# **Ecosystem project**

# **Computational Model**

by

# Inbar koursh

This project is submitted under the scope of 1 unit and as partial fulfillment of the requirements for obtaining a grade in computational science

This work was performed under the guidance of Shlomo Rozenfeld

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### **Abstract**

Studying natural biological systems using a simulated ecosystem and reinforcement learning.

This project attempts to mimic similar simulations of ecosystems to further understand the factors that lead to successful species. This is done by studying the correlation between the recorded variables at any given time.

The model was successfully established. It demonstrated a significant correlation between all the 13 variables that were examined. This demonstrates the long thought idea that all an ecosystem is a highly complex system in which every change impacts the entire system.

# **Background**

It is expected that the reader would hold a fair knowledge of python and understand the concept behind evolutionary-based neural networks.

Neuroevolution is a machine learning technique that applies evolutionary algorithms to construct artificial neural networks, taking inspiration from the evolution of biological nervous systems in nature. Compared to other neural network learning methods, neuroevolution is highly general; it allows learning without explicit targets, with only sparse feedback, and with arbitrary neural models and network structures. Neuroevolution is an effective approach to solving reinforcement learning problems and is most commonly applied in evolutionary robotics and artificial life. <sup>1</sup>

<sup>&</sup>lt;sup>1</sup> Joel Lehman and Risto Miikkulainen (2013) Neuroevolution. Scholarpedia, 8(6):30977., revision #137053

For example, it is next to impossible to say if any one chess move at the start of the game is a good move. However, in neuroevolution, you can train the network by giving it a score on the whole game and not just on 1 move (unlike traditional approaches).

Python is an interpreted, object-oriented, high-level programming language with dynamic semantics. It's high-level built-in data structures, combined with dynamic typing and dynamic binding, make it very attractive for Rapid Application Development, as well as for use as a scripting or glue language to connect existing components. Python is open-sourced and is available for free on all major platforms. <sup>2</sup>

The goal of this work was to create a model based on Neuroevolution that would be used to mimic an ecosystem. The model would then be analyzed to gain insight into the factors that contribute to the success of the ecosystem.

### **Methods**

# The Algorithm

Note: the identifier - agent/creature is used interchangeably.

The idea of this algorithm is to give every creature different traits and different neural networks as a brain with each network tackling a specific type of behavior. Unlike other algorithms, I do not seek to find the optimal solution to the simulation, but rather observe a simulated version of evolution and draw conclusions based on the effects of certain factors on the behavior of the creatures. Because of this, they can do as they wish. They are not forced to breed, nor is the size of their pregnancy hardcoded. They are essentially given "free will". All factors that can be reasonably expected to be subject to evolution, are.

Note that the coordinate system ranges from -1 to +1 but the world is round. This means that agents can move from -0.99 to 0.99 (assuming 0.2 movement speed). This also carries with it the challenges of distance calculations for collision detection and others. This was done as a response to the agents sometimes getting stuck in  $\pm 1/-1$ .

<sup>&</sup>lt;sup>2</sup> What is Python? Executive Summary. (n.d.). Retrieved August 30, 2020, from https://www.python.org/doc/essays/blurb/

### **Structure**

- 1. Dependencies
- 2. Defining global simulation parameters
- 3. Inferring simulation specific variables
- 4. Defining agent traits and properties
- 5. Code Documentation
- 6. Complexity and optimizations
- 7. Code snippets explanations
- 8. Statistics

# **Dependencies**

Pip dependencies.

- 1. configparser
- 2. copy
- 3. gc
- 4. math
- 5. os
- 6. random
- 7. time
- 8. typing
- 9. NumPy
- 10. matplotlib
- 11. Pillow
- 12. openpyxl
- 13. savReaderWriter
- 14. pickle
- 15. json
- 16. argparse

Apt dependencies.

1. FFmpeg

Custom dependencies.

1. Modular neural network library (nn.py)

For quick setup, you can use the included requierments.txt file.

### **Global Simulation Parameters**

These are the global constants that define the simulation and their default values:

- 1. INT\_CONST = 1 Energy cost for intelligence (IQ, EQ)
- 2. MOV\_CONST = 5 Energy cost for moving
- 3. ENLB\_CONST = 0.6 energy to mass ratio used to determine if an agent is sick, if so, the agent starts to lose health
- 4. ENGB\_CONST = 0.4 energy to mass ratio used to determine if an agent is healthy, used to gain health and to gain mass if the agent is still growing
- 5. ENL\_CONST = 1 the amount of health lost if an agent's energy is under the threshold.
- 6. ENG\_CONST = 4 the amount of energy gained if an agent is above the energy threshold
- 7. MAX\_LIFE\_SPAN = 200 the maximum number of steps an agent with mass 100 can survive after reaching maturity
- 8. AGE\_CONST = ENG\_CONST (100 / MAX\_LIFE\_SPAN) Age suffered every step, derived from the maximum life span and the amount of health an agent can gain per turn.
- 9. POP\_DENCITY = 1 controls how concentrated is the populous (scales the map size)
- 10. AGING\_TIME = 0.99 the mass percentage required before maturity (age starts)

- 11. G\_SPEED\_FACTOR = 10 controls universal speed, provides a way to control gravity and friction
- 12. FOOD\_CONST = 50 how much energy a food item contains
- 13.  $START\_MASS\_P = 0.95$  The mass to final mass ratio for the initial population
- 14. G \_COL\_CONST = 0.1 the global accuracy for collisions (actual is scaled)
- 15. MIN\_IQ = 1 the minimum number of hidden neurons (per layer).
- 16.  $MAX_{IQ} = 10$  the maximum number of hidden neurons (per layer)
- 17.  $MIN_EQ = 1$  the minimum number of hidden neurons (per layer)
- 18.  $MAX_EQ = 10$  the maximum number of hidden neurons (per layer)
- 19. FOOD\_FLUCT = 1 0.3 the percentage the food count is allowed to dip before food is reproduced.
- 20. GROUP\_FACTOR = 100 A factor applied to the collision constant to define the minimum distance between 2 agents that are in the same group

# **Simulation Specific Variables**

- size\_factor = 1 / (agents / POP\_DENCITY)
- col\_const = G \_COL\_CONST \* self.size\_factor

The rest of the simulation specific variables are either used to keep track of statistics or are necessary for internal calculations (e.g.: agents).

# **Agent Traits and Properties**

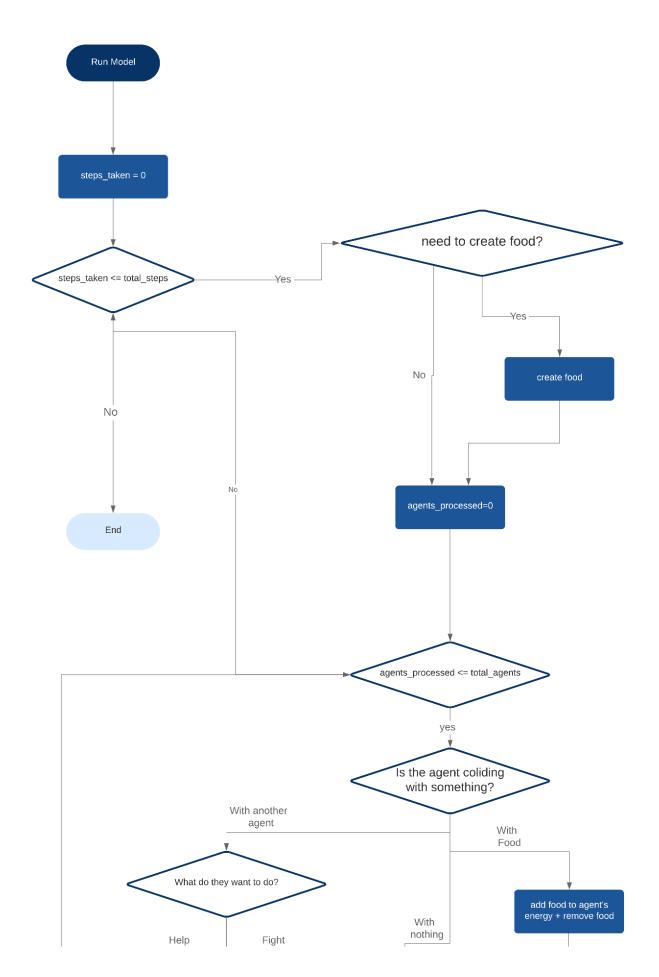
These are the agent traits tracked by the algorithm:

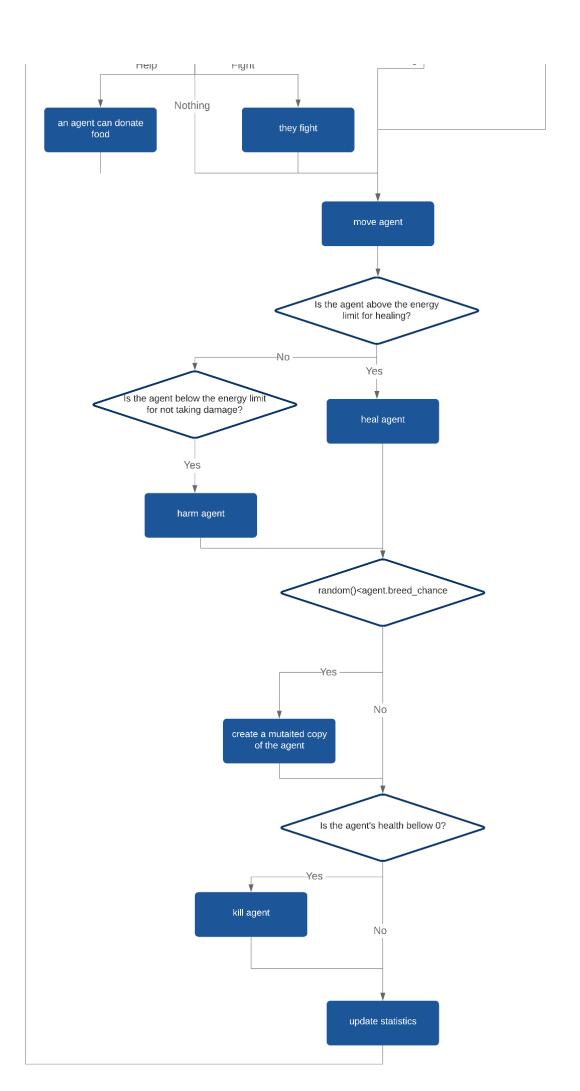
- Movement brain the neural network in charge of moving the agent. Structure [3, IQ, IQ, 1], inputs[distance to the nearest food item, distance to the nearest agent, is the nearest agent stronger], output a value between 0 and one, that is mapped between ±max speed, applied to the agents x coordinate.
- 2. IQ this determines the number of hidden neurons per hidden layer in the neural network in charge of moving the agent.

