

# Spontaneous Mutations in Population Dynamics and Phase Transitions

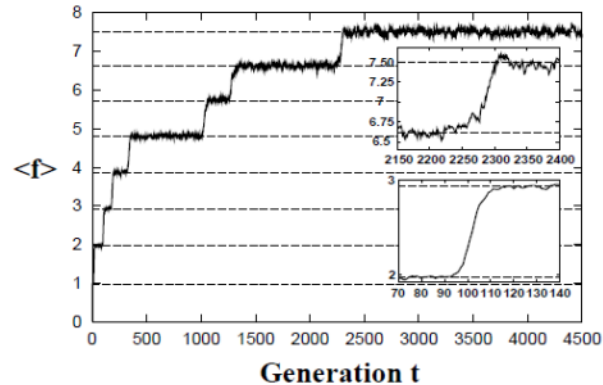
**Physics of Computation  
(Physics 256B, Spring 2022)**

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# Introduction and Previous models

- History of mutation-selection models
- Notion of exploration of the space of fitnesses through random mutations

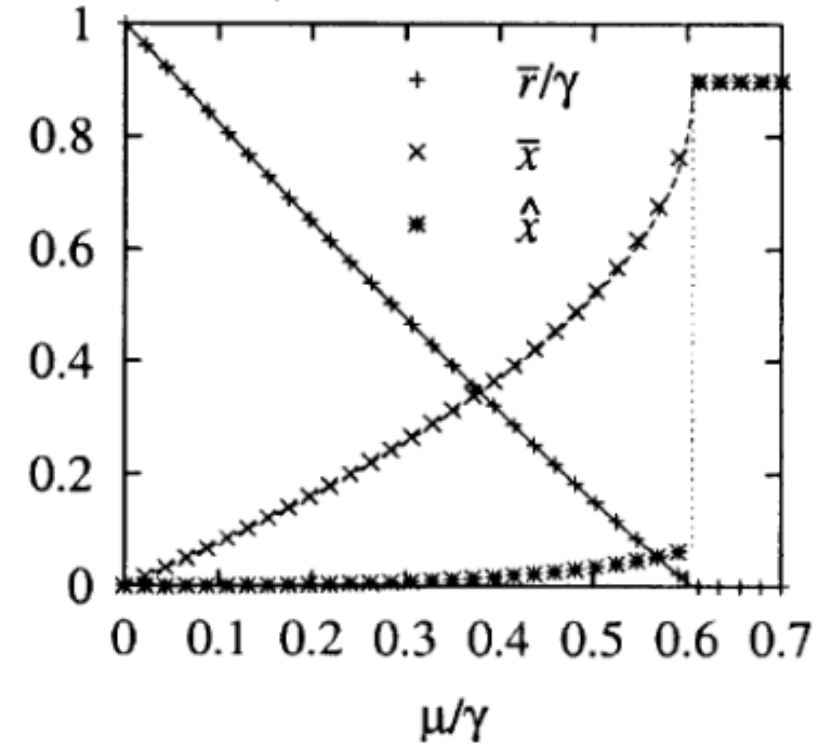


- Mean fitness increases in a jump pattern separated by stagnating periods
- Increasing the population size smoothens the increase in fitness, the stagnating periods are really metastable.

Erik van Nimwegen, James P. Crutchfield, and Melanie Mitchell, "Finite Populations Induce Metastability in Evolutionary Search", *Physics Letters A*, **229**:3 (1997) 144-150.

