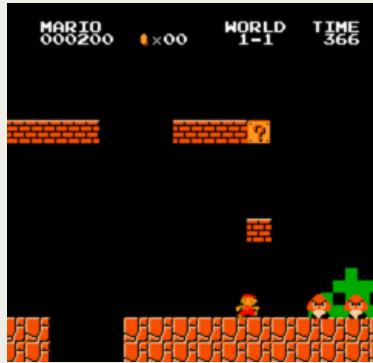
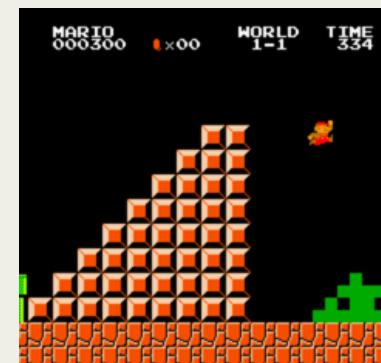
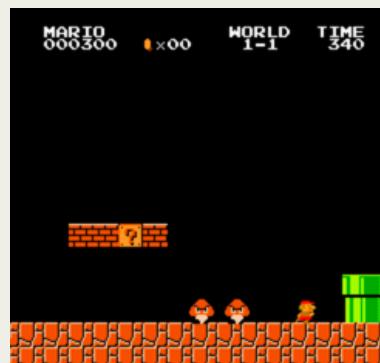
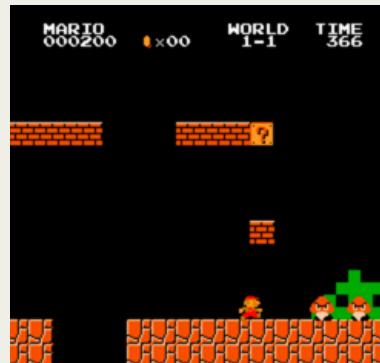


Fig. 3. Matching up genomes for different network topologies using innovation numbers. Although Parent 1 and Parent 2 look different, their innovation numbers (shown at the top of each gene) tell us which genes match up with which without the need for topological analysis.

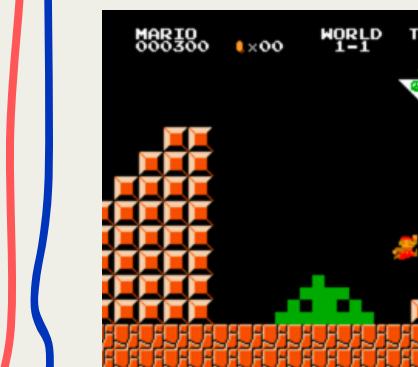
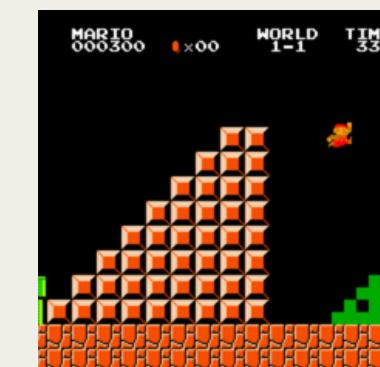
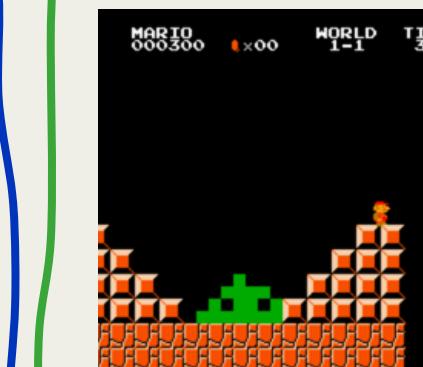
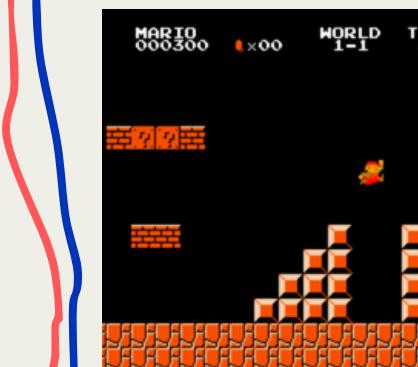
# For hver generasjon: Starter med en populasjon