- 3. Social brain the neural network in charge of agent interactions. Structure [4, EQ, EQ, 2], inputs[if they are close family, the agent's energy to mass ratio, the other agent's food to mass ratio, what agent is stronger], output [probability of the agent attacking, probability of the agent assisting (donating 1 food item)]
- 4. EQ this determines the number of hidden neurons per hidden layer in the neural network in charge of agent interactions
- 5. Mass the mass of the agent. Note that the agent's mass will grow over time as it gets older.
- 6. Final mass indicates when the creature reaches maturity. It is the maximum value a creature mass can take. However, it is also a requirement for reproduction, a creature must reach this a factor of this final mass before it can reproduce.
- 7. Energy the amount of remaining energy the agent has. It starts equal to the agent's mass. Every time it moves or thinks, energy is deducted. It can gain or lose health each step depending on if his energy is above or below a certain level.
- 8. Speed the maximum distance an agent can travel each turn. See Movement Brain (1). Speed has an inverse relation to mass (1/mass). Additionally, speed is scaled by the environment's size factor and global speed factor. See constants.
- 9. Health the creature's "life". If it is below 0, the creature is removed from the simulation. This is effectively the creature's fitness function.
- 10. Breed Mass Divider the initial mass of an agent's child relative to its actual mass.
- 11. Breed Chance the chance of the agent to breed every step, note that breeding requires energy.
- 12. Position (X) the creature's X coordinate on the simulation. This dictates his interaction with other agents, as well as if he is close enough to pick up a food item.
- 13. ID helper variable used by the simulation to keep track of agents as well as to monitor close family.
- 14. Parent ID used to assert if 2 agents are close family (parent/child or siblings)
- 15. Size factor helper variable used by the sim's movement and collision engines to control the size of the simulation.

#### **Model Architecture Outline**

This is a rough overview of a model run





### **Complexity and Optimizations:**

Because a collision detection system needs to know the closest agent and food item to every agent, the original algorithm had an  $O(n^2)$  complexity. It used the following structure:

For every agent:

For every agent:

. . .

For every food item:

. .

Because of the immense amount of time this algorithm required, I optimized it to have an O(n) complexity. This is achieved via the use of sorting functions. A full explanation can be found in the code snippets explanation which provides a detailed explanation for the unintuitive parts of the algorithm.

To test the time improvement, I modeled the time needed for each step of a non-optimized sim (each step is done on a newly generated sim).

Note that these results portray a worst-case scenario where the agents are randomly ordered instead of slightly unordered.

Fig. 1 represents an overview of the newer and older models and the time needed to calculate a step for any number of agents.

Fig. 2 represents a close up of the faster model and shows the average compute time of 5 tests with the top and bottom bars representing the maximum and minimum values in those tests.

*Fig 1.* 

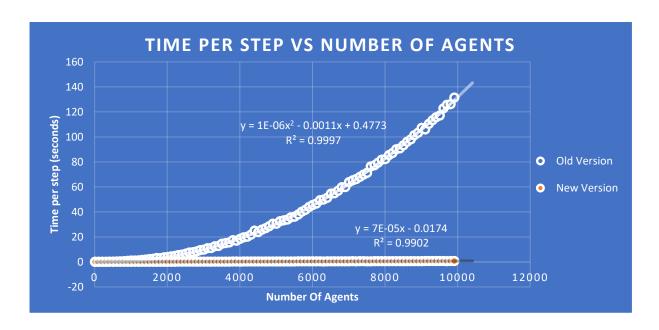
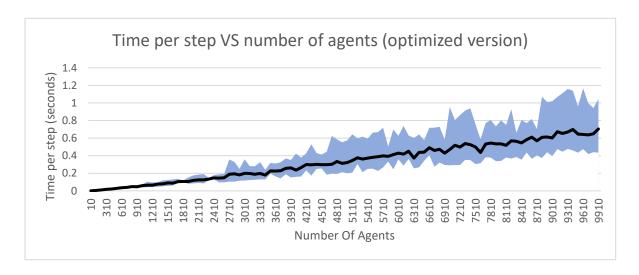


Fig. 2



Even so, it can be seen, that the newer version drastically outperforms its predecessor.

# **Code Snippets Explanations**

These are parts of the code that might require additional explanation to understand

# • Animation projection

Animating the simulation presented a challenge, I needed to map my 1-dimensional coordinate system onto a circle. To start, I created an array with a size proportional to the resolution of the animation and populated it with pixel values corresponding to the

position of food and agents (food was given the value 100 while agents were assigned the value 255). Next, I created a circle with a circumference equal to the length of the 1d array and calculated its radius. Because equal central angles in a circle correspond to equal arch lengths on the circumference, I can map the pixel values from the 1d array to their circle counterparts using basic trigonometric functions. I simply divide 360 by the number of pixels in the 1d array to get  $\Delta\theta$ . Then I loop over the values of the 1d array and by knowing  $\theta$  I can apply a vector to the circle's origin to get the corresponding position on the circle and I apply the corresponding pixel value.

### • Collision detection

As you can see in the complexity section of the algorithm, it is O(n) instead of O(n²). I was able to do this by sorting the agents and the food items by position. Finding the closest agent is easy, simply calculate the distance to the next and previous agent, from there you can check if either of them is smaller than the minimum collision distance. With food items, it's a bit trickier, you keep track of the current food index (starting at 0) then, for every agent, loop from food index until you find a food item with an x coordinate larger than the agents. Then move the food index 1 back. Now, similar to the agent scenario, you compare the distance to the 2 nearest food items. This is, of course, a general overview, in practice, you need to take into consideration what happens if the food is eaten or the around the map. I contemplated whether to allow multiple interactions at once, this would be achieved by searching left and right until no match is detected as opposed to simply searching the closest 2. I decided against this course of action for now, but I leave this as a potential improvement for further versions.

### • Group detection

This is relatively straightforward compared to the others. It is severely limited and cannot act intelligently or keep track of groups as they evolve and clash. Instead it simply defines a maximum distance to the previous agent where you are still considered part of the group

#### **Statistics**

SPSS was used to run statistical correlations and general analysis on the data generated by 3 successive simulations.

The bivariate correlation was done using Person's r.

# **Results**

The following results are the processed outputs of 3 simulations. The initial agent count was set at 500. Each simulation ran for 1,000,000 steps.

Tables 1, 3, and 5 present the bivariate correlation between the following variables: number of steps, number of agents, average agent mass, amount of food consumed, average agent IQ, average agent EQ, average agent breeding mass divider, average agent breed chance, fight count relative to population size, help count relative to population size, ignore count relative to population size, number of groups, and close family ratio in groups for sims 1,2 and 3 respectively.

Tables 2, 4, and 6 present basic statistical analysis for each of the variables described in tables 1, 3, and 5. Specifically, they track: mean, median, Std. Deviation, range, minimum, and maximum.

Figures 3, 4, and 5 present the values of the previously mentioned variables, over the span of the simulation.

Figure 6 represents an animation of a separate simulation spanning 1,000 steps. It is only presented as a proof of concept to allow for intuitive visual understanding and is not meant for meaningful analysis.

The raw SPSS files will be available in the GitHub repository accompanying this project.