# Transition between order and disorder

- Influence of the mutation probability
- Notion of mutation threshold
  
- Above the threshold, 0 mean fitness and large fluctuations
- High mutation rate leads to disorder



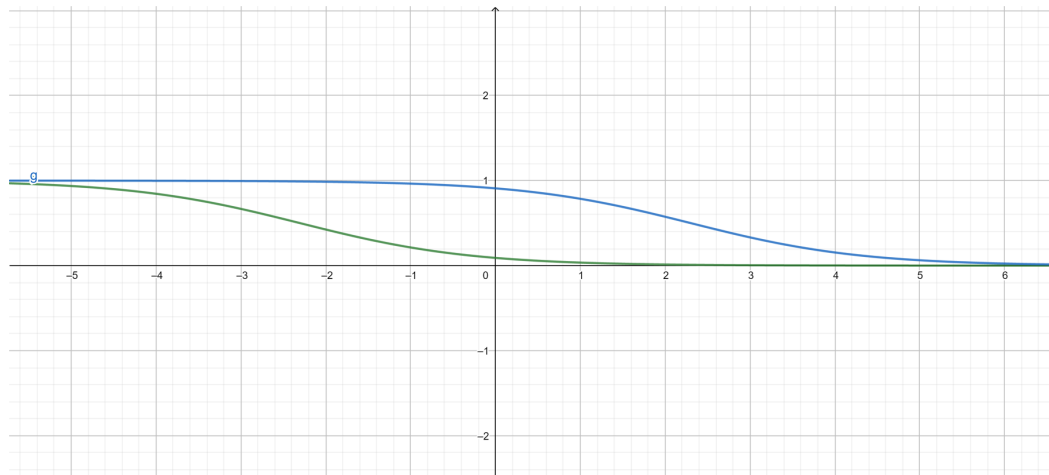
Hermisson, J., Redner, O., Wagner, H., & Baake, E. (2002). Mutation–selection balance: ancestry, load, and maximum principle. *Theoretical population biology*, 62(1), 9-46.

# Motivations and Goals

- Study population growth through mutation-selection
- Build a suited model with time-varying population size
- Influence of the mutation rate on the growth (or decay) of populations
- Can we relate disorder (entropy) and growth ?

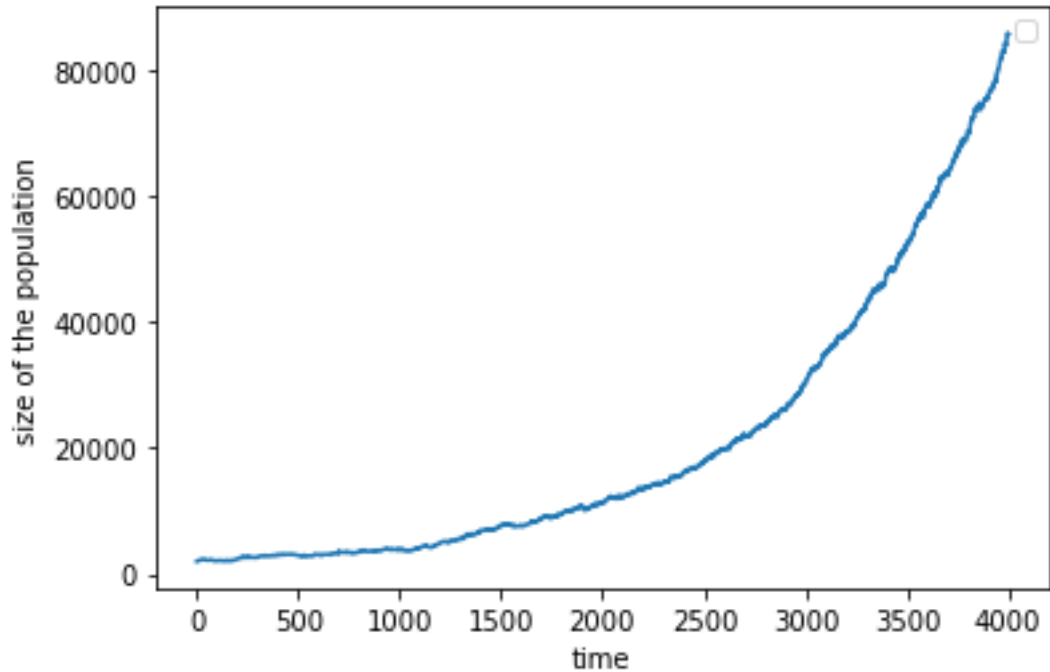
# The model

- Iterative mutation selection model
- Mutation step : Every individual has a fixed mutation rate. The new fitness is taken from a given distribution (2 choices). The parameters of the model quantify this step.
- Selection step : 3 outcomes possible for each individual : death, reproduction (actually cloning), or neither of the two.



