

Primjena koncepta makrokariotipa u modeliranju evolucije stanica kvasca

Ina Mihalj

Mentor: Nenad Pavin





- biološka motivacija
- problem

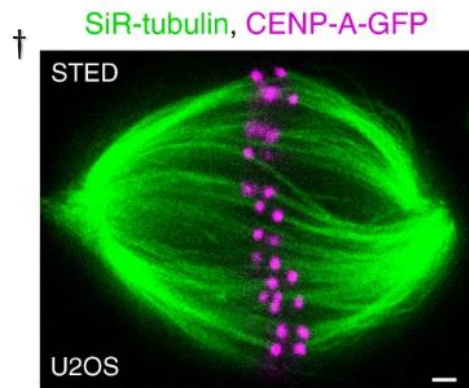
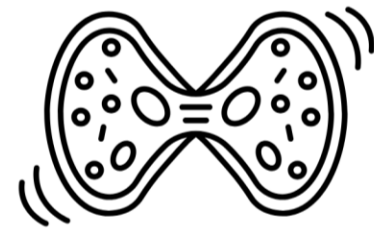


- koncept makrokariotipa
- Monte Carlo simulacija
- teorija srednjeg polja



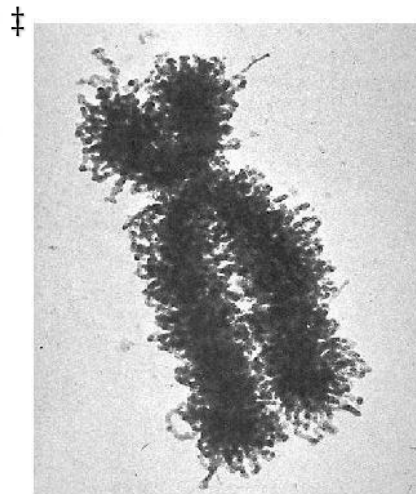
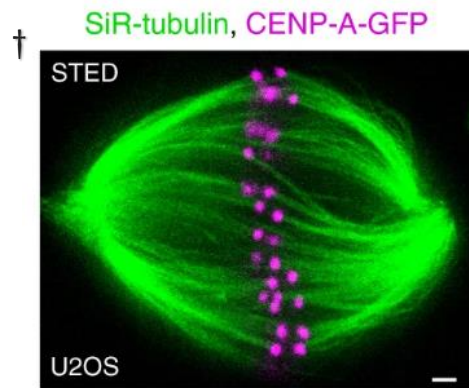
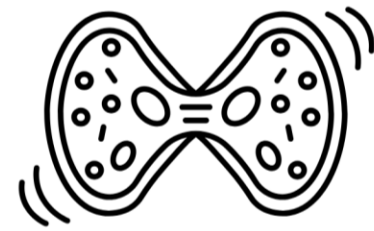
- rezultati
- zaključak

Dioba stanica



† Novak, M., Polak, B., Simunić, J. *et al.* The mitotic spindle is chiral due to torques within microtubule bundles. *Nat Commun* 9, 3571 (2018).

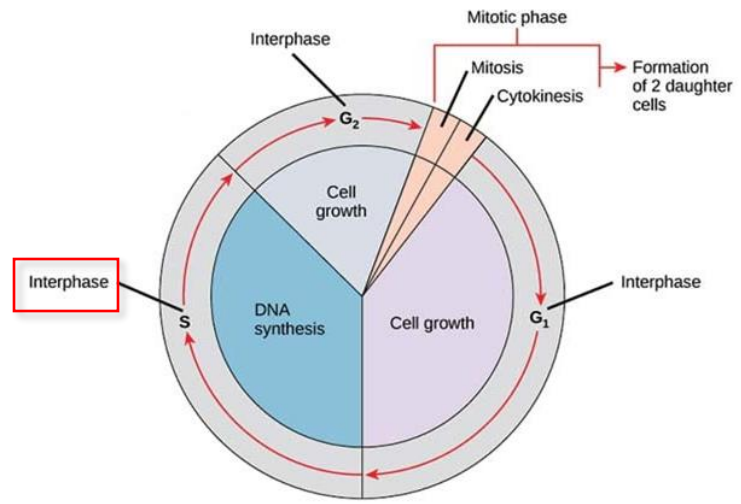
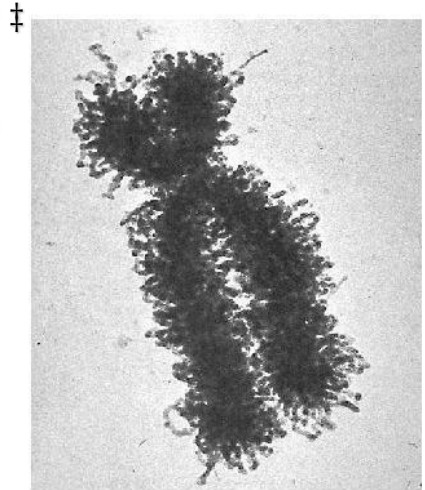
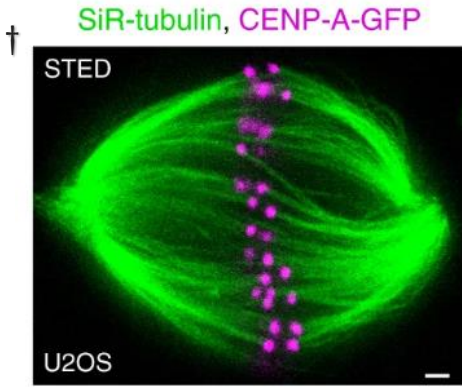
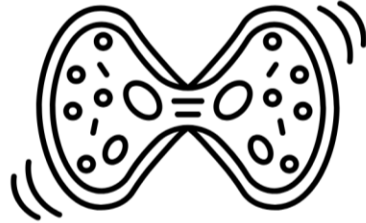
Dioba stanica