### Sim 1

Elapsed time (spell free CPU server) - 18D 11H 9M 37S

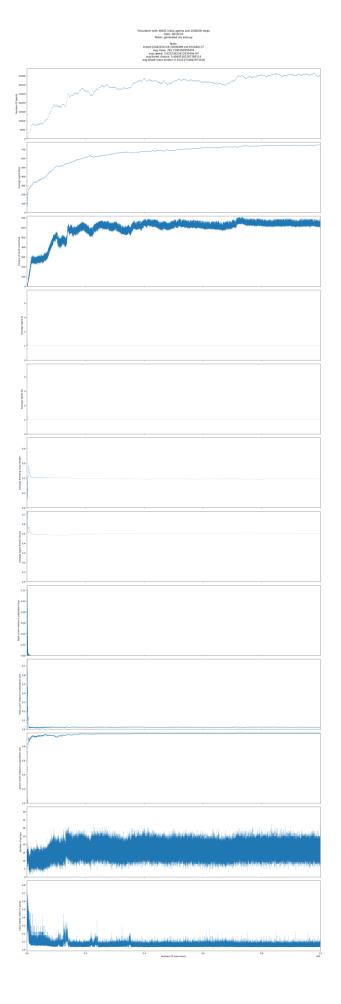
Table. 1

		Steps	Number Of Agents	Average Agent Mass	Amount of Food Consumed	Average Agent IQ	Average Agent EQ	Average breeding mass divider	Average Agent Breed Chance	Fight count relative to population size	Help count relative to population size	Ignore count relative to population size	Number of groups	Close family ration in group
Steps	Pearson Correlation	1	.749	.730	.523	025	025	.241	220	055	075	.245	.096	269
	Sig. (2-tailed)		0.000000E+0	0.000000E+0	0.000000E+0	2.0676E-134	2.0676E-134	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Number Of Agents	Pearson Correlation	.749	1	.973	.921	070	070	168	058	157	164	.451	.254	506
	Sig. (2-tailed)	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Average Agent Mass	Pearson Correlation	.730**	.973	1	.909**	096	096	208	065	196**	195	.490**	.227**	516
	Sig. (2-tailed)	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Amount of Food Consumed	Pearson Correlation	.523	.921	.909	1	104	104	297	113	231	245	.527	.304	589
	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Average Agent IQ	Pearson Correlation	025	070	096	104	1	1.000	.357	.151**	.290**	.047	382**	.057**	.146
	Sig. (2-tailed)	2.0676E-134	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Average Agent EQ	Pearson Correlation	025	070	096	104	1.000	1	.357	.151**	.290**	.047	382**	.057**	.146
	Sig. (2-tailed)	2.0676E-134	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Average breeding mass	Pearson Correlation	.241	168	208	297	.357	.357	1	373	254	186	.099	085	.192
divider	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Average Agent Breed	Pearson Correlation	220**	058**	065	113	.151	.151	373	1	.593**	.443**	607**	020	.306
Chance	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	4.43363E-87	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Fight count relative to	Pearson Correlation	055	157	196**	231	.290	.290**	254	.593	1	.554	789**	014	.339
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	1.13972E-45	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Help count relative to	Pearson Correlation	075	164	195	245	.047	.047	186	.443	.554	1	722	033	.503
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	1.3573E-237	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Ignore count relative to	Pearson Correlation	.245	.451	.490	.527	382	382	.099	607	789	722	1	.085	569
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Number of groups	Pearson Correlation	.096	.254	.227**	.304**	.057**	.057**	085	020	014	033	.085	1	573
-	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	4.43363E-87	1.13972E-45	1.3573E-237	0.000000E+0		0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000
Close family ration in group	Pearson Correlation	269	506	516	589	.146	.146	.192	.306	.339	.503	569	573	1
,	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000

<sup>\*\*.</sup> Correlation is significant at the 0.01 level (2-tailed).

 $Table.\ 2$ 

		Number Of Agents	Average Agent Mass	Amount of Food Consumed	Average Agent IQ	Average Agent EQ	Average breeding mass divider	Average Agent Breed Chance	Fight count relative to population size	Help count relative to population size	Ignore count relative to population size	Number of groups	Close family ratio in group
N	Valid	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	0
	Missing	0	0	0	0	0	0	0	0	0	0	0	1000000
Mean		32645.80	707.3483662	610.0481370	1.000523417	1.000523417	.1975626697	.4992347051	.0000476249	.0231473024	.9749230900	16.10122700	
Media	n	34788.00	745.9694299	631.0000000	1.000000000	1.000000000	.1962600826	.4995908947	.0000000000	.0226543177	.9794606486	16.00000000	
Std. D	eviation	6595.006	95.17685043	82.65808613	.0367404014	.0367404014	.0059629664	.0084273072	.0014921066	.0138740283	.0363477334	3.432059713	
Range		39104	727.2776330	716.0000000	3.915625000	3.915625000	.4161827403	.2501035345	.1290322581	.7763157895	.9865538366	42.00000000	
Minim	um	44	51.47084233	.0000000000	1.000000000	1.000000000	.0563782873	.4817307825	.0000000000	.0000000000	.0000000000	1.000000000	
Maxim	um	39148	778.7484754	716.0000000	4.915625000	4.915625000	.4725610277	.7318343171	.1290322581	.7763157895	.9865538366	43.00000000	



**Sim 2**Elapsed time (spell free CPU server) - 3H 12M 9S *Table 3*.

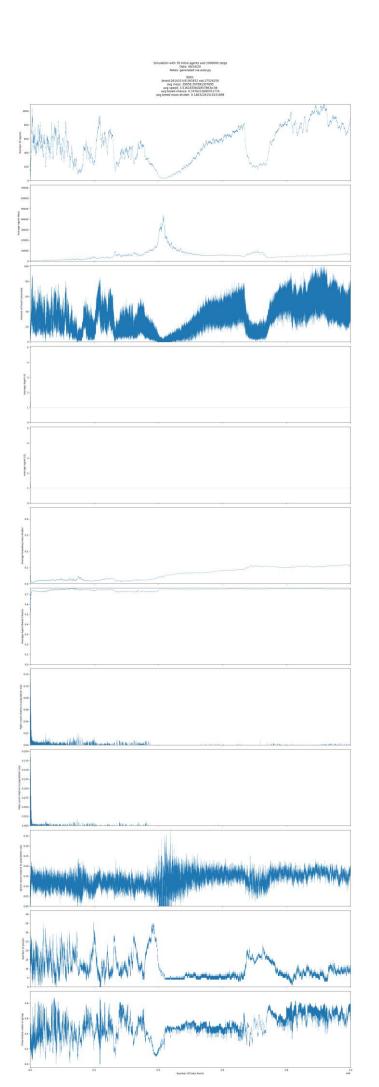
		Steps	Number of Agents	Average Agent Mass	Amount of Food Consumed	Average Agent	Average Agent EQ	Average breeding mass divider	Average Agent Breed Chance	Fight count relative to population size	Help count relative to population size	Ignore count relative to population size	Number of groups	Close family ratio in group
Steps	Pearson Correlation	1	089	.519	047	027	027	.937	300	087	066	.140	498	.280
	Sig. (2-tailed)		0.000000E+0	0.000000E+0	0.000000E+0	2.8897E-165	2.8897E-165	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Number of Agents	Pearson Correlation	089	1	630	.934	015	015	.088	.020	017	023	.482	070	.505
	Sig. (2-tailed)	0.000000E+0		0.000000E+0	0.000000E+0	1.95804E-52	1.95804E-52	0.000000E+0	3.78940E-85	2.37500E-67	6.7438E-119	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Average Agent Mass	Pearson Correlation	.519	630	1	575	017	017	.318	.004	052	040	323	335	157
	Sig. (2-tailed)	0.000000E+0	0.000000E+0		0.000000E+0	5.66859E-62	5.66859E-62	0.000000E+0	1.070452E-4	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Amount of Food Consumed	Pearson Correlation	047	.934	575	1	013	013	.120	.008	015	020	.476	111"	.491
	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0		2.87516E-40	2.87516E-40	0.000000E+0	1.23063E-15	9.40253E-53	8.69632E-87	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Average Agent IQ	Pearson Correlation	027	015	017	013	1	1.000	.061	322**	.350	.396	047	.056	017
	Sig. (2-tailed)	2.8897E-165	1.95804E-52	5.66859E-62	2.87516E-40		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	6.53072E-66
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Average Agent EQ	Pearson Correlation	027	015	017	013	1.000	1	.061	322	.350	.396	047	.056	017
	Sig. (2-tailed)	2.8897E-165	1.95804E-52	5.66859E-62	2.87516E-40	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	6.53072E-66
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Average breeding mass	Pearson Correlation	.937	.088	.318	.120	.061	.061	1	243	054	027	.261	469	.358
divider	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	4.7170E-165	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Average Agent Breed	Pearson Correlation	300	.020	.004	.008	322	322	243	1	354	364	009	.022	048
Chance	Sig. (2-tailed)	0.000000E+0	3.78940E-85	1.070452E-4	1.23063E-15	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	7.52374E-20	2.4885E-103	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Fight count relative to	Pearson Correlation	087	017	052	015	.350	.350	054	354	1	.962	046	.093	061
population size	Sig. (2-tailed)	0.000000E+0	2.37500E-67	0.000000E+0	9.40253E-53	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Help count relative to	Pearson Correlation	066	023	040	020	.396	.396	027	364	.962	1	048	.079	049
population size	Sig. (2-tailed)	0.000000E+0	6.7438E-119	0.000000E+0	8.69632E-87	0.000000E+0	0.000000E+0	4.7170E-165	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Ignore count relative to	Pearson Correlation	.140	.482	323	.476	047	047	.261	009	046	048	1	222	.347
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	7.52374E-20	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Number of groups	Pearson Correlation	498	070	335	111	.056	.056	469	.022**	.093**	.079	222	1	403
	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	2.4885E-103	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
Close family ratio in group	Pearson Correlation	.280	.505	157	.491	017	017	.358	048	061	049	.347	403	1
	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	6.53072E-66	6.53072E-66	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	
	N	999550	999550	999550	999550	999550	999550	999550	999550	999550	999550	999550	999550	999550

<sup>\*\*.</sup> Correlation is significant at the 0.01 level (2-tailed).

# Table 4.

		Number of Agents	Average Agent Mass	Amount of Food Consumed	Average Agent	Average Agent EQ	Average breeding mass divider	Average Agent Breed Chance	Fight count relative to population size	Help count relative to population size	Ignore count relative to population size	Number of groups	Close family ratio in group
N	Valid	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999550
	Missing	0	0	0	0	0	0	0	0	0	0	0	450
Mean		424.5197310	13972.28757	27.52425000	1.000549055	1.000549055	.0988433827	.7403312553	.0000739787	.0000897025	.1303282436	7.463551000	.5356712952
Media	n	421.0000000	9502.153108	25.00000000	1.000000000	1.000000000	.1076129657	.7440492087	.0000000000	.0000000000	.1343638526	5.000000000	.5256024096
Std. D	eviation	274.3366793	13233.26920	19.60046632	.0347066764	.0347066764	.0474813391	.0129776352	.0014084915	.0023309005	.0402967390	5.703148779	.1840700737
Range	:	1073.000000	72632.91304	101.0000000	4.084507042	4.084507042	.4629336375	.3951758832	.1294117647	.2052401747	.3797468354	42.00000000	.9113190731
Minim	um	23.00000000	52.22489083	.0000000000	1.000000000	1.000000000	.0081897683	.3665264462	.0000000000	.0000000000	.0000000000	.0000000000	.0000000000
Maxim	ium	1096.000000	72685.13793	101.0000000	5.084507042	5.084507042	.4711234058	.7617023294	.1294117647	.2052401747	.3797468354	42.00000000	.9113190731

Fig 4.