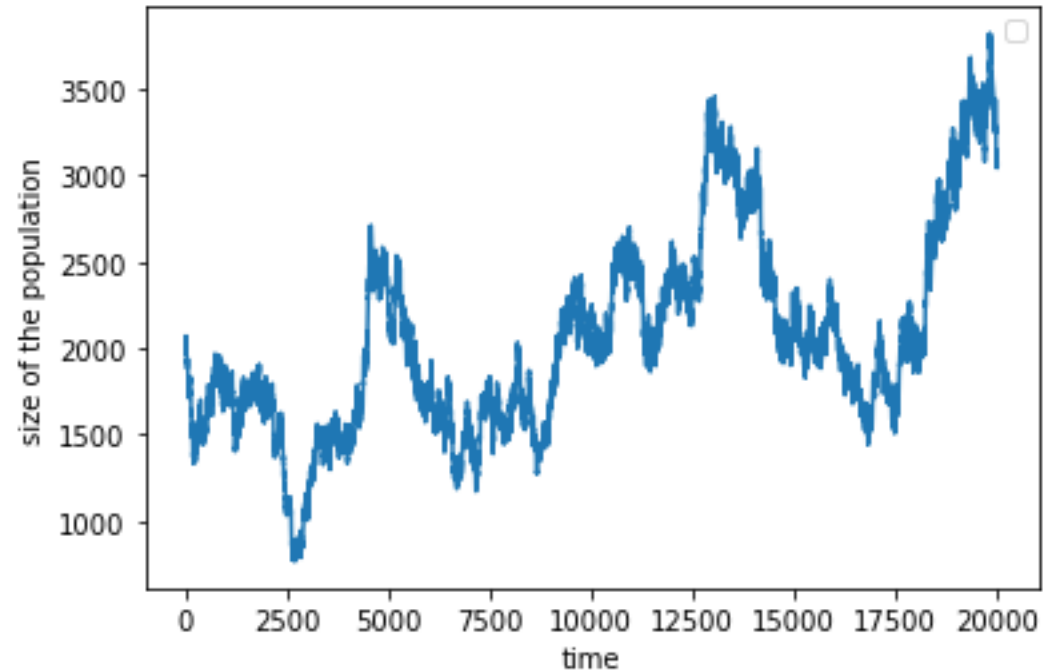
The probabilities for the selection step depend on the gap between the fitness of the individual and the mean fitness

