Competition and space structure in an efficient population genetics model of carcinogenesis

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University of Trieste

Introduction

Original paper

Title: Efficient simulation under a population genetics model of

carcinogenesis

Authors: Zhu Tianqi, Hu Yucheng, Ma Zhi-Ming, Zhang De-Xing, Li

Tiejun, Yang Ziheng

Year: 2011

Journal: Bioinformatics

Volume: 27

Number: 6

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Genotypic space

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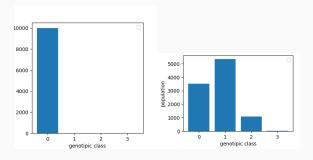


Figure 1: example of population distribution in the genotypic space at t=0 (left) and t= waiting time (right).

· Fixed population.

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- Fitness landscape: $f_i = 1.01^i$.

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- Fitness landscape: $f_i = 1.01^i$.
- Constant and uniform mutation rate (μ).
- · Possible events:
 - 1. a type j cell is **replaced** by a type j' one, with rate:

$$a_{jj'} = \frac{x_j f_{j'} x_{j'}}{\sum_{l=0}^{m-1} f_l x_l};$$

2. a type j cell **mutates** into a type j + 1 one, with rate:

$$a_j = \mu x_j$$
.

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· Variable population.

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- · Possible events:
 - 1. a type *j* cell is **born**, with rate:

$$a_j^b = \frac{\tilde{N}(t)f_jx_j}{\sum_{l=0}^{m-1}f_lx_l},$$

where $\tilde{N}(t)$ is a parameter capable of controlling the population size:

2. a type *j* cell **dies**, with rate:

$$a_j^d = x_j;$$

3. a type j cell **mutates** into a type j + 1 one, with rate:

$$a_j^m = \mu x_j$$
.

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Exact algorithm

Pseudocode:

- 1. Compute events rates.
- Apply Gillespie algorithm to sample time to next event e and next event.
- 3. Update time to t' = t + e and system's state.
- 4. If population of the m-th genotypic class is different from zero, set waiting time to t' and stop, otherwise go back to step 1.

Hybrid algorithm

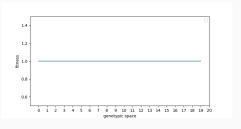
Pseudocode:

- 1. Compute events rates.
- 2. Partition events in critical and non-critical ones.
- 3. Determine tau-leaping step length τ .
- 4. Determine (by Gillespie's method) time to next critical event e.
- 5. If $e < \tau$, update system's state according to Gillespie algorithm, otherwise go directly to next step.
- 6. Let $h = min(e, \tau)$; update system's state applying tau-leaping algorithm over time h.
- 7. Update time to t' = t + h.
- 8. If population of the m-th genotypic class is different from zero, set waiting time to t' and stop, otherwise go back to step 1.

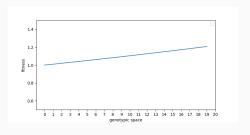
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My humble contribute

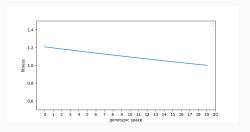
1. Flat: $f_i = 1$



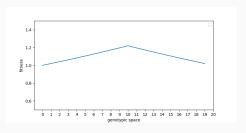
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- 4. Static 'mountain': $f_i = 1.01^{m-|m-2i|}$



Modelling competition

Two **assumptions**:

- the closer two organisms in the phenotypic space, the stronger the competition for resources
- the genotype is the only factor determining the phenotype

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- · the genotype is the only factor determining the phenotype

Result:

The more 'crowded' a portion of the genotypic space, the lower the fitness of the organisms in that region.

- 1. Flat: $f_i = 1$
- 2. Static increasing: $f_i = 1.01^i$
- 3. Static decreasing: $f_i = 1.01^{m-i-1}$
- 4. Static 'mountain': $f_i = 1.01^{m-|m-2i|}$
- 5. Static 'mountain' + dynamic: $f_i = 1.01^{m |m-2i|} \sum_{j=0}^{m-1} \frac{x_j}{N} \frac{|i-j|}{m}$
- 6. static decreasing + dynamic: $f_i = 1.01^{m-i-1} \sum_{j=0}^{m-1} \frac{x_j}{N} \frac{|i-j|}{m}$

Spatial structure (naive)

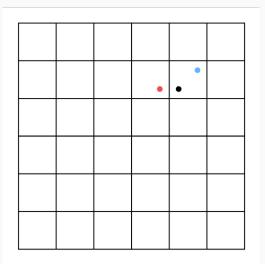


Figure 2: 'world' of the spatial model with resolution=16.

Spatial structure (less naive)

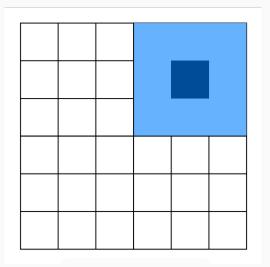


Figure 3: 'world' of the spatial model with resolution=16.