# Sim 3

# Elapsed time (spell free CPU server) - 4D 9H 36S

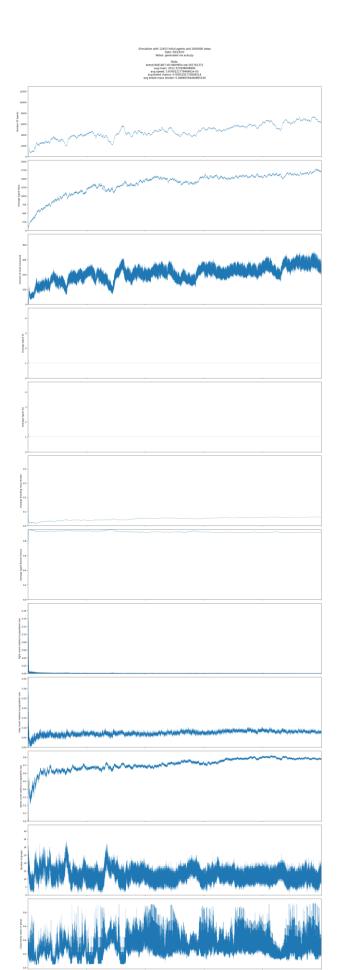
Table. 5

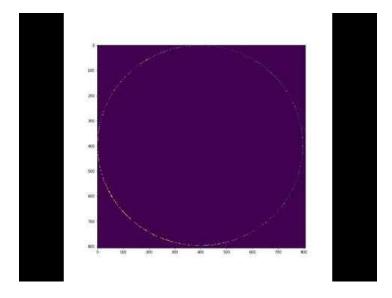
			Number of	Average Agent	Amount of Food	Average Agent	Average Agent	Average	Average Agent	Fight count relative to	Help count relative to	Ignore count relative to	Number of	Close family
		Steps	Agents	Mass	Consumed	IQ	EQ EQ	divider	Breed Chance	population size	population size	population size	groups	ration in group
Steps	Pearson Correlation	1	.962	.888	.884	022	022	.776	.095	084	.752	.866	092	271
	Sig. (2-tailed)		0.000000E+0	0.000000E+0	0.000000E+0	1.5228E-111	1.5228E-111	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Number of Agents	Pearson Correlation	.962	1	.872	.952	032	032	.732	.120	106	.759	.855	153	359
	Sig. (2-tailed)	0.000000E+0		0.000000E+0	0.000000E+0	1.9828E-230	1.9828E-230	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Average Agent Mass	Pearson Correlation	.888	.872	1	.849	058	058	.841	125	176	.767	.930	175	124
	Sig. (2-tailed)	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Amount of Food	Pearson Correlation	.884	.952	.849	1	043	043	.696	.058	129	.716	.816	204	375
Consumed	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Average Agent IQ	Pearson Correlation	022	032	058	043	1	1.000	.404	470	.182	034	111	.062	.009
	Sig. (2-tailed)	1.5228E-111	1.9828E-230	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	1.3298E-257	0.000000E+0	0.000000E+0	3.67404E-18
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Average Agent EQ	Pearson Correlation	022	032	058	043	1.000	1	.404	470	.182	034	111	.062	.009
	Sig. (2-tailed)	1.5228E-111	1.9828E-230	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	1.3298E-257	0.000000E+0	0.000000E+0	3.67404E-18
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Average breeding mass	Pearson Correlation	.776	.732	.841	.696	.404	.404	1	427	.018	.707	.775	078	047
divider	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	8.00933E-69	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Average Agent Breed	Pearson Correlation	.095	.120	125	.058	470	470	427	1	113	115	090	.083	227
Chance	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Fight count relative to	Pearson Correlation	084	106	176	129	.182	.182	.018	113	1	.073	277	.055	.004
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	8.00933E-69	0.000000E+0		0.000000E+0	0.000000E+0	0.000000E+0	8.626659E-5
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Help count relative to	Pearson Correlation	.752	.759	.767	.716	034	034	.707	115	.073	1	.812	007	169
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	1.3298E-257	1.3298E-257	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	1.10530E-12	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Ignore count relative to	Pearson Correlation	.866	.855	.930	.816	111	111	.775	090	277	.812	1	061	133
population size	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0		0.000000E+0	0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Number of groups	Pearson Correlation	092	153	175	204	.062	.062	078	.083	.055	007	061	1	135
	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	1.10530E-12	0.000000E+0		0.000000E+0
	N	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
Close family ration in	Pearson Correlation	271	359	124	375	.009	.009	047	227	.004	169**	133	135	1
group	Sig. (2-tailed)	0.000000E+0	0.000000E+0	0.000000E+0	0.000000E+0	3.67404E-18	3.67404E-18	0.000000E+0	0.000000E+0	8.626659E-5	0.000000E+0	0.000000E+0	0.000000E+0	
	N	999998	999998	999998	999998	999998	999998	999998	999998	999998	999998	999998	999998	999998

<sup>\*\*.</sup> Correlation is significant at the 0.01 level (2-tailed).

Table. 6

		Number of Agents	Average Agent Mass	Amount of Food Consumed	Average Agent	Average Agent	Average breeding mass divider	Average Agent Breed Chance	Fight count relative to population size	Help count relative to population size	Ignore count relative to population size	Number of groups	Close family ration in group
N	Valid	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	1000000	999998
	Missing	0	0	0	0	0	0	0	0	0	0	0	2
Mean		6814.29	1582.573095	261.7613720	1.000458672	1.000458672	.0556022766	.9228355159	.0001227537	.0802587796	.7728563072	12.13154100	.2611748256
Media	n	6833.00	1649.639629	263.0000000	1.000000000	1.000000000	.0594981904	.9210438871	.0000000000	.0827962938	.7974492364	12.00000000	.2563353998
Std. D	eviation	2663.685	335.2913388	76.58496167	.0353978979	.0353978979	.0111601734	.0101707889	.0017326552	.0118894032	.0898872933	3.532779988	.1105391846
Range	:	12846	1982.265633	472.0000000	3.674518201	3.674518201	.4766288511	.4725134421	.1789473684	.3578947368	.8775796599	44.00000000	.9211668104
Minim	um	41	50.43897216	.0000000000	1.000000000	1.000000000	.0157544666	.4812349089	.0000000000	.0000000000	.0000000000	.0000000000	.0307851415
Maxim	ium	12887	2032.704605	472.0000000	4.674518201	4.674518201	.4923833176	.9537483510	.1789473684	.3578947368	.8775796599	44.00000000	.9519519520





### **Discussion**

My objective was to create a computational model that can be used to describe the evolution of an ecosystem. As such, a large part of my results is the algorithm itself, that can be used to obtain much greater insights into the behavior of populations than I was able to.

Getting a stable populous is more challenging than it seems. At first, the population skyrocketed into the tens of thousands and quickly overwhelmed my computer's capabilities. Other times, they quickly died, leaving no obvious reason for their sudden demise. As it turns out I had broken one of the most basic laws of physics, and as such created a "hack". I had forgotten to regulate the reproduction mechanic and allowed for blatant violations of the conservation of mass and energy. Yet even with those fixed, A lot of time was spent fine-tuning the parameters to allow for stable growth. This included implementing a mechanism that allowed for scaling the simulation. Because the simulation's spatial coordinates range from -1 to +1, I had to adjust the necessary distance for collision, as well as the maximum speed of the agents as to virtually scale the coordinate system without actually doing so. Another such alteration was the addition of many variables to the creature's traits, most prominently "breed chance" and "breed mass divider". This had the added benefit of making the simulation much more realistic and indicative of the real world, but its primary objective was to remove the need to manually tune these parameters.

To analyze the results from the simulations I performed a bivariate correlation between all the variables using Person's r. It appears that they are all at least loosely correlated to one another (correlation is significant at the 0.01 level - 2-tailed). This is fascinating. Even though the correlation itself might be weak, this demonstrates that all factors in an ecosystem drastically affect the others.

I would argue that simulations a and c are more successful than simulation b. This is because the total number of agents, as well as food consumed, is higher. Because they use more of their available resources, I consider then to be better. A factor that I found works well for measuring the success of the simulation is the time it takes to complete. Sim b measured at ~3H, simulation c was ~4D while simulation a took a staggering ~18D to complete.

The differences between simulations a and c to simulation b will become more apparent once we start examining the correlations.

Some of these correlations are expected. For example:

• The correlation between the number of agents and the food consumed

$$\rho_a$$
=0.921  $\rho_b$ =0.934  $\rho_c$ =0.952

This is mostly self-explanatory. The more food consumed the higher the agent's health, the more children it can have.

• Average agent mass and close family ratio in group

$$\rho_a$$
=-0.516  $\rho_b$ =-0.124  $\rho_c$ =-0.157

This is a loose correlation between the average agent mass and the close family ratio in the group. While they may seem unconnected at first, the more massive an agent is the more offspring it has. This effect is however counteracted by the dilution of family members by the increased number of agents in groups. To get a better understanding of the strength of this correlation we can look to the STD. deviation of the mass in each simulation. Unfortunately, STD over time was not tracked and is a potential improvement for the future.

These correlations indicate that the simulation is at least partially successful as it produces results that are expected using common intuition.

I will now cover some more interesting correlations.

