

Lab G6: Natural selection (simplified)

Emanuele Pietropaolo

Politecnico di Torino

Student id: s319501

emanuele.pietropaolo@studenti.polito.it

Abstract—Natural selection is a key mechanism in evolution, whereby a population evolves due to environmental factors that favour some random mutations over others. This paper presents a first approach to simulating such a process over a simplified population. Some assumptions have been made, such as the choice of model and the type of inheritance that the sons will have.

I. PROBLEM OVERVIEW

In nature, a population starts to evolve because of environmental factors. These factors affect the chances of an individual surviving. There can be many different variables that affect this chance, such as scarcity of resources, predators, disease, etc. Some individuals can develop resistance to one or more of these factors, improving their ability to survive and reproduce, and pass this resistance on to new generations.

II. PROPOSED APPROACH

Simulating such a complex process can be a challenging task. For this reason, some assumptions have been made:

A. Assumptions

- **Reproduction:** in nature there are many different forms of reproduction. Most require two individuals of the same species. In this simplified version, only asexual reproduction is considered, a form of reproduction that requires a single individual. This means that each individual can give birth to
- **Reproduction rate (λ):** each individual will reproduce with a specified rate of reproduction. In this simplified

version, is considered that all the individuals will share the same reproduction rate (λ) and that this rate is always the same no matter the age of the individual.

- **Reproduction process:** it is assumed that this process follows a Poisson process. In fact, each birth event can be considered as independent and occurring with a given frequency in time. The reproduction process of the whole population can be considered as following a Poisson distribution with a given $\lambda_{tot} = \sum_0^i \lambda_i$ where i = individual. This assumption with the previous one means that the process of the whole population follows a Poisson distribution with $\lambda_{tot} = \lambda \cdot n_{pop}$ where n_{pop} = size of the population.
- **Improvement over generation:** this is the process that models how resistance has been passed on to new generations. Each newborn receives a life expectancy from its parents, which is used to generate its lifetime variable. There is a possibility that this lifetime will be greater than the lifetime of the parent (p_i). In this case, the lifetime is randomly generated from a uniform distribution between the lifetime of the parent and the lifetime of the parent multiplied by a factor $(1 + \alpha)$. Otherwise it is generated from a uniform distribution between 0 and the lifetime of the parent.

III. RESULTS

A. Lifetime

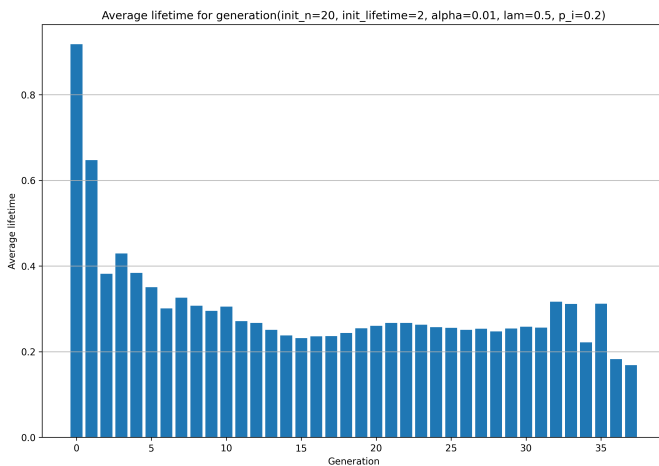


Fig. 1. Lifetime evolution with low p_i and low λ

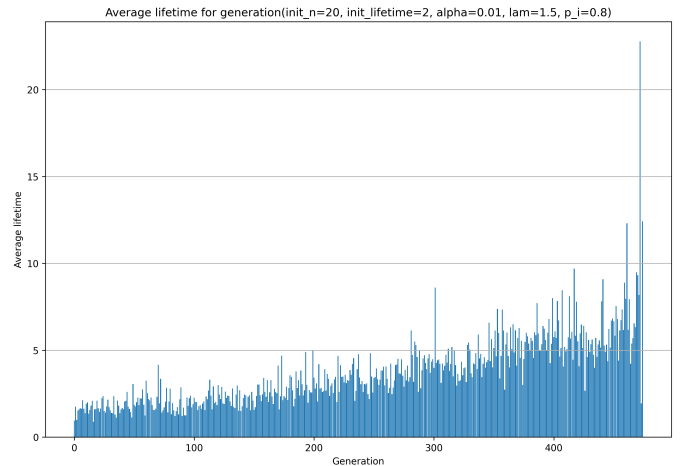


Fig. 2. Lifetime evolution with high p_i and low λ

In terms of lifetime, we can see in Fig. 1 that it starts to decrease when the probability of improvement (p_i) is low, while in Fig. 2 it increases otherwise.

B. Average number of children per generation

The average number of children is also an interesting measure, as it also depends on p_i . It can be seen in Fig. 3 that it starts to decrease with a low value of p_i , while it remain stable with a high value of p_i (Fig. 4).