Spatial structure (less naive)

Three possible events:

1. a type *j* cell is born, with rate:

$$a_{j}^{b} = \frac{\tilde{N}(t)}{4} \frac{f_{j} X_{j}'}{\sum_{l=0}^{m-1} f_{l} X_{l}'}, X_{i}' = X_{i} + \frac{1}{4} \sum_{\{ngb\}} X_{k}$$

where *res* is the resolution and #ngb is the number of neighbour areas;

2. a type *j* cell dies, with rate:

$$a_j^d = \frac{N}{N + \frac{1}{4} \sum_{\{ngb\}} N_k} X_j';$$

3. a type j cell mutates into a type j + 1 one, with rate:

$$a_j^m = \frac{N}{N + \frac{1}{4} \sum_{\{nqb\}} N_k} \mu x_j'.$$

Spatial structure (less naive)

Time to next event was **heuristically** fixed by taking at each iteration:

$$\tilde{e} = \frac{\min\left(\{e\}_{areas}\right) + \max\left(\{e\}_{areas}\right)}{2},$$

where $\{e\}_{areas}$ is the set of times to next critical event computed on each area.

Simulation results

Reproducing results of the original paper

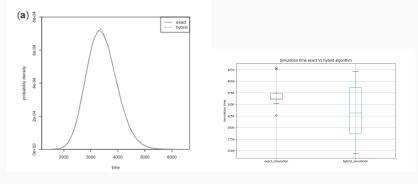


Figure 4: comparison of waiting time obtained with fixed-population exact and hybrid algorithms in original paper (left) and our work (right); parameters: $N = 10^3$, m = 20, $\mu = 10^{-3}$, fitness: static increasing.

Reproducing results of the original paper

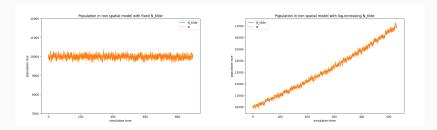


Figure 5: behaviour of *N* and *N_tilde* in time in our dynamic-population model with constant (**left**) and increasing (**right**) \tilde{N} ; parameters: $N = 10^4$, m = 4, $\mu = 10^{-4}$, fitness: static increasing.

Changing the fitness landscape

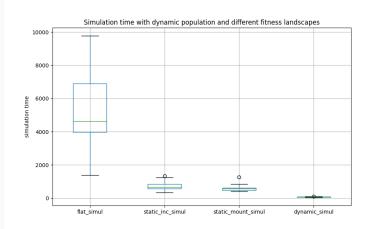


Figure 6: waiting time obtained with fixed-population hybrid model for different fitness landscapes; parameters: $N=10^4$, m=4, $\mu=10^{-4}$; note: static *increasing* fitness did not converge within an acceptable time.

Changing the fitness landscape

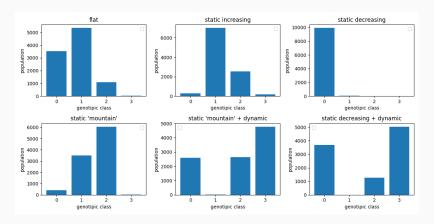


Figure 7: final state obtained with fixed-population hybrid model for different fitness landscapes; parameters: $N = 10^4$, m = 4, $\mu = 10^{-4}$.

Introducing space

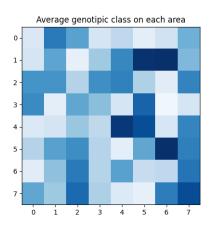


Figure 8: final distribution of the average genotypic class in space obtained with spatial model; parameters: $\tilde{N}=10^4$ (constant), m=4, resolution = 64, $\mu=10^{-4}$, fitness: static increasing.

Waiting time in spatial model

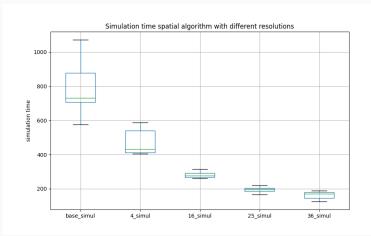


Figure 9: waiting time obtained with spatial model for different resolutions (from 1 to 36); parameters: $\tilde{N}=10^4$ (constant), m=4, $\mu=10^{-4}$, fitness: static increasing.

Population in spatial model

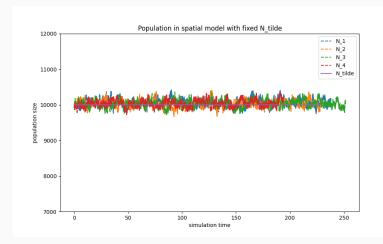
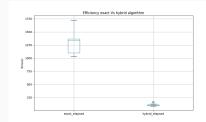


Figure 10: N's behaviour in time for four simulations of the same spatial model; parameters: $\tilde{N}=10^4$, m=4, $\mu=10^{-4}$, resolution = 16, fitness: static increasing.

An efficiency study



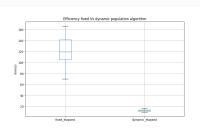


Figure 11: efficiency (in seconds) for fixed-population exact and hybrid models; parameters: $N=10^3$, m=20, $\mu=10^{-3}$, fitness: static increasing.

Figure 12: efficiency (in seconds) for fixed-population and dynamic-population hybrid models; parameters: $N\left(\tilde{N}\right)=10^3,\,m=20,\,\mu=10^{-3},\,\text{fitness: static increasing.}$

An efficiency study

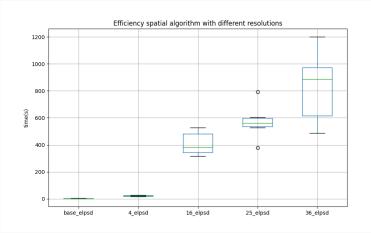


Figure 13: efficiency (in seconds) of spatial model for different resolutions (from 1 to 36); parameters: $\tilde{N}=10^4$ (constant), m=4, $\mu=10^{-4}$, fitness: static increasing.

Thank you!