# Spill mario og få fitness



# Del inn i arter

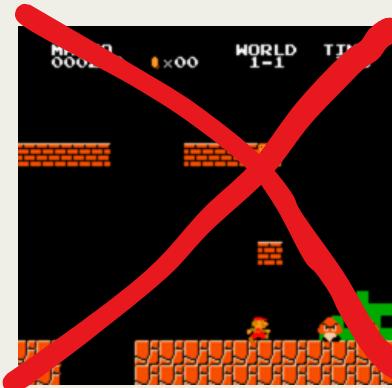


Art nummer 1

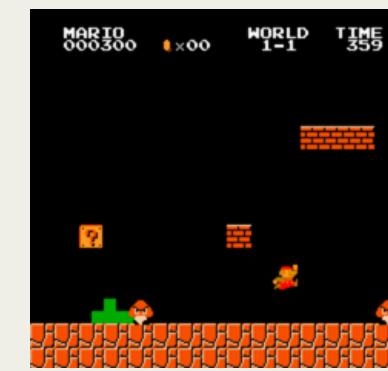
Art nummer 2

Art nummer 3

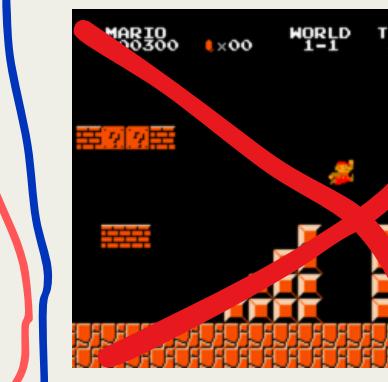
# DREP DE DÅRLIGSTE



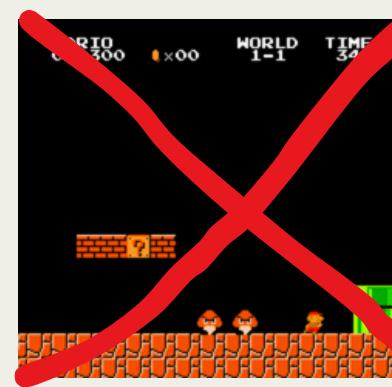
Fitness: 69



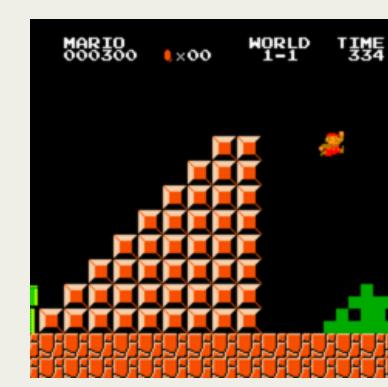
Fitness: 420



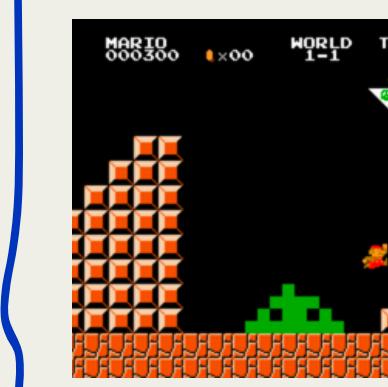
Fitness: 3



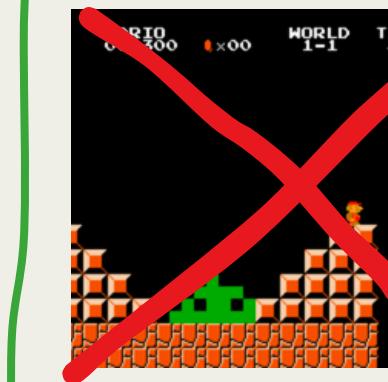
Fitness: 74



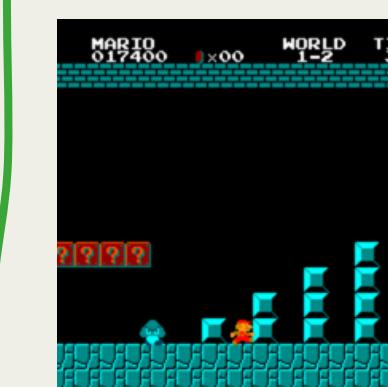
Fitness: 79



Fitness: 300



Fitness: 12



Fitness: 500



Fitness: 42



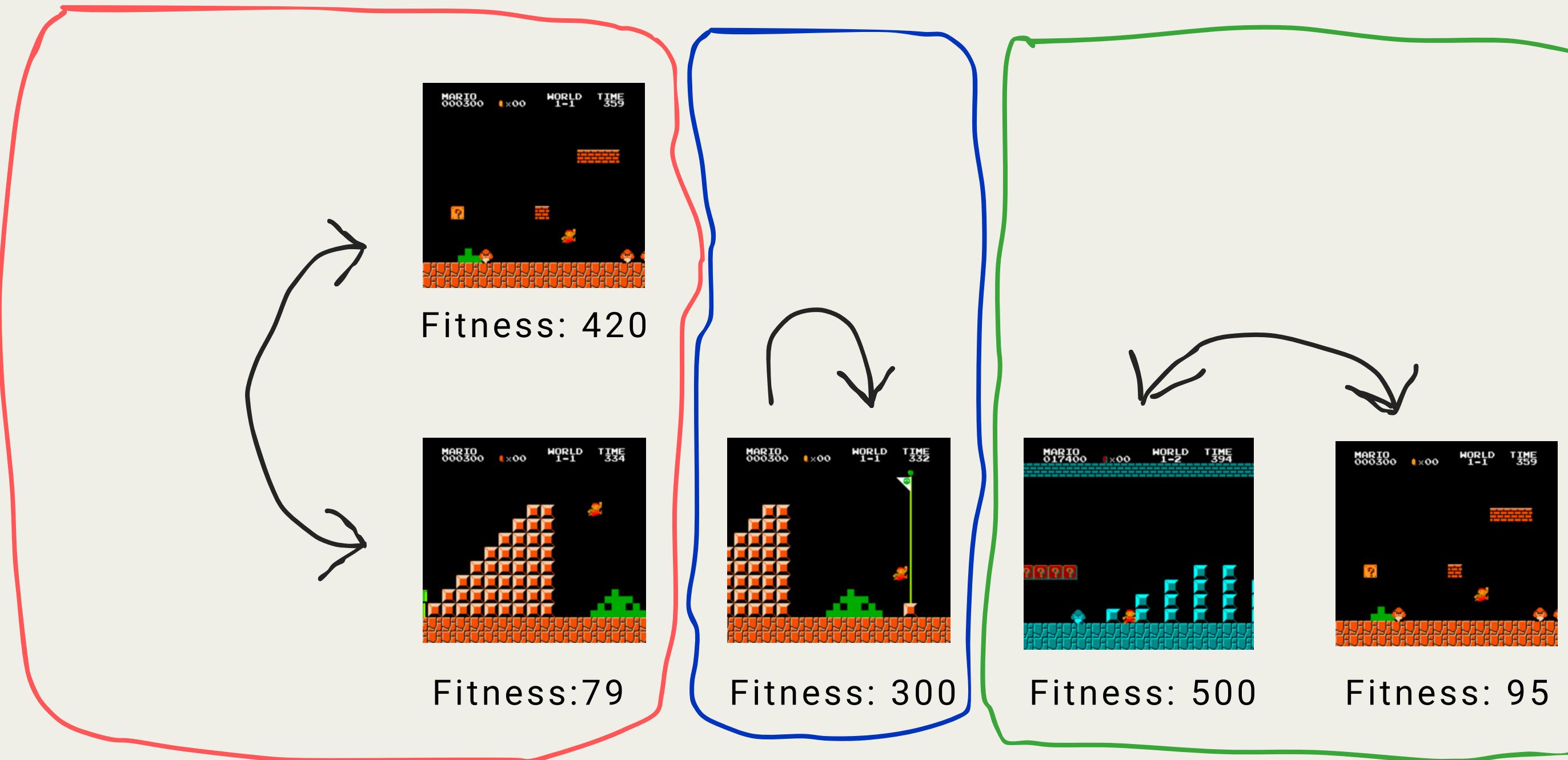
Fitness: 95

Art nummer 1

Art nummer 2

Art nummer 3

# De som overlever parrer seg

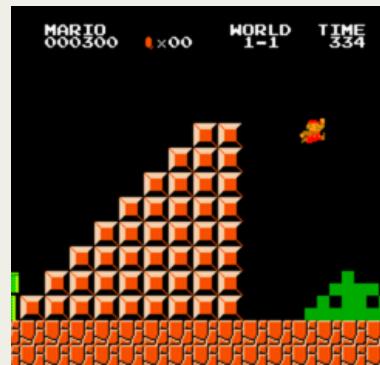
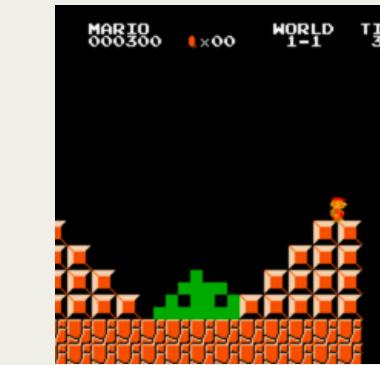