# First results with correlation, some possible behaviours



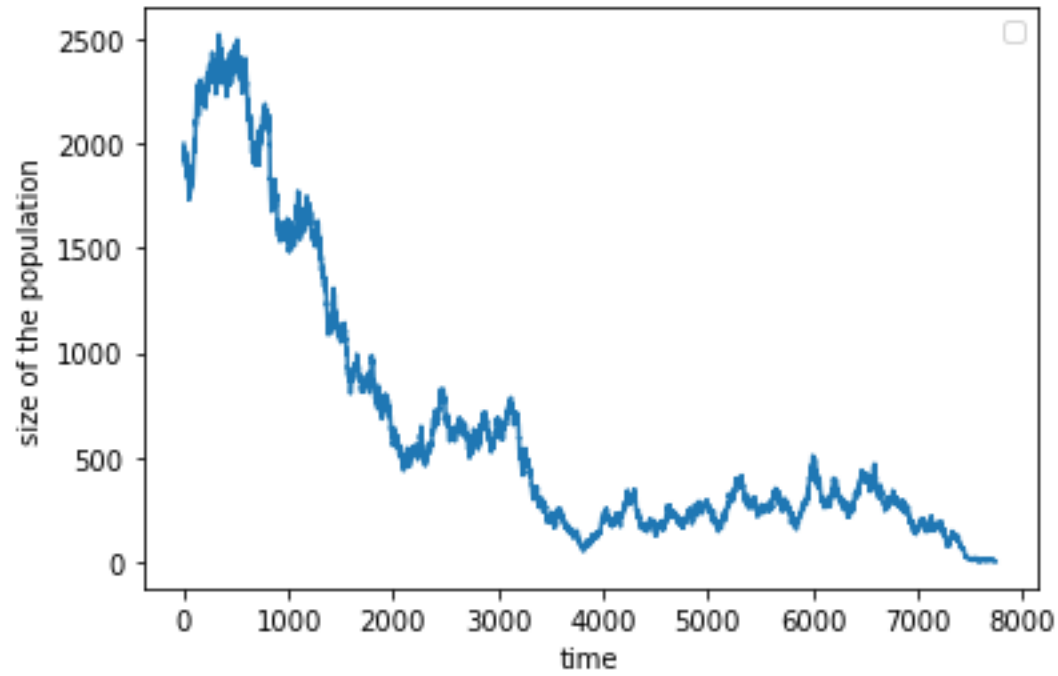
$N = 2000, p = 0.05, \sigma = 1$

Exponential growth  
This leads to the second choice  
for the mutation step



$N = 2000, p = 10^{-6}, \sigma = 1$

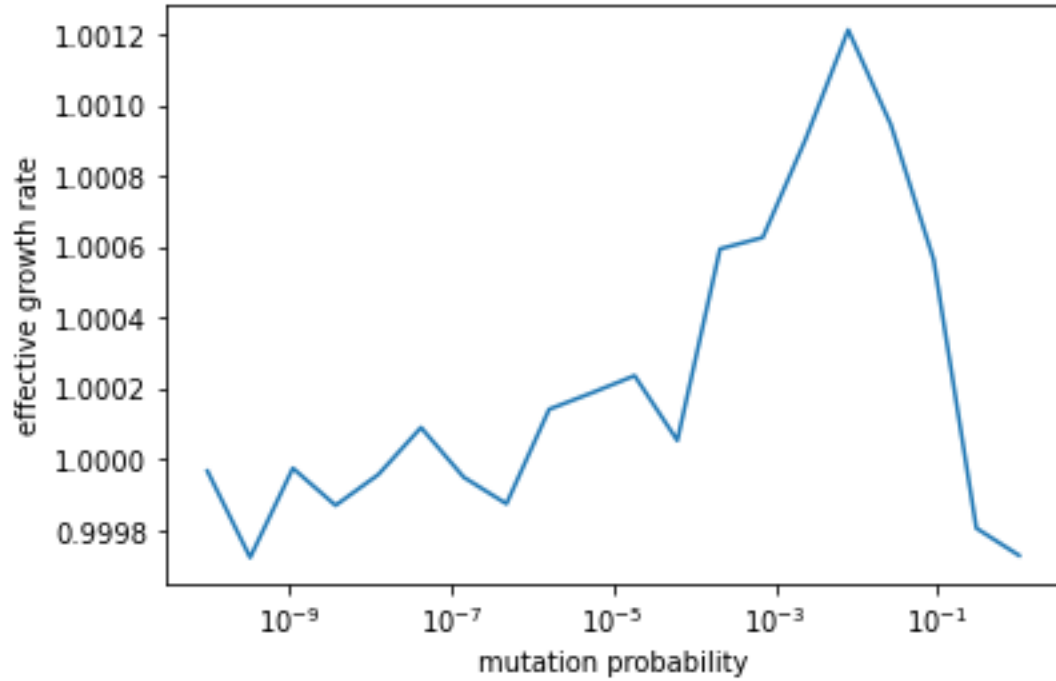
Low mutation rate, no exploration and  
limited growth