# • Agent breed mass divider with the progress of the simulation (Steps)

$$\rho_a$$
=0.241  $\rho_b$ =0.937  $\rho_c$ =0.776

This makes a lot of sense to me. I propose that at the beginning of the simulation, before agents were able to optimize their search for food, having children was a big risk and so the new children were given limited mass. But as time went on the risk decreased, and agents could now afford to gamble a lot of their health. It is important to note that although I talk about the agent taking risks, I am referring to the concept of the selfish gene. The continuation of the gene is dependent on its current carrier but also on its offspring. Although genes are not used directly in this model, the general concept still applies.

The reason that the first simulation does not fall under the same pattern would be because of its lower breed chance compared to the other two. This allowed it to mostly start on 0.2 instead of reaching it after a prolonged period.

### • Close family and number of agents

$$\rho_a$$
=-0.506  $\rho_b$ =0.505  $\rho_c$ =-0.359

This is interesting because the correlations appear to be reversed.  $\rho_b$  is expected as having more close family seems to increase the chances of survival as intuitively family members are less likely to attack other family members as well as are more likely to help them. However, I propose that the reason for the correlation we observe in sims a and c is because of the increased number of agents diluting the close family ratio. The more agents, the less close family.

I believe that the reason close family affects the number of agents and not the other way around in sim b is because of the fewer number of agents that increase the effect of each agent and its family. This is contrary to sims a and c where a large number of agents appear to dilute most of the advantages.

### Amount of food consumed and average agent mass

$$\rho_a$$
=0.908  $\rho_b$ =-0.575  $\rho_c$ =0.849

Here the correlations between sim a and sim c to sim b are also reversed. While  $\rho_c$  and  $\rho_a$  seem expected, the more food consumed the higher the average mass,  $\rho_b$  is the exact opposite.

The number of agents does not seem to be the cause because they both have similar ratios regarding the number of agents and food consumed:  $\rho_a$ =0.921  $\rho_b$ =0.934  $\rho_c$ =0.952. In fact, simulation c seems to have more agents when there is more food compared to sim b.

I propose that the reason for this is that the second simulation was too erratic for such a correlation. Because food only affects an agent's mass directly if he is still growing and agent mass itself is constantly changing due to childbirth, this correlation might only show on more stable simulations with a high agent count. The number of agents in sim2 hover around 424 on average, while sims 3 and 1 boast 6,833 and a staggering 32,645 average agents respectively.

This is in addition to factors like the high agent mass in sim  $2 \sim 9,502$  compared to sim  $3 \sim 1,582$  which would reduce the effects of factors like food consumed on the average mass.

Another interesting observation is that agents chose to mostly ignore one another instead of fighting. I have been unable to determine the reason for this behavior although I suspect it arises from a mistake in the fight mechanic where fighting might have more drawbacks than rewards. It is however possible that the agents with a higher tendency to fight simply killed themselves. This is in line with the idea that animals refrain from cannibalism.

I would suggest that in the future, more analysis should be done on the differences between the simulations. Additionally, I would recommend the use of real-world experiments to approximate the actual values of the constants used.

### **Assumptions and Limitations**

Throughout the project, I have made several key assumptions.

- Agents cannot learn. At first glance, this may seem like a glaring flaw, as I'm treating
  agents as individuals yet without the ability to learn. However, I would argue that the
  learning process of an individual is largely irrelevant because unlike natural agents,
  these agents can pass on much more information via their mutated cloned brains to
  their offspring.
- I make several assumptions concerning the way that mass governess the ability of the creatures. I make no allowances for muscle mass to body mass ratio and seem to be giving a large mass a great advantage. In the future simulation should make use of 2 different variables instead of relying solely on one.

Here is proof that acceleration is defined by muscle mass to body mass ratio instead of mass: (This is of course still a major oversimplification, it does not take into account heat dissipation, internal machinery and much more)

$$a = \frac{F}{m}$$

$$F = m k^*$$

$$a = \frac{mk}{m} = k$$

As you can see, an agent's acceleration is not related to its mass but rather to its muscle mass to body mass ratio. In future versions, this can be resolved by using 2 variables for mass.

- \* k is the muscle mass to body mass ratio
- Agents age in a constant fashion, regardless of their mass. This, in turn, diminishes the effect of aging on larger creatures. I would argue that ageing is a vague process,

because its less of a biological process and more of a physical one. In spite of this, life expectancy does seem to be linearly correlated to mass.<sup>3</sup>

- The simulation only uses 1 axis. While a 2 or even 3 axis system might represent reality better (e.g.: allow for advance fighting tactics), I believe that the computational drawbacks outweigh the advantages for this project. To partially compensate for this, agents can "jump" over other agents, in other words the simulation doesn't check to see if there is an agent in the way before a creature moves. This would, however, be an avenue for further improvement should anyone express interest in improving the model.
- Because the simulation attempts to work via the macro world and not to simulate physics, it necessarily is less accurate. The assumptions I made are closely connected to this fact. And while I have done my best to minimize them, as well as justify them to a certain degree, they still present limitations to the accuracy of the simulation. For example, all creatures are equally efficient. While this does not accurately describe the natural world, because all creatures strive for maximum efficiency, I believe that this is acceptable. This assumption allows me to ignore the physics side of things as creating a physics simulation and allowing agents to develop efficient solutions for aerodynamics as well as cell level reactions would be a near-complete waste of computational resources for modeling macro events as well as way beyond my technical capabilities.
- IQ and EQ are immutable. Because of the nature of neural networks, IQ and EQ are completely static. They do not change from parent to child, unlike other traits. I believe that this is the reason that all the agents in every simulation I ran, were reduced to 1 EQ and IQ after the first couple of steps. IQ and EQ produce unwanted strain on the agent's energy repository. In the early development stages, IQ and EQ produce no meaningful advantage, yet I believe that in later stages this may change.

This poses an interesting challenge, one that is beyond my ability to solve. However, a possible partial solution would be to allow for a steady supply of new random

<sup>&</sup>lt;sup>3</sup> Szekely, P., Korem, Y., Moran, U., Mayo, A., & Alon, U. (2015). The Mass-Longevity Triangle: Pareto Optimality and the Geometry of Life-History Trait Space. PLOS Computational Biology, 11(10), e1004524. https://doi.org/10.1371/journal.pcbi.1004524

agents throughout the simulation's life cycle. This would not fully resolve the issue because these agents would need to start from scratch, however, it would allow for the introduction of agents with high EQ and IQ.

- The constants of the simulation, while not arbitrary, are not necessarily indicative of the natural world and as such they misrepresent reality. This could probably be overcome by dedicating time to each constant and attempting to approximate it.
- While the simulation does keep track of numerous variables and statistics, it does not present the full picture. To partially compensate for this, I created the animate function that creates a rendering of the simulation. However, it is difficult to ascertain any meaningful amount of information beyond the events that occur on the macro scale. This is the reason its sparsely missioned.
- The food class in the simulation always boats the same food reward. This is an interesting avenue for further discussion. Should the food amounts be generated randomly the agent might be provided with multiple food items and it will have to pick intelligently between distance and food amount. Furthermore, the introduction of seasons would also be an interesting factor.
- Agents are not able to maintain a memory of past interactions and communicate
  information concerning them, nor are they able to identify other agents with whom
  they interact. This might be fixed by using LSTM or RNN instead of a simple neural
  network, in addition to ids being passed while interacting.
- Agents can only interact with 1 agent and 1 food item per step. I detail in my code snippet explanation a possible solution that should mostly maintain the efficiency improvements from the sorting solution. Even so, the step function will become an O(n²) algorithm.

There are of course other limitations and assumptions that I did not mention here or realize.

### Conclusion

In conclusion, a simulated ecosystem can grant us new insights into the inner workings of complex interactions between innumerable values. As I have demonstrated there exist loose correlations between all the variables I have studied, most of whom are verified at beyond the

5-sigma level. I was able to gain interesting insights into the mechanics of these interactions' causes and implications.

# **Appendix 1 – Using the simulation**

# **Initial installation (for all use cases)**

1. Clone the code from <a href="https://github.com/ikoursh/EcoSystemProject">https://github.com/ikoursh/EcoSystemProject</a> by running:

git clone https://github.com/ikoursh/EcoSystemProject

2. Then enter the model directory:

cd ecosystemproject/proj

3. Finally, install the dependencies:

pip install -r requierments.txt

If you want to add support for animation you will need an installation of FFmpeg. To do so you can go <a href="here">here</a> (don't forget to add it to PATH, <a href="this">this</a> is a great explanation for windows)

Note that pip install ffmpeg does not seem to work, however, Linux users may use apt.

# **GUI** (Windows only officially supported)

This option might be a bit buggy and has not been tested on Linux or Mac. It is however the preferred option for beginners.

Simply install the program using the included installer (ecosystemproject/GUI/dist/EcoSystem-Project-GUI Setup 1.0.0.exe)

Then open the program, and in the settings tab enter the path to auto.py.

You can now create simulations using the new simulation menu. Simulations will automatically display in your "my simulations" menu. Once a simulation has finished you can click on it to see the output files.

Note that EnvSim will attempt to run in the background and can be terminated using the task manager or the slightly buggy tray bar. Once the program is quit the simulations will too and they cannot be resumed.

Note, if you choose to enable spell integration you must commit all local changes before trying to use Spell.

### **CLI**

This option is easy and flexible. Run "python auto.py —help" for more information about the parameters. It is very easy to use and user friendly.

Examples can be found in the GitHub repository and the API documentation.

### As a Library

Please check the API documentation for more information.

\* if you attempt to process close family ratio via the generated SAV(SPSS) file you might encounter multiple errors. This can be fixed by rounding the Close\_family\_ratio\_in\_group values to fewer decimal points. To do this click transform => compute variable, enter the name of the new variable and enter  $\frac{\text{RND}(\text{Close\_family\_ratio\_in\_group *10**x})}{10**x}$  (where x is the precision, 16 is the recommended value).