Art nummer 1

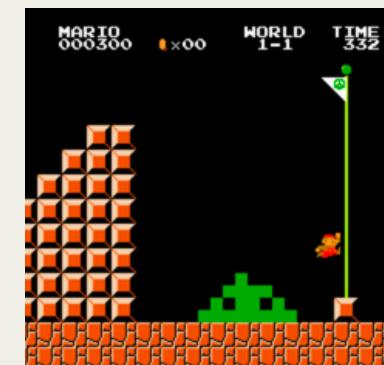
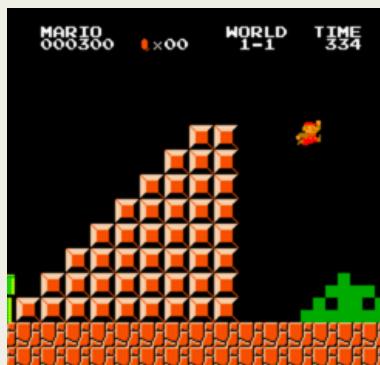
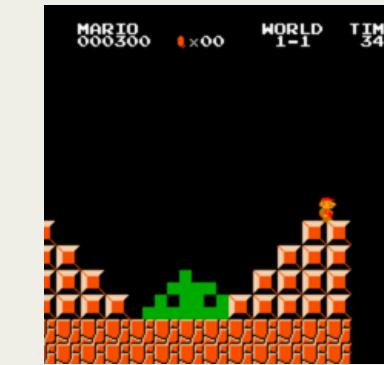
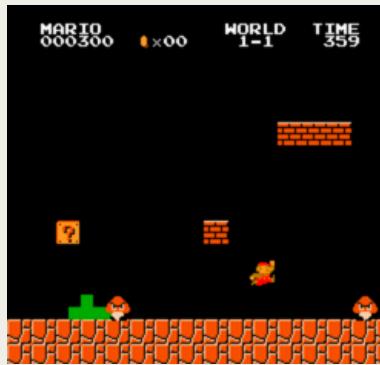
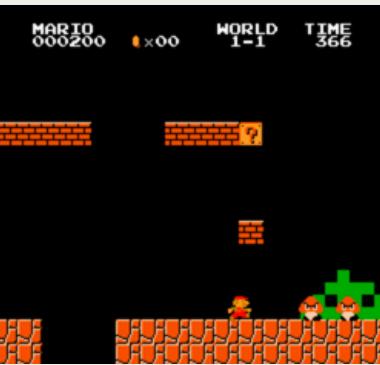
Art nummer 2

Art nummer 3

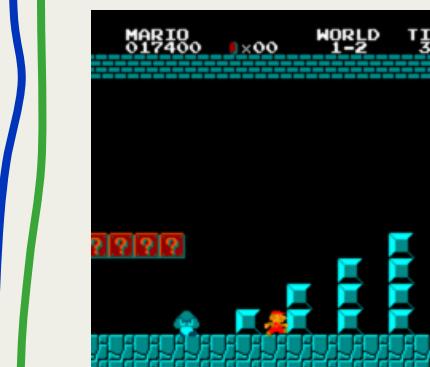
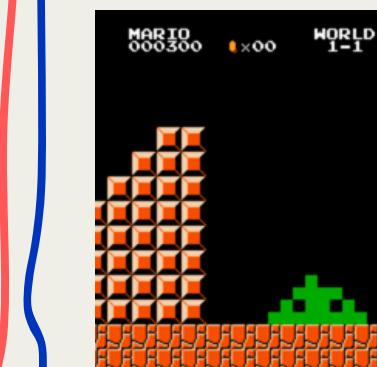
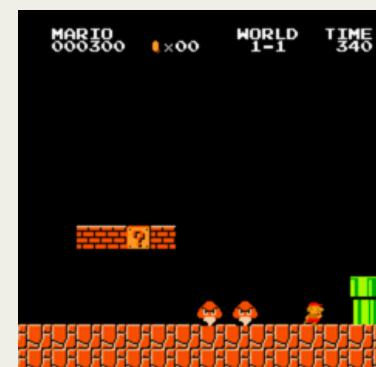
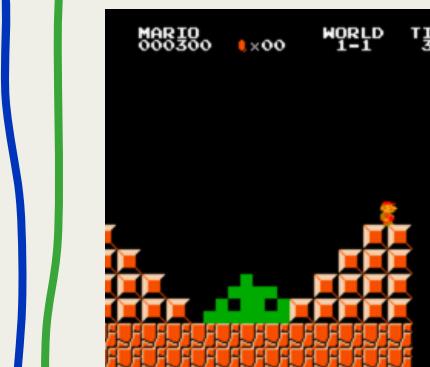
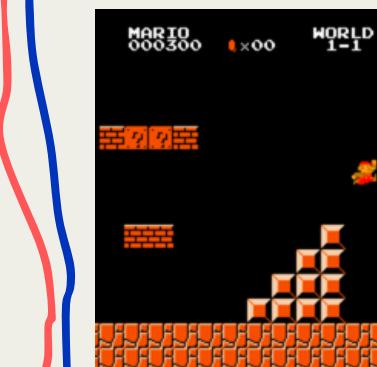
# Ny generasjon blir laget