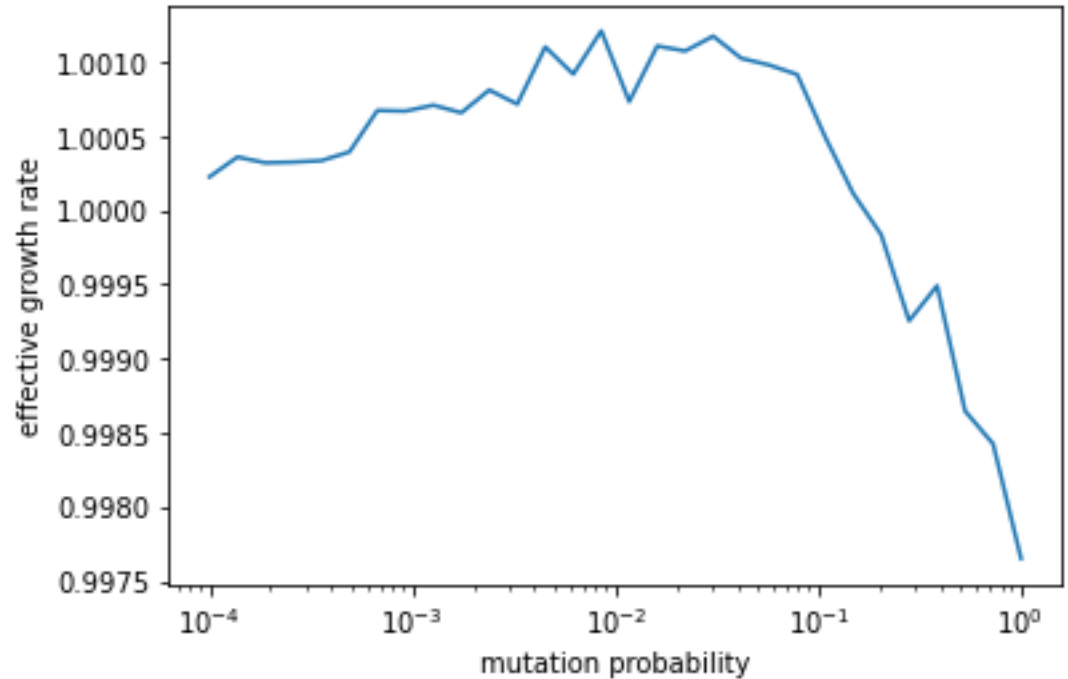
$N = 2000, p = 0.3, \sigma = 1$

High enough mutation rate leads to the decay of the population and long term extinction.