# Appendix 2 – GitHub Repository

All of the source code is available for free under GPL V3 at this project's <u>GitHub repository</u>.

# **Appendix 3 - Code Documentation**

The core code used in the project has been documented via Google's API doc guidelines.

The documentation has been compiled using SPHINX and is available here: <a href="https://inbarkoursh.com/ecosystemproject/">https://inbarkoursh.com/ecosystemproject/</a> as well as on this project's GitHub repository here: <a href="https://github.com/ikoursh/EcosystemProject/tree/master/docs/">https://github.com/ikoursh/EcosystemProject/tree/master/docs/</a>

# **Appendix 4 - Code**

It is recommended that all code be viewed using this project's <u>GitHub repository</u> and not here.

The code for the GUI as well as the documentation is only available in the GitHub repository.

### nn.py:

```
def mutate(a: numpy.ndarray) -> numpy.ndarray:
 Mutate Function
 return a + xp.random.normal(0, 0.1)
class NNLayer:
 def __init__(self, nodes: int, prev_nodes: int) -> None:
   self.nodes = nodes
   self.prev_nodes = prev_nodes
   self.weights = xp.random.random((prev_nodes, nodes))
   self.bias = xp.random.random(nodes)
 def feed_forward(self, xs: numpy.ndarray) -> numpy.ndarray:
   if len(xs) != self.prev_nodes:
     raise Exception(
```

```
self.prev_nodes, len(xs)))
   return sigmoid(xp.add(xp.matmul(xp.array(xs), self.weights), self.bias))
 def mutate(self) -> None:
   self.weights = mutate(self.weights)
   self.bias = mutate(self.bias)
 def __repr__(self) -> str:
   return "weights: {} bias: {}".format(self.weights, self.bias)
class NeuralNetwork:
 def __init__(self, nodes: list) -> None:
   if len(nodes) < 2:</pre>
     raise Exception(
   self.layers = []
   for i in range(len(nodes) - 1):
     self.layers.append(NNLayer(nodes[i + 1], nodes[i]))
 def feed_forward(self, xs: numpy.ndarray) -> numpy.ndarray:
```

### model2.py:

```
import copy
import gc
import math
import os
import random
import time
from typing import Tuple
import matplotlib
import matplotlib import animation
from nn import NeuralNetwork

from PIL import PngImage Plugin
import openpyxl # work with excel
import savReaderWriter # work with spss
import pickle # support serialization
```

```
matplotlib.get_backend()
import sys
if sys.version_info[0] < 3 or sys.version_info[1] < 4:</pre>
 raise Exception(
if sys.version_info[0] > 3:
 print("WARNING this program was write to support python 3. Use any future versions at your
config = configparser.ConfigParser(inline_comment_prefixes="#")
config.read('config.ini')
INT_CONST = float(config["VARIABLES"]["INT_CONST"])
MOV_CONST = float(config["VARIABLES"]["MOV_CONST"])
ENLB_CONST = float(config["VARIABLES"]["ENLB_CONST"])
ENGB_CONST = float(config["VARIABLES"]["ENGB_CONST"])
ENL_CONST = float(config["VARIABLES"]["ENL_CONST"])
ENG_CONST = float(config["VARIABLES"]["ENG_CONST"])
MAX_LIFE_SPAN = float(config["VARIABLES"]["MAX_LIFE_SPAN"])
AGE_CONST = config["VARIABLES"]["AGE_CONST"]
if AGE_CONST == "auto":
 AGE_CONST = ENG_CONST - (100 / MAX_LIFE_SPAN)
 AGE_CONST = float(AGE_CONST)
POP_DENCITY = float(config["VARIABLES"]["POP_DENCITY"])
AGING_TIME = float(config["VARIABLES"]["AGING_TIME"])
G_SPEED_FACTOR = float(config["VARIABLES"]["G_SPEED_FACTOR"])
FOOD_CONST = float(config["VARIABLES"]["FOOD_CONST"])
START_MASS_P = float(config["VARIABLES"]["START_MASS_P"])
G_COL_CONST = float(config["VARIABLES"]["G_COL_CONST"])
MIN_IQ = float(config["VARIABLES"]["MIN_IQ"])
MAX_IQ = float(config["VARIABLES"]["MAX_IQ"])
MIN_EQ = float(config["VARIABLES"]["MIN_EQ"])
MAX_EQ = float(config["VARIABLES"]["MAX_EQ"])
FOOD_FLUCT = float(config["VARIABLES"]["FOOD_FLUCT"])
GROUP_FACTOR = float(config["VARIABLES"]["GROUP_FACTOR"])
def map_from_to(x: float, a: float, b: float, c: float, d: float) -> float:
```

```
return (x - a) / (b - a) * (d - c) + c
GRAPHS_FOLDER = "/graphs-0.3"
ANIMATION_FOLDER = '/animations-0.1'
if not os.path.isdir('.' + GRAPHS_FOLDER):
 os.mkdir('.' + GRAPHS_FOLDER)
if not os.path.isdir('.' + ANIMATION_FOLDER):
 os.mkdir('.' + ANIMATION_FOLDER)
if not os.path.isdir("./saved"):
 os.mkdir("./saved")
class Agent:
```

breed\_chance(float): The chance for the agent to breed at every step. See :attr:`model2.Agent.breed\_chance`

size\_factor(float): A variable used by the sim's movement and collision engines to control the size of the simulation. See :attr:`model2.Sim.size\_factor`

move\_brain(NeuralNetwork): The neural network the agent will use to decide where to move. This is either provided by the creature's parent (via a mutated brain) or is generated from scratch (for the initial population) See :attr:`model2.Agent.move\_brain`.

social\_brain(NeuralNetwork): The neural network the agent will use to interact with other agents. This is either provided by the creature's parent (via a mutated brain) or is generated from scratch (for the initial population) See :attr:`model2.Agent.social\_brain`.

parent\_id(int): This variable is used to detect close family by the simulation. It is then used as a variable in agent interactions. See :attr:`model2.Agent.parent\_id`

#### Attributes

energy(float): Tracks the amount of energy an agent has. energy is acquired by eating and lost by moving or thinking. Energy allows agents to grow and regenerate health. If an agent has low energy, he will take damage. See :meth:`model2.Agent.think`.:meth:`model2.Agent.move`.:meth:`model2.interact`

speed(float): The maximum velocity at which the creature can move. Inversely related to mass (1/mass) health(float): How much damage can a creature sustain. When it reaches 0, the creature dies.

x (float): The agents position. See :meth: `model2.Agent.move`

id(int): An id used by the simulation class to keep track of agents. See :attr:`model2.Agent.parent\_id` for use case.

iq (int): The number of neurons in every hidden layer of the agent's neural network that is responsible for movement. See :attr:`model2.Agent.move\_brain` for use case.

eq (int): The number of neurons in every hidden layer of the agent's neural network that is responsible for agent to agent interactions. See :attr:`model2.Agent.social\_brain` for use case

mass(int): The mass of the agent, represents its size final\_mass(int): The final mass of the agent.

breed\_mass\_div(float): The initial mass to final mass ratio of the agent's children. See meth:`model2 Agent breed`

breed\_chance(float): The chance for the agent to breed at every step. See :meth:`model2.Agent.breed'

size\_factor(float): A variable used by the sim's movement and collision engines to control the size of the simulation. See :attr:`model2.Sim.size\_factor`

move\_brain(NeuralNetwork): The neural network the agent will use to decide where to move. Structure: [3, iq, iq, 1] See :meth:`model2.Agent.think` and :attr:`model2.Agent.iq`.

social\_brain(NeuralNetwork): The neural network the agent will use to interact with other agents. Structure [4, eq, eq, 2] See :meth:`model2.interact` and :attr:`model2.Agent.eq`.

```
parent_id(int): This variable is used to detect close family by the simulation. It is then used as a variable
def __init__(self,
      eq,
      final_mass,
      breed_mass_div,
      breed_chance,
      size factor.
      move_brain=None,
      social_brain=None,
      parent_id=None):
  self.iq = iq
  self.eq = eq
 if move_brain is not None:
    self.move_brain = move_brain
    if iq == 1: # if the size of the hidden layers is 1, the amount of hidden layers doesn't matter
      self.move_brain = NeuralNetwork([3, 1])
      self.move_brain = NeuralNetwork([3, iq, iq, 1])
 if social brain is not None:
    self.social_brain = social_brain
    if eq == 1: # if the size of the hidden layers is 1, the amount of hidden layers doesn't matter
      self.social_brain = NeuralNetwork([4, eq, 2])
      self.social_brain = NeuralNetwork([4, eq, eq, 2])
  self.parent_id = parent_id
  self.mass = mass
  self.energy = mass
  self.speed = (1 / mass) * size_factor * G_SPEED_FACTOR
  self.health = mass
  self.final_mass = final_mass
  self.breed_mass_div = breed_mass_div
  self.breed_chance = breed_chance
```