# Muter den nye generasjonen



# Del inn i nye arter



Art nummer 1

Art nummer 2

Art nummer 3

## **Formel for å regne genetisk distanse mellan to nettverk**

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

**Formel for å regne ut hvor mange individer hver  
art skal ha i neste generasjon.**

$$N'_j = \frac{\sum_{i=1}^{N_j} f_{ij}}{\bar{f}},$$

# Filosofiske og etiske avveiinger

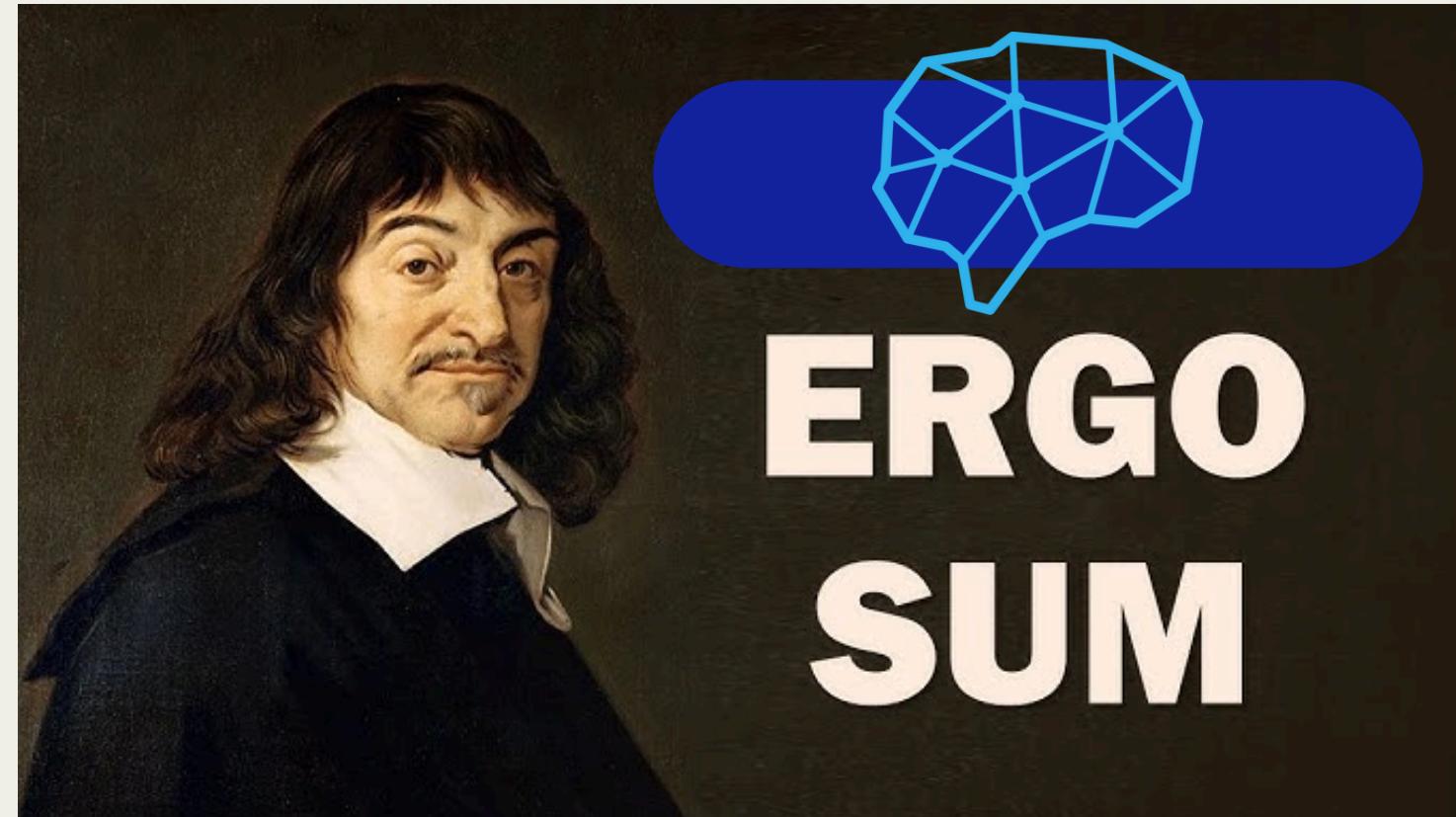
Sosialdarwinisme

Dele inn i arter og segregere dem

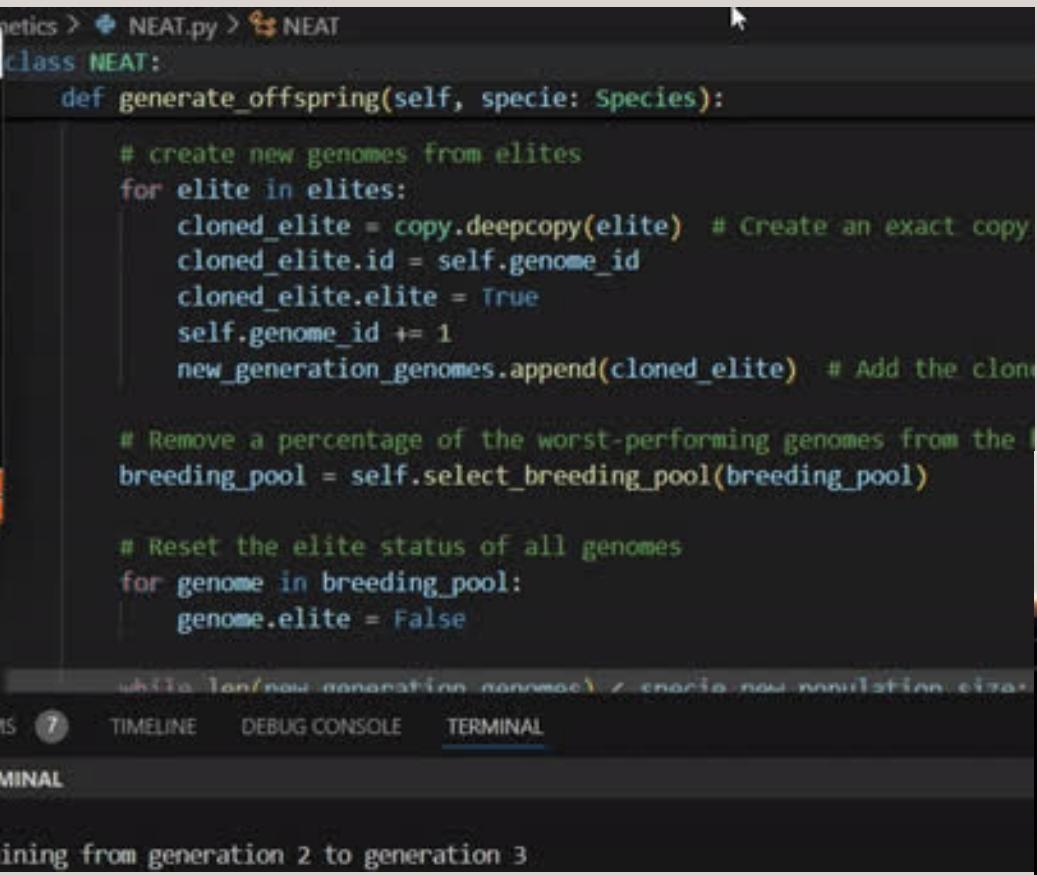
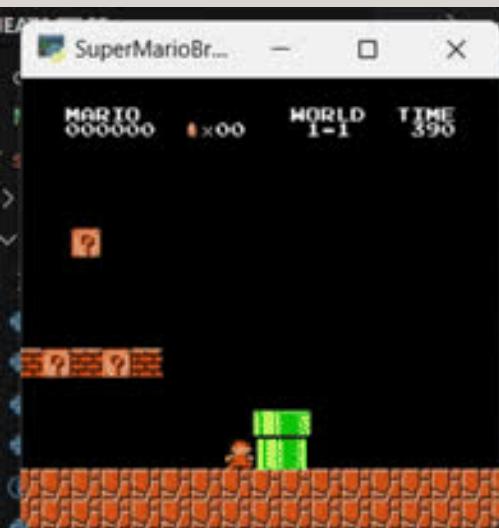
Drep svakeste barn