# Optimum for population growth

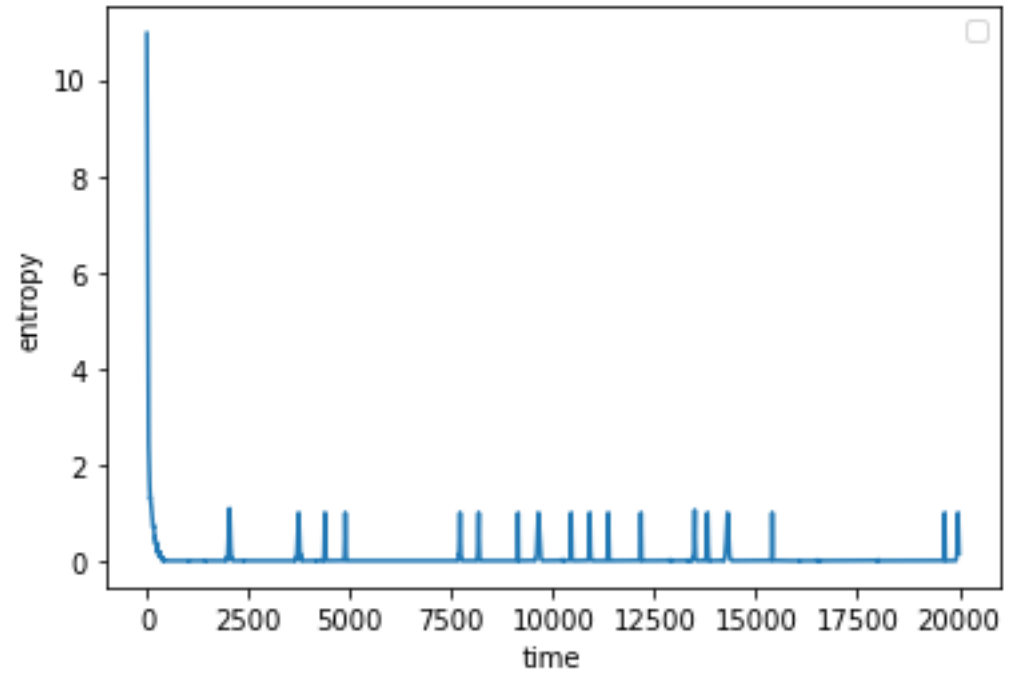
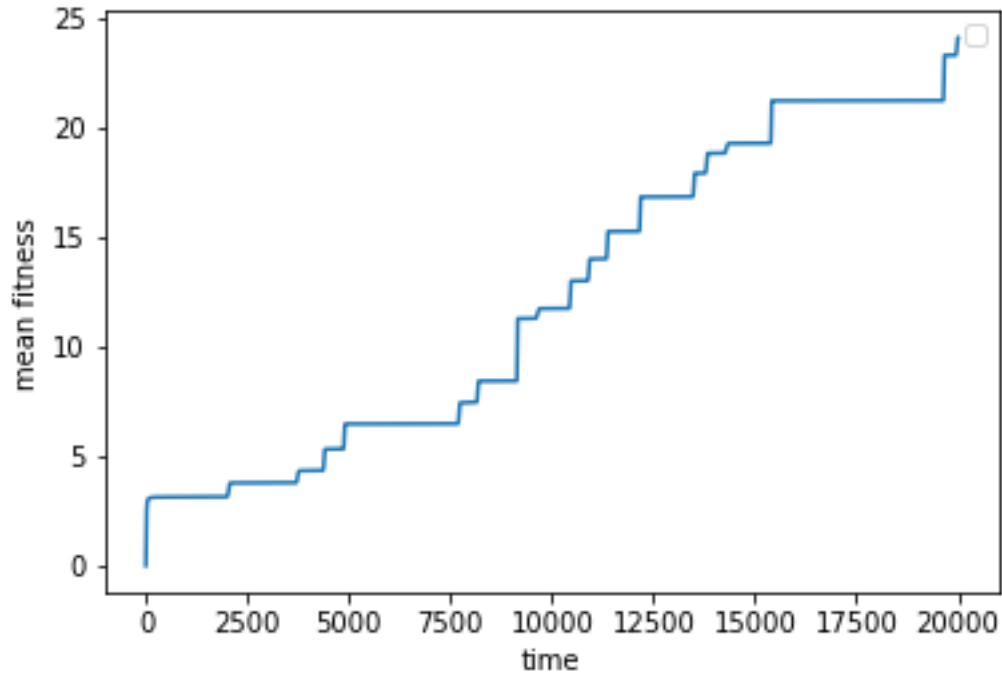