self.x = x

self.id = id

```
self.size_factor = size_factor
def age(self) -> None:
  if self.mass > self.final_mass * AGING_TIME:
    self.health -= AGE_CONST
def think(self, d_food: float, d_agent: float, s_agent: 'Agent') -> float:
    d_food(float): The distance to the nearest food item
  out = self.move_brain.feed_forward(
    [map_from_to(d_food, -1, 1, 0, 1), map_from_to(d_agent, -1, 1, 0, 1),
    int(s_agent.mass > self.mass)])
  self.energy -= self.iq * INT_CONST
  return map_from_to(float(out[0]), 0, 1, -self.speed, self.speed)
def move(self, dx: float) -> None:
    dx: Distance to travel
  self.x += dx
  if self.x > 1: # make the world round
    self.x = 1 - self.x
```

```
if self.x < -1:
    self.x = -self.x - 1
  self.energy -= MOV_CONST * dx
def breed(self, nid: int) -> 'Agent':
   nid(int): The id of the new child
  nb = copy.deepcopy(self.move_brain)
  nb.mutate()
  nsb = copy.deepcopy(self.social_brain)
  nsb.mutate()
  nm = math.ceil((self.mass + random.randrange(-10, 10)) * self.breed_mass_div)
  self.health -= nm
  self.mass -= nm
 if self.health <= 0 or self.mass < 1 or nm < 1:
    self.health = -1
  return Agent(self.iq, self.eq, nm, self.x + random.uniform(0.001, -0.001), nid,
         math.ceil(nm / self.breed_mass_div),
         self.breed_mass_div + random.uniform(0.01, -0.01), self.breed_chance + random.uniform(0.01,
         self.size_factor, nb, nsb, self.id)
def eat(self, food: int) -> None:
 if self.mass < self.final_mass and self.energy / self.mass > ENGB_CONST:
    self.mass += food # update mass and speed
    self.speed = (1 / self.mass) * self.size_factor * G_SPEED_FACTOR
    self.energy += food
```

```
def __str__(self) -> str:
   return str(self.id)
def fight(a1: Agent, a2: Agent) -> None:
 a1.health -= a2.mass
 a2.health -= a1.mass
 a1.energy += a2.mass
 a2.energy += a1.mass
def interact(a1: Agent, a2: Agent, s) -> None:
 a1.energy -= a1.eq * INT_CONST # subtract energy used for interaction thought
 a2.energy -= a2.eq * INT_CONST
 close_family = a1.parent_id == a2.id or a2.parent_id == a1.id
 s1 = a1.social_brain.feed_forward(
    [1 if close_family else 0, a1.energy / a1.mass, a2.energy / a2.mass, 1 if a1.mass > a2.mass else 0])
 s2 = a2.social_brain.feed_forward(
    [1 if close_family else 0, a2.energy / a2.mass, a1.energy / a1.mass, 1 if a1.mass < a2.mass else 0])
 if s1[0] > 0.5 or s2[0] > 0.5: # if either agent wants to fight
    fight(a1, a2)
    s.fight += 1
    a1.energy -= FOOD_CONST
   a2.energy += FOOD_CONST
```

```
s.help += 1
 if s2[1] > 0.5:
   a1.energy += FOOD_CONST
   a2.energy -= FOOD_CONST
   s.help += 1
   s.nothing += 1
class Food:
 def __init__(self, x: float) -> None:
   self.x = x
def mk_round(d: float) -> float:
   return d - 2
   return -d - 1
   return d
```

```
breed(int): The number of times agents breed in total
  help(int): The number of times agents help one another in total
  gcsteps(int): The number of total steps taken (if run/animate is used more than once)
  fight_OT(list[int]): A list of the amount of fighting over time. See :meth:`model2.fight`,
def __init__(
    self,
```

agents: int = 500,

```
food_count: int = None,
if food_count is None:
  food_count = 5 * agents
if (not isinstance(food_count, int)) or (not isinstance(agents, int)):
  raise TypeError
self.size_factor = 1 / (agents / POP_DENCITY)
self.col_const = G_COL_CONST * self.size_factor
self.food_count = food_count
self.breed = 0
self.kill = 0
self.fight = 0
self.help = 0
self.nothing = 0
self.id = 0
self.eat = 0
self.agents = []
self.food = []
self.gcsteps = 0
self.dataPoints = 0
self.interactions = 0
for i in range(agents): # create initial population
  mass = math.ceil(random.randrange(1, 100))
  self.agents.append(
    Agent(int(random.randrange(MIN_IQ,
                  MAX_IQ)), int(random.randrange(MIN_EQ, MAX_EQ)),
       math.ceil(mass * START_MASS_P),
       random.uniform(-1, 1), self.id, mass, random.random(), random.random(), self.size_factor))
  self.id += 1
self.group()
self.i_OT = []
self.number_of_agents_OT = []
self.mass_OT = []
self.eat_OT = []
self.iq_OT = []
self.eq_OT = []
self.breed_mass_div_OT = []
self.breed_chance_OT = []
self.interactions_OT = []
self.fight_OT = []
self.help_OT = []
```

```
self.nothing_OT = []
  self.relative_groups_OT = []
  self.close_family_in_group_OT = []
  self.cfood()
def get_fn(self) -> str:
  return '{}-{}-{}'.format(
    len(self.agents), self.gcsteps,
    time.strftime("%d%M%Y%H%M%S", time.localtime()))
def save(self, file: str = None) -> None:
  if file is not None and file.endswith(".envs"):
    raise ValueError("File must end with .envs")
  pickle.dump(self, open(file if file is not None else "save/" + self.get_fn() + ".envs", "w"), 4)
@classmethod
def load(cls, filename: str) -> "Sim":
   the loaded sim
  with open(filename, 'rb') as f:
    return pickle.load(f)
def run(self, steps: int = 1000, print_freq: int = None, max_attempts: int = 1, data_point_freq: int = 10,
```

```
gui: bool = False) -> Tuple[bool, int]:
  See also:
  if max_attempts == -1: # if maximum attempts is -1, make it effectively infinite
    max_attempts = 2 ** 32
  sim_copy = copy.deepcopy(self) # create a copy of the sim to use as a restore point
 if print_freq is None: # if print frequency is not specified, set it so that the model prints every 1% of
    print_freq = steps / 100
  for a in range(max_attempts):
    failed = False
    for i in range(steps):
      if not self.step(): # call step, if it failed, stop this attempt
        self._dict_.update(copy.deepcopy(sim_copy)._dict_) # resetting the sim to its original state
        failed = True
      if i % print_freq == 0:
        self.progress(steps, i, gui)
      if i % data_point_freq == 0:
        self.update_stats() # update statistics
    if not failed:
      self.progress(steps, steps, gui)
  return False, max_attempts
def group(self) -> int:
```

```
prev_a = self.agents[0]
  prev_a.group = 0
  for a in self.agents:
    a.group = prev_a.group + (0 if abs(a.x - prev_a.x) < self.col_const * GROUP_FACTOR else 1)
    prev_a = a
 return prev_a.group
def progress(self, steps: int, csteps: int, gui: bool) -> None:
 if gui:
    print(json.dumps({
      "steps": csteps,
      "food": len(self.food),
      "agents": len(self.agents)
    sys.stdout.flush()
    print(
        .format(round((csteps / steps) * 100, 2),
            csteps, steps,
            len(self.agents),
            len(self.food))) # print status
def update_stats(self) -> None:
  agent_count = len(self.agents) # save time
  self.gcsteps += 1
  self.dataPoints += 1
  self.i_OT.append(self.gcsteps)
  self.number_of_agents_OT.append(agent_count)
  self.interactions_OT.append(self.interactions)
  self.interactions = 0
  self.relative_groups_OT.append(self.group())
  helper_mass = []
  helper_iq = []
  helper_eq = []
```

```
helper_bmd = []
  helper_bch = []
  helper_group_i = -1
  helper_group_size = []
  helper_close_family_group = []
  for a in self.agents:
    helper_mass.append(a.mass)
    helper_iq.append(a.iq)
    helper_eq.append(a.eq)
    helper_bmd.append(a.breed_mass_div)
    helper_bch.append(a.breed_chance)
    if a.group != helper_group_i:
      a.group = helper_group_i
      helper_close_family_group.append([])
      helper_group_size.append(0)
      helper_group_i += 1
    helper_close_family_group[helper_group_i].extend([a.id, a.parent_id])
    helper_group_size[helper_group_i] += 1
  self.close_family_in_group_OT.append(np.mean(
    [(len(helper_close_family_group[i]) - len(set(helper_close_family_group[i]))) / helper_group_size[i]
    range(helper_group_i)])) # the average amount of duplicates is the amount of close family
 self.mass_OT.append(np.mean(helper_mass))
  self.eat_OT.append(self.eat)
 self.eat = 0
  self.iq_OT.append(np.mean(helper_iq))
  self.eq_OT.append(np.mean(helper_eq))
  self.breed_mass_div_OT.append(
    np.mean(helper_bmd))
  self.breed_chance_OT.append(
    np.mean([a.breed_chance for a in self.agents]))
  self.fight_OT.append(self.fight / agent_count)
  self.help_OT.append(self.help / agent_count)
  self.nothing_OT.append(self.nothing / agent_count)
  self.help = 0
 self.fight = 0
  self.nothing = 0
def cfood(self) -> None:
  self.food = []
  for i in range(int((1 - FOOD_FLUCT) * self.food_count)):
   self.food.append(Food(random.uniform(-1, 1)))
```

```
def step(self) -> bool:
  self.gcsteps += 1
  if len(self.food) < FOOD_FLUCT * self.food_count:</pre>
    self.cfood()
  if len(self.agents) <= 1:</pre>
    print("ALERT: the model has died")
  self.agents.sort(
    key=lambda ag: ag.x
  ) # sort agents by position, allows to quickly determine the closest agent with low complexity
  self.food.sort(key=lambda ag: ag.x)
  food_index = 0 # used to find closest food item with low complexity
  debug = []
  for a in range(len(self.agents)):
    if a >= len(self.agents): # agents can be removed but the range isn't updated
    ax = self.agents[a].x
    lf = None
    tmp_fi = food_index # crete debug data
    debug_line = "food length: " + str(len(self.food)) + " start search index: " + str(food_index)
    for i in range(food_index, len(self.food)):
        if self.food[i].x > ax:
          food_index = i - 1
          tf = self.food[i]
          debug_line += " found food index " + str(i) + " new food index set to " + str(i - 1)
          lf = self.food[i]
        print("there was an unexpected crash")
        print("Agent number " + str(a) + " out of " + str(len(self.agents)))
        print("Food index at crash was " + str(i))
        print("Length of food list is " + str(len(self.food)))
        print("The food index at the start of the search was " + str(tmp_fi))
        print("\n\n debug: ")
        for p in debug:
          print(p)
        exit(-1)
```

```
if tf is None:
  tf = self.food[0]
if If is None:
  lf = self.food[-1]
dtf = mk_round(tf.x - ax)
dlf = mk_round(lf.x - ax)
dfood = dtf if dtf < dlf else dlf
if abs(
    dfood
) < self.col_const: # if the abs distance is smaller than the required collision const
  self.food.remove(tf if dtf < dlf else lf) # remove food</pre>
  self.agents[a].eat(FOOD_CONST) # eat food
  self.eat += 1 # update food statistic
  food_index -= 1
  debug_line += " food was consumed, new food index is now " + str(food_index)
debug.append(debug_line)
if a == 0:
  d1 = abs(ax - self.agents[-1].x)
  d1 = abs(ax - self.agents[a - 1].x)
  d2 = abs(ax - self.agents[a + 1].x)
  d2 = abs(ax - self.agents[0].x)
if d2 < d1:
    min_d = ax - self.agents[a + 1].x
  a_s = a + 1
  min_d = ax - self.agents[a - 1].x
  a_s = a - 1
if (abs(ax - self.agents[a_s].x) <</pre>
    self.col_const
  self.interactions += 1
  interact(self.agents[a], self.agents[a_s], self)
self.agents[a].move(
  self.agents[a].think(
```

```
dfood, min_d, self.agents[a_s]
    self.agents[a].age() # applying age effect
    if self.agents[a].energy < ENLB_CONST * self.agents[a].mass: # if the agent is sick, lose health
      self.agents[a].health -= ENL_CONST
    if self.agents[a].energy > ENGB_CONST * self.agents[a].mass: # if the agent is healthy, gain health
      self.agents[a].health += ENG_CONST
    if (random.random() < self.agents[a].breed_chance</pre>
        and self.agents[a].health > 0
        and self.agents[a].mass >= self.agents[a].final_mass): # determine if an agent will breed
      nk = self.agents[a].breed(self.id)
      if nk is not None: # if the agent did not die in childbirth, append the newborn to the sim
        self.breed += 1
        self.id += 1
        self.agents.append(nk)
    if self.agents[a].health < -1e-5: # if the agent's health is <=0, kill it
      self.kill += 1
      self.agents.remove(self.agents[a])
  return True
def stats(self) -> str:
  Get basic statistics
  return "breed:{} kill:{} eat:{}\n".format(
    self.breed, self.kill, sum(self.eat_OT)
  ) + "avg mass: {}\n".format(np.mean([
    a.mass for a in self.agents
  ])) + "avg speed: {}\n".format(np.mean([
    a.speed for a in self.agents
  ])) + "avg breed chance: {}\n".format(
    np.mean([a.breed_chance for a in self.agents
         )) + "avg breed mass divider: {}\n".format(
    np.mean([a.breed_mass_div for a in self.agents]))
def graph(self, info: str = None, output=("plt", "excel")) -> str:
```

```
compatible_out = ["plt", "excel", "spss"]
   e = False
   for ro in output:
     if ro not in compatible_out:
        print("WARNING, output format {} is not supported, it will be skipped".format(ro))
     print("We currently support " + str(compatible_out))
   if info is None:
     info = input("Enter additional information about the sim: ")
   titles = [
population size",
     self.number_of_agents_OT, self.mass_OT, self.eat_OT, self.iq_OT, self.iq_OT,
     self.breed_mass_div_OT, self.breed_chance_OT, self.fight_OT, self.help_OT, self.nothing_OT,
     self.relative_groups_OT, self.close_family_in_group_OT
   extention = "png"
   fn = "graphs-0.3/" + self.get_fn()
   os.mkdir(fn)
     if "plt" in output:
       if len(titles) != len(values):
          raise Exception("Error len of titles must match len of vars")
        fig, axs = plt.subplots(len(values), sharex='all', figsize=(20, 60))
        metadata = dict()
        for i in range(len(values)):
          axs[i].plot(self.i_OT, values[i], linewidth=0.25)
          axs[i].axes.set_ylim([0, max(values[i])])
          axs[i].set_ylabel(titles[i])
```