Dominant og undertrykket  
nettverk

Genetisk determinisme



# Så var det training



```
SuperMarioBros - □ X
genetics > NEAT.py > NEAT
class NEAT:
    def generate_offspring(self, specie: Species):
        # Create new genomes from elites
        for elite in elites:
            cloned_elite = copy.deepcopy(elite) # Create an exact copy
            cloned_elite.id = self.genome_id
            cloned_elite.elite = True
            self.genome_id += 1
            new_generation_genomes.append(cloned_elite) # Add the clone

        # Remove a percentage of the worst-performing genomes from the breeding pool
        breeding_pool = self.select_breeding_pool(breeding_pool)

        # Reset the elite status of all genomes
        for genome in breeding_pool:
            genome.elite = False

    while len(new_generation_genomes) < specie.new_population_size:
        # Create new genomes from elites
        for elite in elites:
            cloned_elite = copy.deepcopy(elite) # Create an exact copy
            cloned_elite.id = self.genome_id
            cloned_elite.elite = True
            self.genome_id += 1
            new_generation_genomes.append(cloned_elite) # Add the clone

        # Remove a percentage of the worst-performing genomes from the breeding pool
        breeding_pool = self.select_breeding_pool(breeding_pool)

        # Reset the elite status of all genomes
        for genome in breeding_pool:
            genome.elite = False

    return new_generation_genomes
```

PROBLEMS 7 TIMELINE DEBUG CONSOLE TERMINAL

Training from generation 2 to generation 3





20879714	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879712	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879710	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879708	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879706	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879704	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879702	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879700	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879698	GPUQ	cost_eff	jcrogers	PD	0:00	1 (Priority)
20879756	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879754	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879752	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879750	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879748	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879746	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879864	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879862	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879860	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879858	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879856	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879854	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879852	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879850	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879848	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879846	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879844	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879842	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879816	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879814	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879812	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879810	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879808	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879806	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879804	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879802	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879800	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879798	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879796	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879794	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879768	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879766	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879764	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879762	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879760	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879758	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879912	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879910	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879908	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879906	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879904	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879902	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879900	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879898	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879896	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879894	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)
20879892	GPUQ	carbon_e	jcrogers	PD	0:00	1 (Priority)