The optimal growth seems to be obtained for a mutation rate close to 0.01





# A stepwise increase in fitness

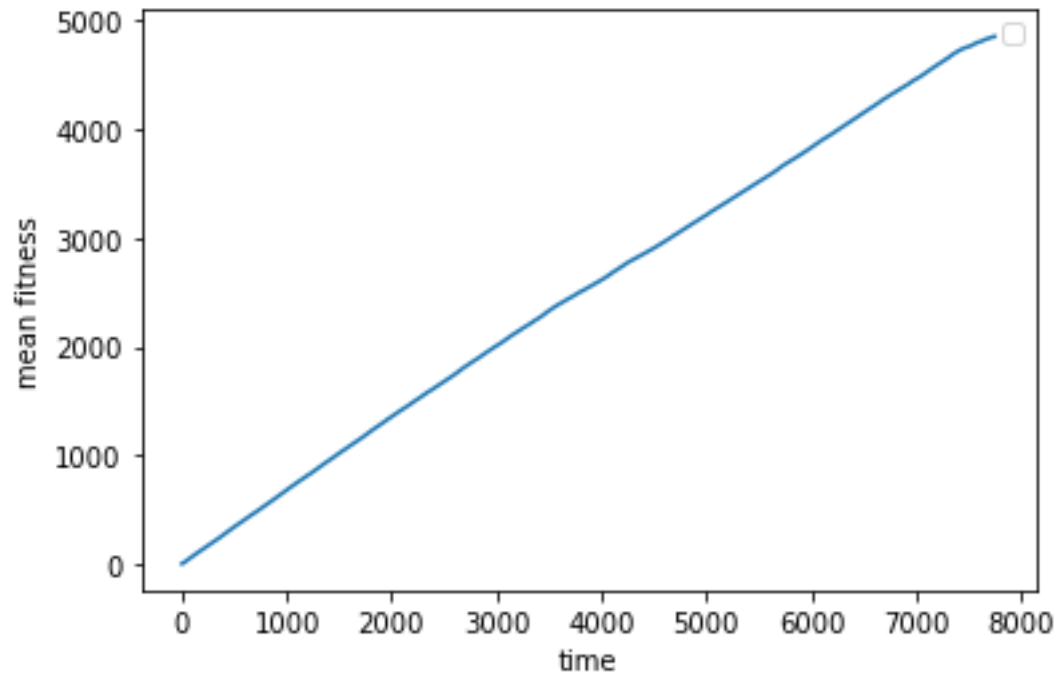


We recover this behaviour for  $p \ll 10^{-5}$   
Here,  $p = 10^{-6}$  and all parameters stay the same.

Each instance of steep growth is visible by a brief increase in entropy. When we are close to perfect homogeneity, we have a correlation between entropy and growth

# Some issues with this first version of the model

- Exponential growth is not realistic, even if this concerns a small region of the parameters
- No stationary regime, the average fitness increases forever

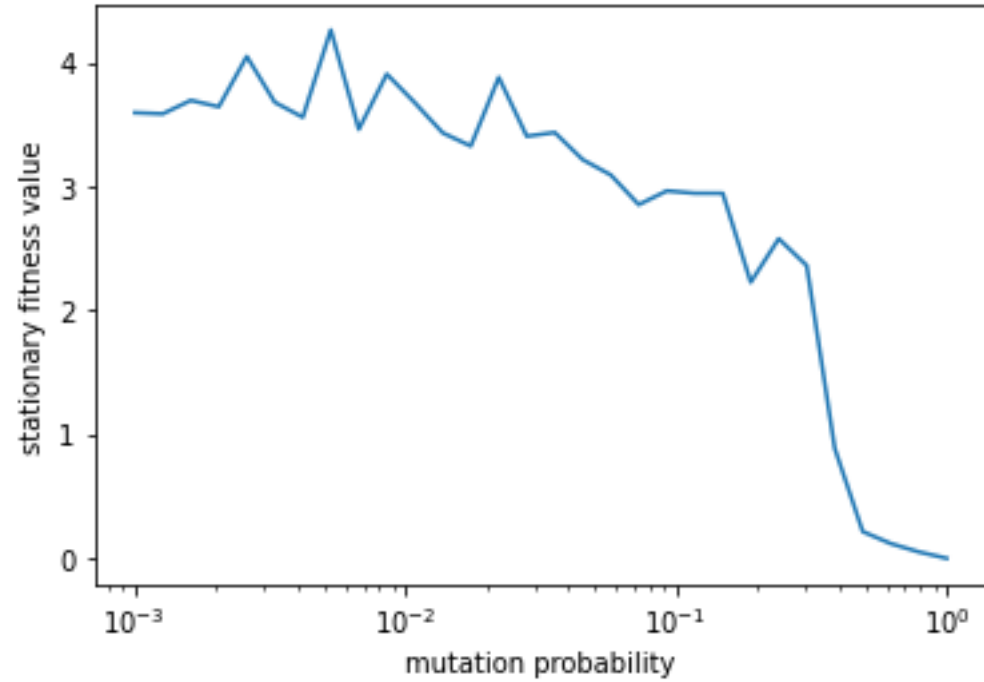


This actually corresponds to a decaying population ( $p=0.3$ ), despite the very high fitness.