metadata["Final" + titles[i]] = values[i][-1]

axs[0].axes.set\_xlim([0, self.dataPoints])

```
axs[0].set_title(
          .format(len(self.agents), self.gcsteps, time.strftime("%D"), info,
              self.stats()), )
      axs[-1].set_xlabel("Number Of Data Points")
      plt.tight_layout()
      plt.autoscale()
      pltfn = fn + "/plt." + extention
      fig.savefig(pltfn, bbox_inches='tight') # save graph
      im = Image.open(pltfn)
      meta = PngImagePlugin.PngInfo()
      for x in metadata:
        meta.add_text(x, str(metadata[x]))
      im.save(pltfn, extention, pnginfo=meta)
    print("error in generating plt file")
  transposed_data = []
  for i in range(self.dataPoints):
    transposed_data.append([j[i] for j in values])
    if "excel" in output:
      if len(values[0]) > 1048576:
        print("to manny data points, skipping excel")
        wb = openpyxl.Workbook(write_only=True)
        sheet = wb.create_sheet()
        sheet.append(titles)
        for i in transposed_data:
          sheet.append(i)
        wb.save(fn + "/excel.xlsx")
    print("error in generating excel file")
 if "spss" in output:
    savFileName = fn + '/spss.sav'
    varNames = [i.replace(" ", "_") for i in titles]
    varTypes = dict()
    for t in varNames:
      varTypes[t] = 0
    with savReaderWriter.SavWriter(savFileName, varNames, varTypes) as writer:
      for i in range(self.dataPoints):
        writer.writerow(transposed_data[i])
  return os.getcwd() + "\\" + fn.replace("/", "\\");
def animate(self, steps, res_mult=5, fps=10, bitrate=20000, print_freq=10, data_point_freq: int = 10,
      gui: bool = False) -> str:
```

```
ims = []
fig = plt.figure(figsize=(res_mult, res_mult))
for i in range(steps):
  if not self.step():
  if i % data_point_freq == 0:
    self.update_stats()
  if i % print_freq == 0:
    self.progress(steps, i, gui)
  acu = self.size_factor / 25
  row = [0 \text{ for i in np.arange}(0, 1, acu)]
  for f in self.food:
      row[round(f.x / acu)] = 100
      if round(f.x / acu) > 1 / acu:
        row[-1] = 100
        row[0] = 100
  for a in self.agents:
      row[round(a.x / acu)] = 255
      if round(a.x / acu) > 1 / acu:
        row[-1] = 255
        row[0] = 255
  r = len(row) / (2 * math.pi)
  p = np.zeros((math.ceil(2 * r) + 10, math.ceil(2 * r) + 10))
```

```
dist_proj = 2 * math.pi / len(row)
  angle = 0
   x = round(r * math.cos(angle) + r)
   y = round(r * math.sin(angle) + r)
    p[y][x] = v
    p[y+1][x] = v
    p[y-1][x] = v
    p[y][x+1] = v
    p[y][x-1] = v
    angle += dist_proj
 ims.append([
    plt.imshow(p,
         interpolation='nearest',
         aspect='auto')
  acu = None
  dist_proj = None
  p = None
 angle = None
  gc.collect()
self.progress(steps, steps, gui)
ani = animation.ArtistAnimation(fig,
                interval=100.
                blit=False,
                repeat_delay=1000)
Writer = animation.writers['ffmpeg']
writer = Writer(fps=fps, metadata=dict(artist='Me'), bitrate=bitrate)
fn = "animations-0.1/" + self.get_fn() + '.mp4'
ani.save(fn, writer=writer)
# de-initialize variables:
fig = None
ani = None
writer = None
Writer = None
gc.collect()
```

## auto.py:

```
from model2 import Sim
import argparse
import math
parser = argparse.ArgumentParser(description='Run the default simulation')
parser.add_argument("-s", type=int, dest="steps",
parser.add_argument("-p", type=int, dest="pop",
parser.add_argument("-f", type=int, dest="food",
          help='Account of food')
parser.add_argument("-a", help="enable animation", dest="animate", action='store_true')
parser.add_argument("-v", help="enable verbose", dest="v", action='store_true')
max e = 1048576
parser.add_argument("-dp", type=int, dest="dp",
            max_e) + ') if excel is used. Else, defaults to the number of steps', default=max_e)
parser.add_argument("--spss", help="output data in SPSS data format. Note that this will NOT force data
parser.add_argument("--no-excel",
           max_e),
         dest="no_excel", action='store_true')
parser.add_argument("--no-plt", help="Don't generate plt preview",
          dest="no_plt", action='store_true')
parser.add_argument("--gui", help="Used to output progress in json format for GUI (in beta)",
args = parser.parse_args()
ms = Sim(args.pop, args.food)
the maximum of 18277
# total_data_points = steps/data_point_freq
```

```
args.dp = min(args.dp, args.steps) # make sure that data points is smaller than steps
if args.dp is None:
 args.dp = args.steps if args.no_excel else min(args.steps, max_e)
data_point_freq = math.floor(args.steps / args.dp)
data_point_freq += 1 if data_point_freq == 0 else 0
if args.v:
  print("data point frequency selected {}".format(data_point_freq))
  print("expected data points: {}".format(args.steps / data_point_freq))
if args.animate:
 ms.animate(args.steps, data_point_freq=data_point_freq, gui=args.gui)
  ms.run(args.steps, max_attempts=-1, data_point_freq=data_point_freq, gui=args.gui)
req_formats = ()
if not args.no_plt:
 req_formats += ("plt",)
if not args.no_excel:
 req_formats += ("excel",)
if args.spss:
 req_formats += ("spss",)
print("Simulation complete. Requested files are stored at: " + ms.graph(info="generated via auto.py",
output=req_formats))
```

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