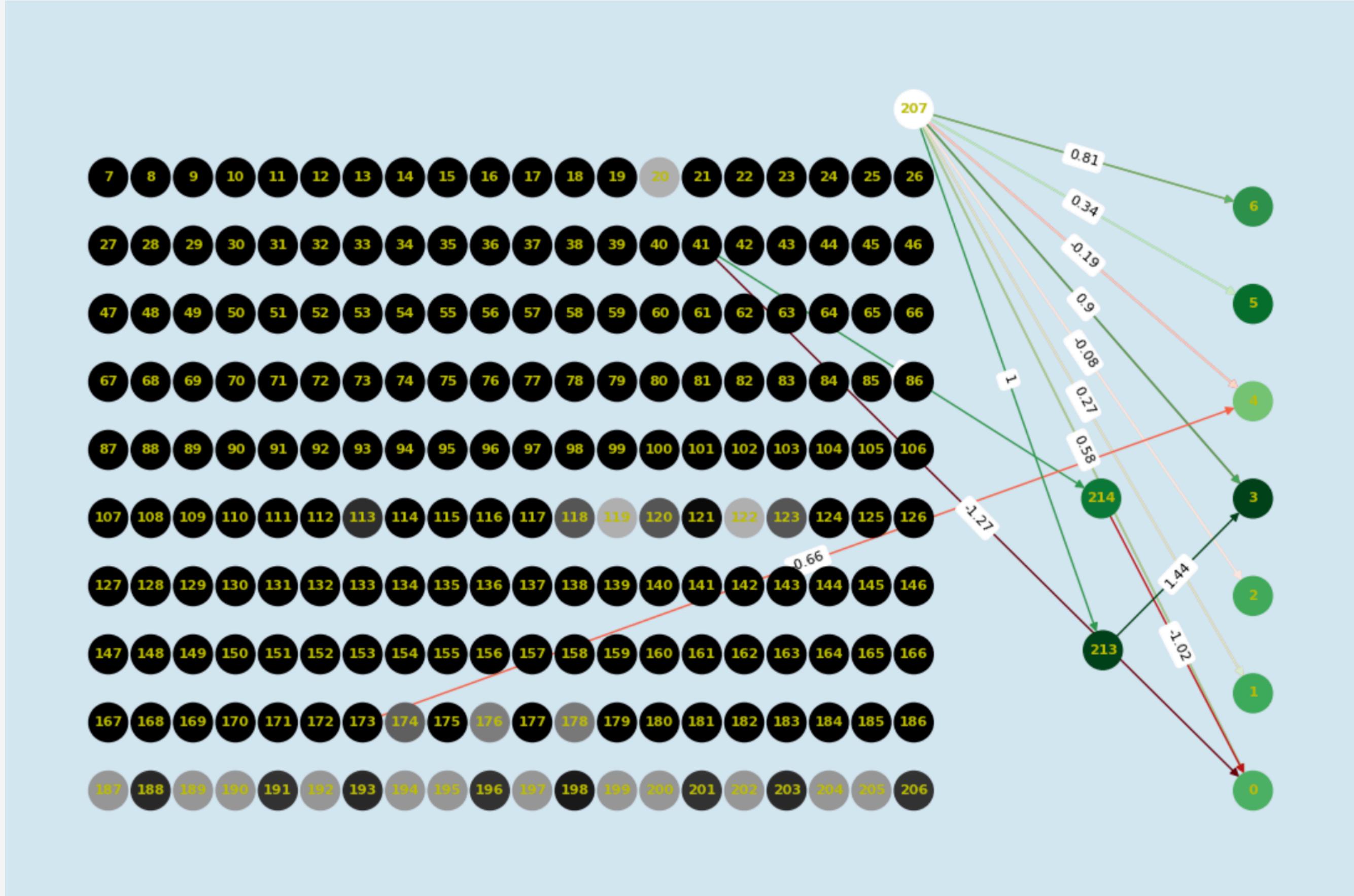


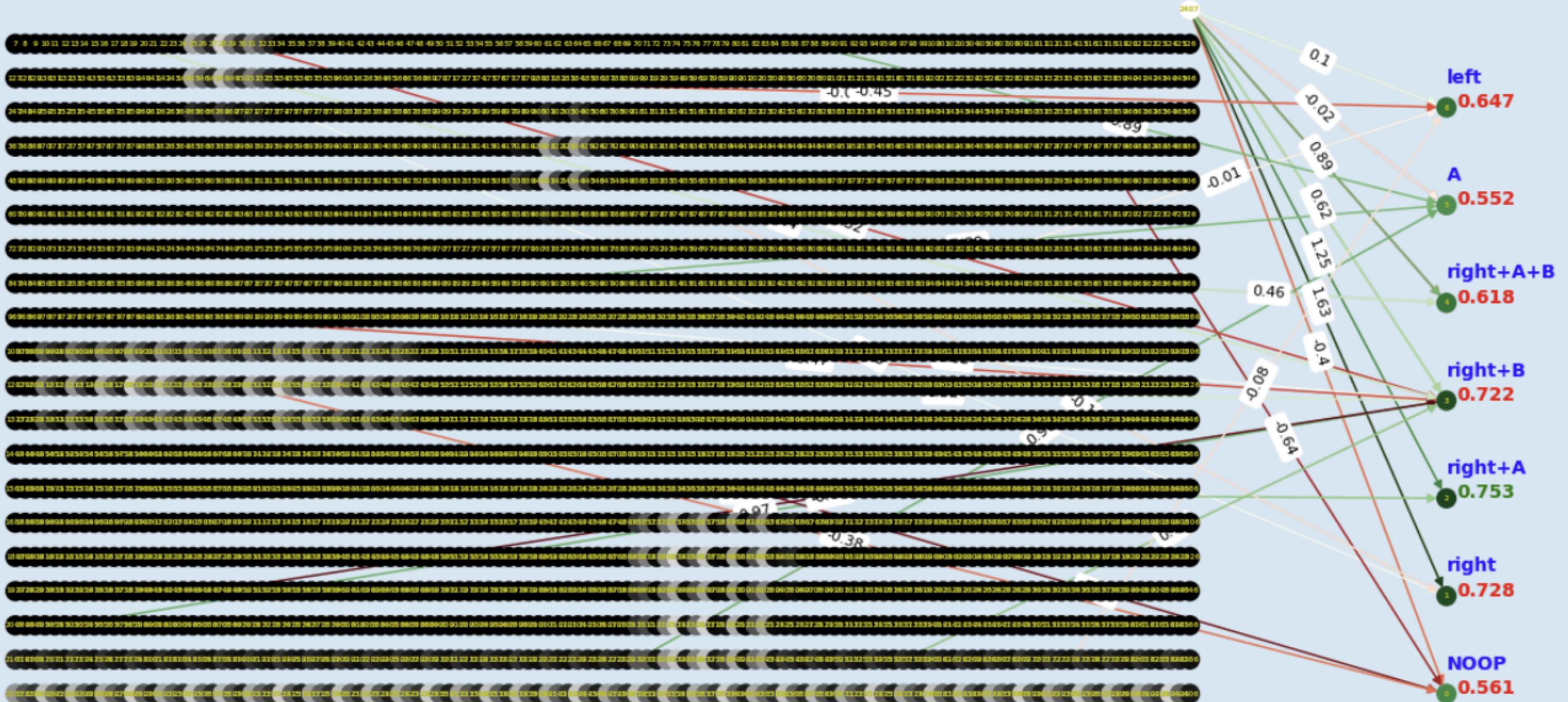