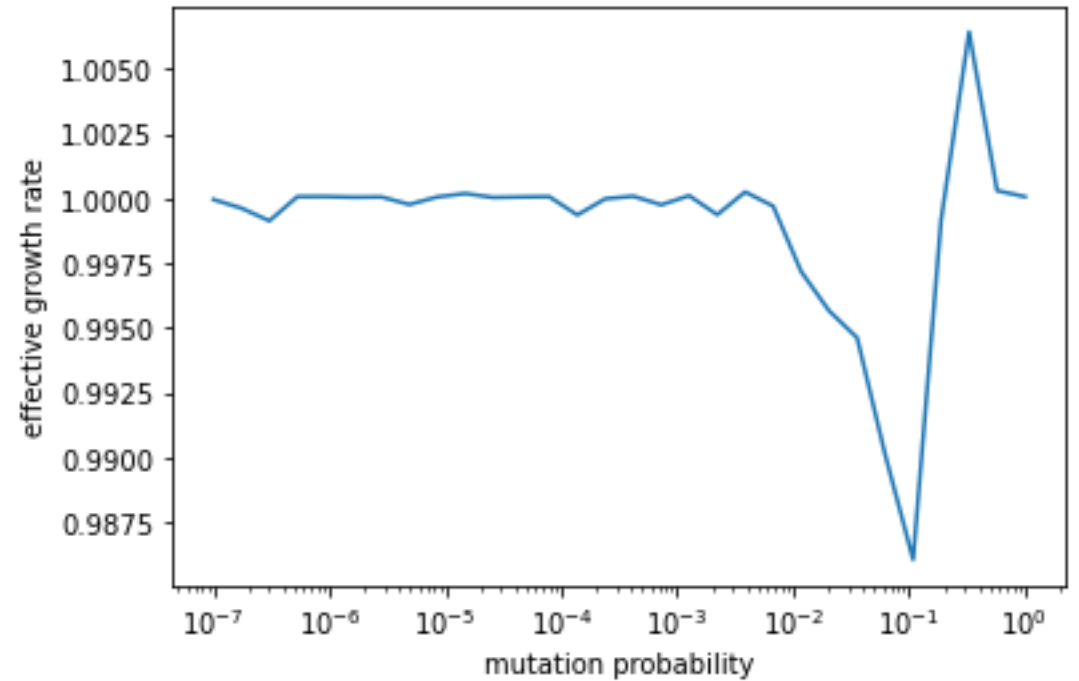
# Suppressing correlations

- The new fitness is taken around 0 instead of the previous fitness.
- This especially reduces the probability of a mutation increasing the fitness if the previous fitness was too high. More realistic
- For example,  $P_r(1) = 0.16$  instead of 0.5 in the previous model.
- We also have  $P_r(+\infty) = 0$  : once the fitness is too high, we cannot improve it anymore.

# Equilibrium fitness



And More ...



- References :

- Van Nimwegen, E., Crutchfield, J. P., & Mitchell, M. (1997). Finite populations induce metastability in evolutionary search. *Physics letters A*, 229(3), 144-150.
- Hermisson, J., Redner, O., Wagner, H., & Baake, E. (2002). Mutation–selection balance: ancestry, load, and maximum principle. *Theoretical population biology*, 62(1), 9-46.
- Baake, E., & Wagner, H. (2001). Mutation–selection models solved exactly with methods of statistical mechanics. *Genetics Research*, 78(1), 93-117.
- Manrubia, S. C., Domingo, E., & Lázaro, E. (2010). Pathways to extinction: beyond the error threshold. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 365(1548), 1943-1952.