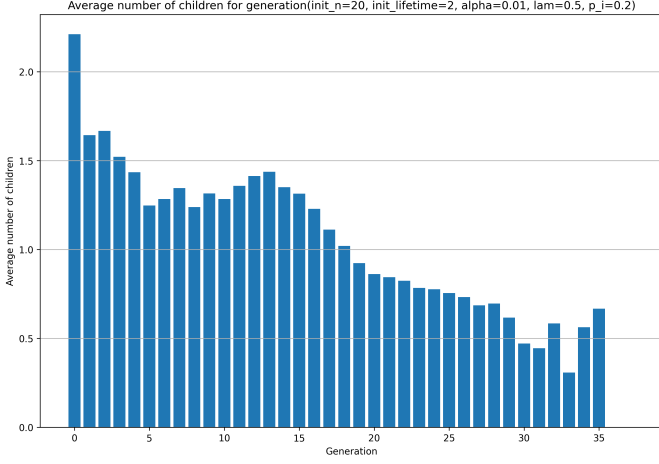


Fig. 3. Avg. number of children evolution with low p_i and low λ

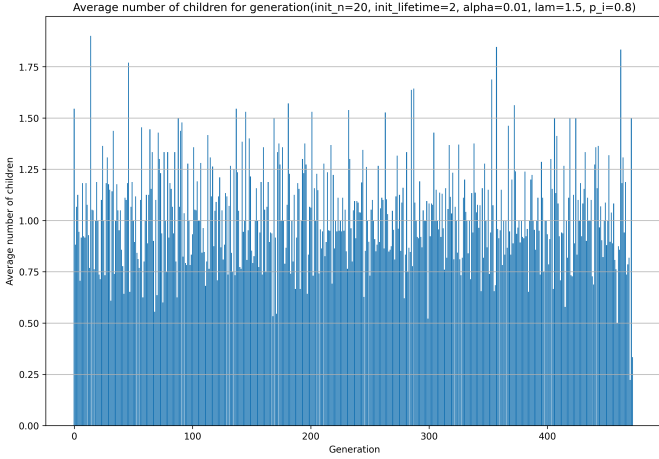


Fig. 4. Avg. number of children evolution with high p_i and low λ

C. Total number of children per generation

Another interesting measure is the total number of children per generation. In Fig. 6 it can be seen that there is a generation that has given birth to the highest number of children, and this happen in the configuration with the low value of p_i .

While Fig. 5 shows the configuration with an high value of p_i and it can be observed that the total number of children it doesn't show any tendency over time.

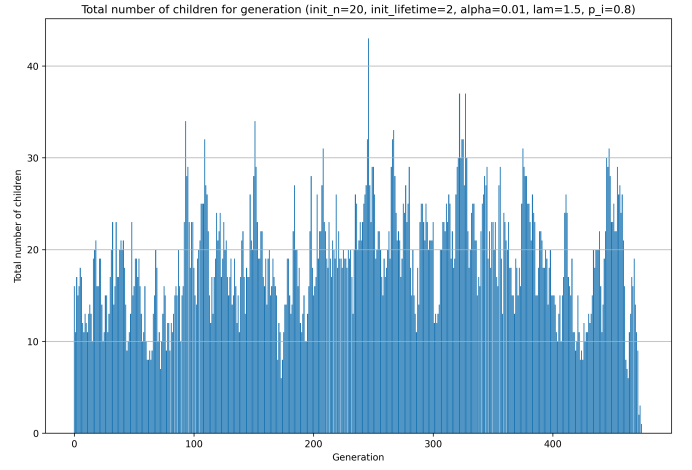


Fig. 5. Avg. number of children evolution with high p_i and low λ

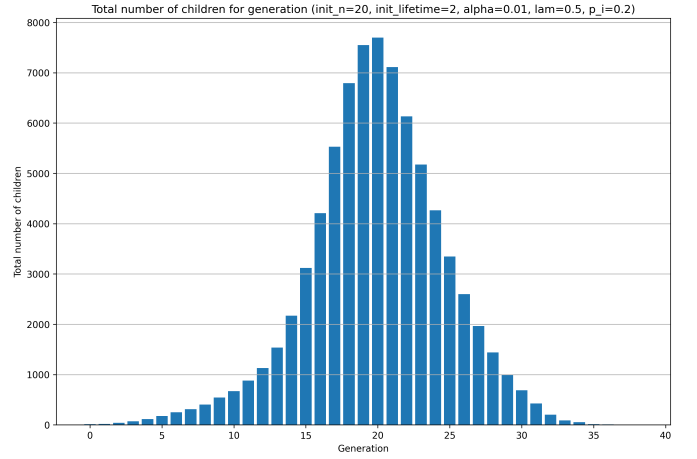


Fig. 6. Avg. number of children evolution with low p_i and low λ

IV. CONCLUSION

In conclusion, the proposed approach seems to provide some information about the model it wants to simulate. This will serve as baseline for a future approach.