**BRUKE IDUN**

mgflip.com

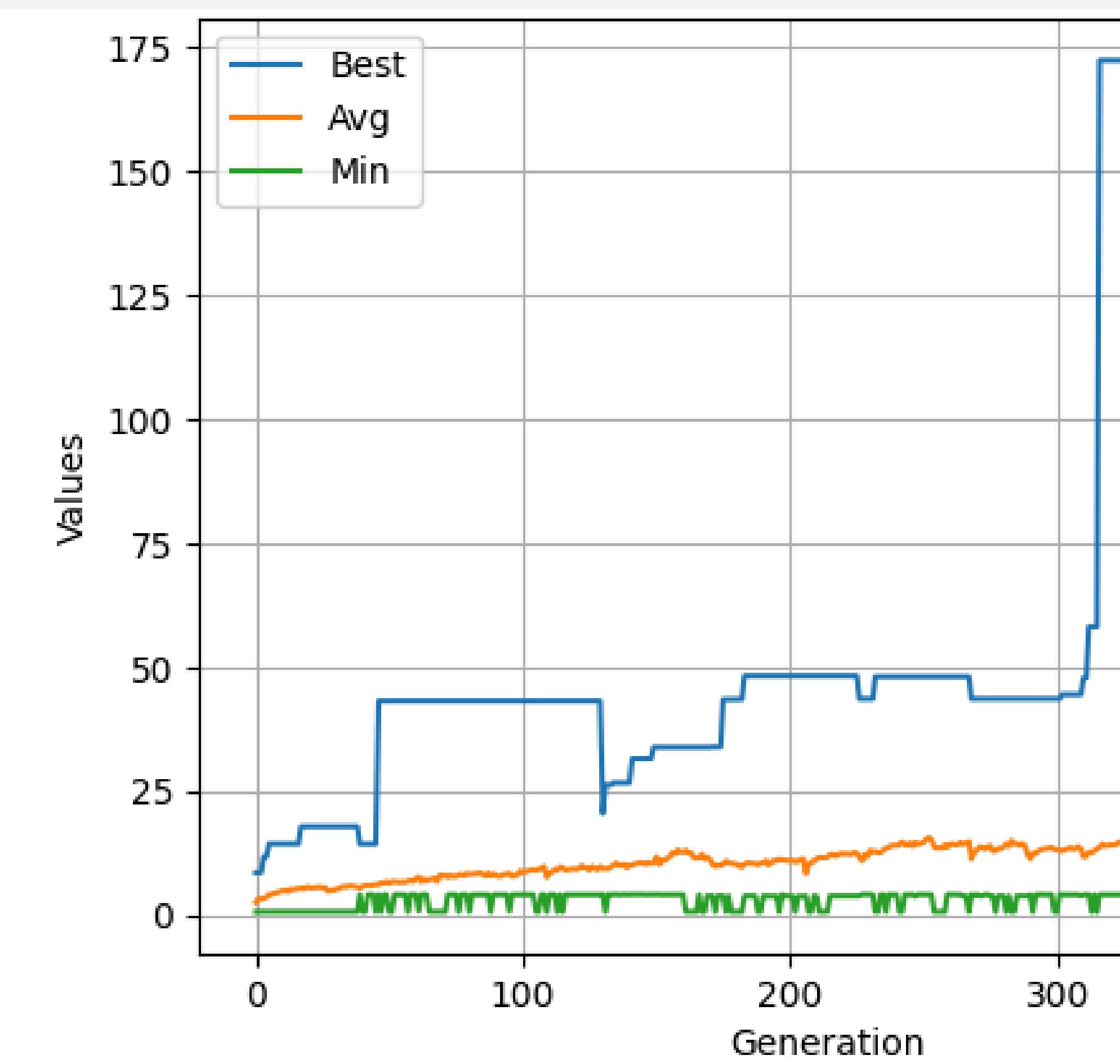
# Hva nettverket ser:



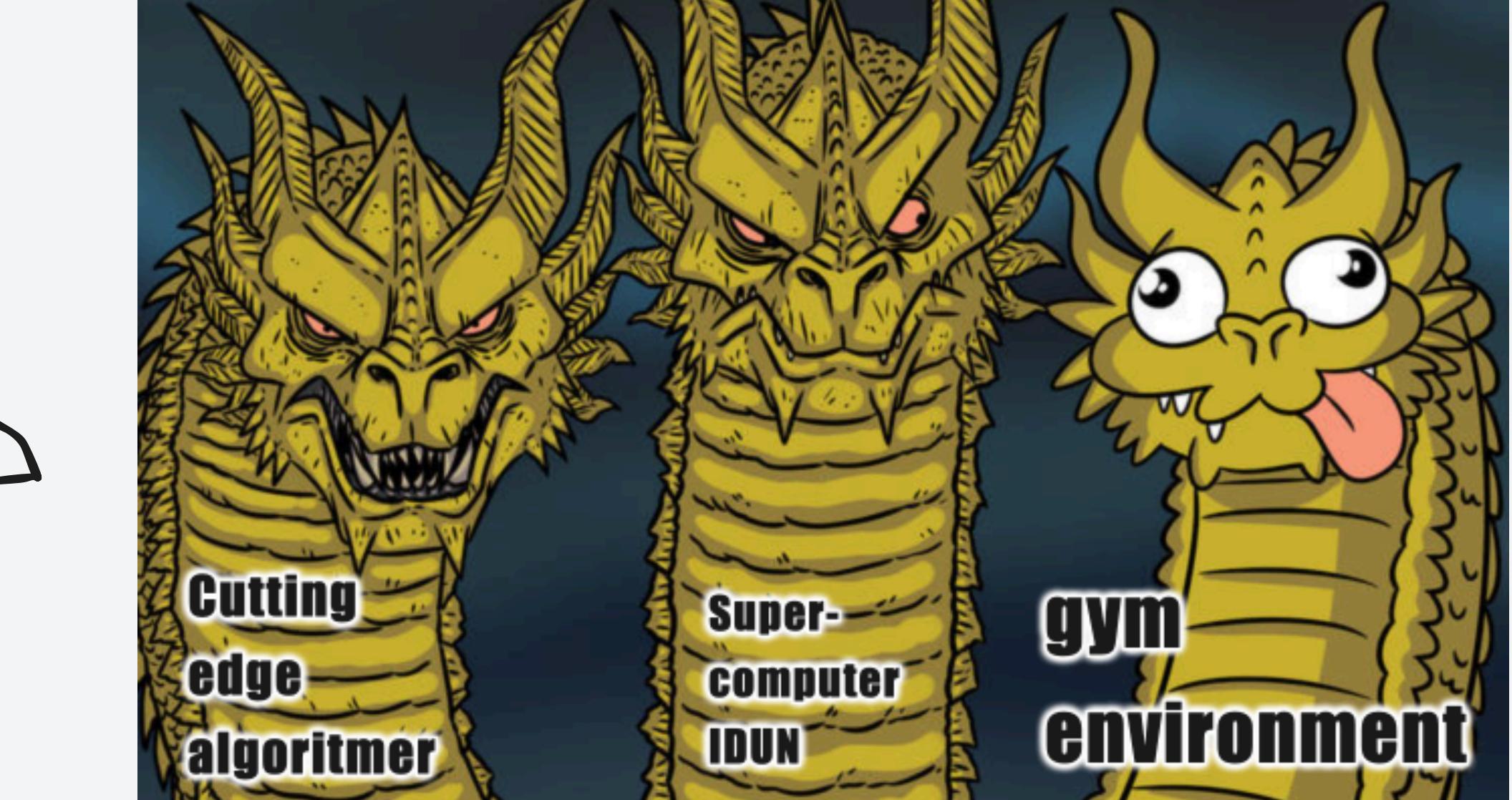




# Fitness



**Problemer og ting vi kunne  
gjort annerledes**





# Carlenius Consulting

Org nr 931 467 220

Telefon 954 83 129 Adresse Huldreveien 19, 1388 Borgen

Sammenlign

Overvåk

Finn lignende

Del profil

Oversikt

Regnskap

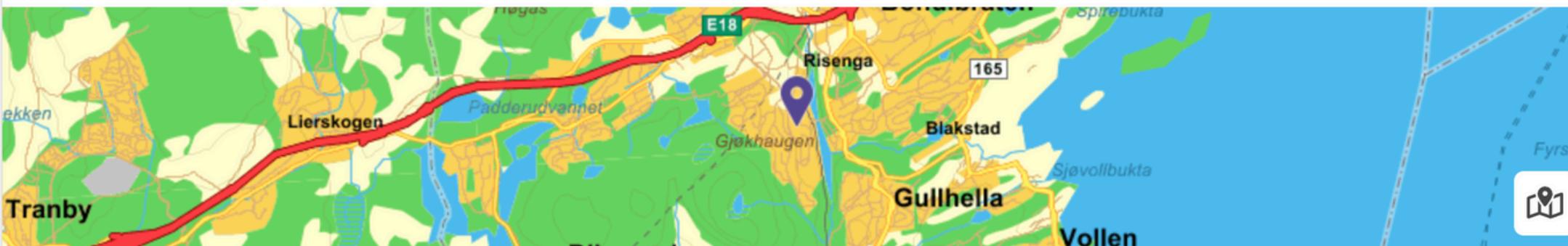
Nøkkeltall

Roller og Eiere

Organisasjon

Kunngjøringer

## Kontaktopplysninger



Telefon

954 83 129

Adresse

Huldreveien 19, 1388 Borgen

Postadresse

Huldreveien 19, 1388 Borgen



## Offisiell foretaksinformasjon

Juridisk navn

CARLENIUS CONSULTING

Adresse

c/o Signy Gautefall Huldreveien 19, 1388  
Borgen

Org nr

931 467 220

Postadresse

c/o Signy Gautefall Huldreveien 19, 1388  
Borgen

Registreringsdato

30.05.2023

Registrert i

MVA

Selskapsform

Enkeltpersonforetak

Foretaksregisteret

NACE-bransje

62.010 Programmeringstjenester

NAV aa-registeret

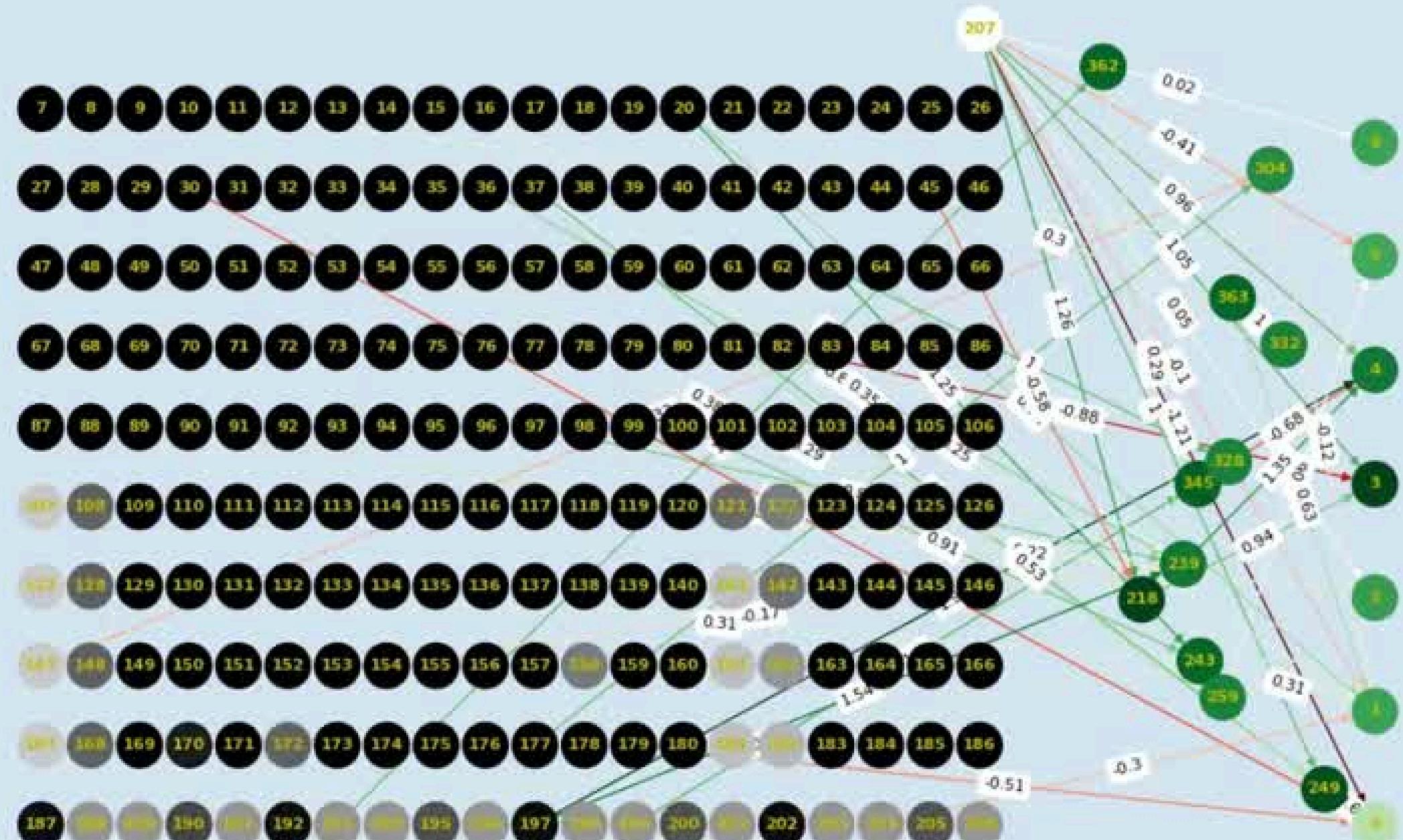
Innehaver

Kristian Gautefall Carlenius

Frivillighetsreg

Kilde: Brønnøysundregistrene

# EXAMPLE NETWORK



# Demo

# Spørsmål?

## CogitoNTNU/ NEATactics



NEAT is neat! Neuroevolution of augmenting topologies is a genetic algorithm.

8 Contributors 1 Issue 8 Stars 1 Fork



**CogitoNTNU/NEATactics: NEAT is neat! Neuroevolution of augmenting topologies is a genetic algorithm.**

NEAT is neat! Neuroevolution of augmenting topologies is a genetic algorithm. - CogitoNTNU/NEATactics

 GitHub