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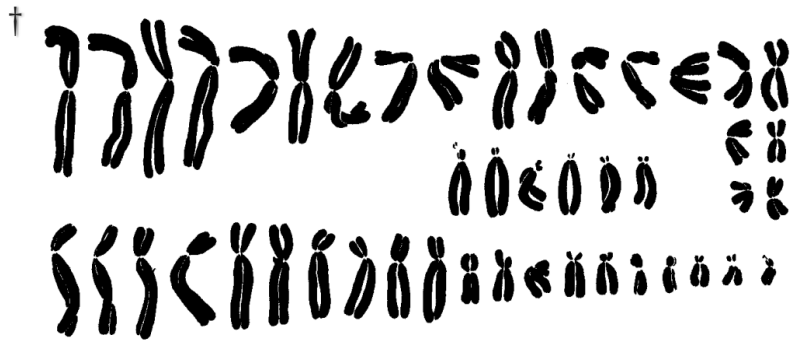
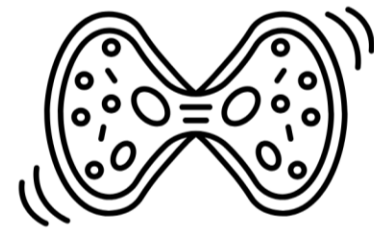
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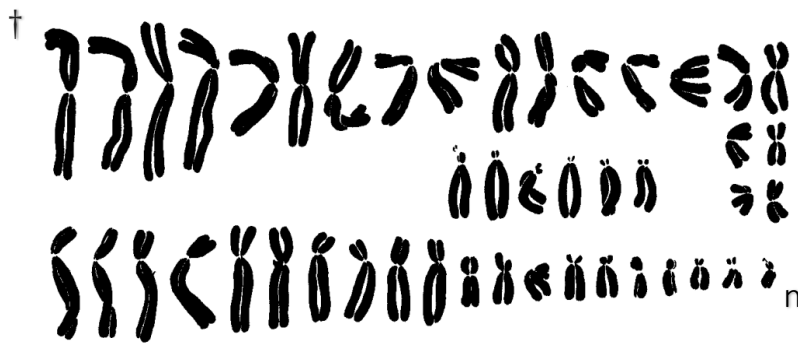
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Dioba stanica



† Tjio, J.H., Levan, A. (1956). The Chromosome Number of Man. In: Persaud, T.V.N. (eds) Problems of Birth Defects. Springer, Dordrecht.

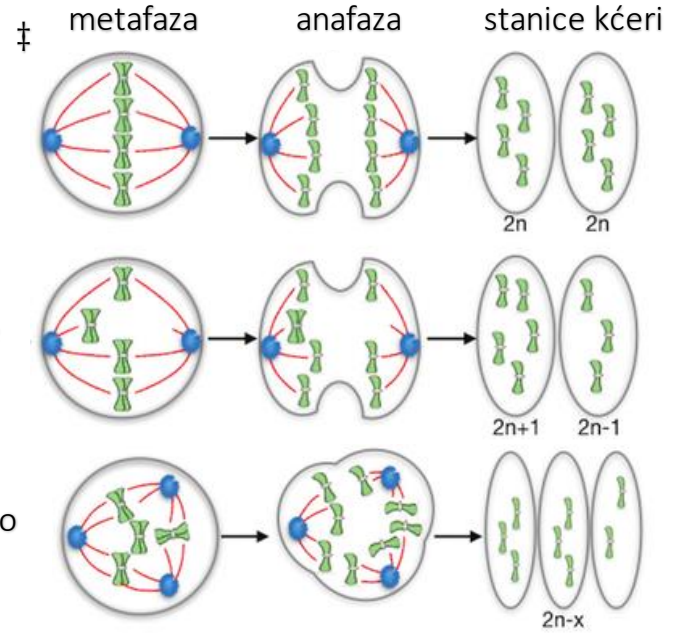
Dioba stanica



ispravna dioba

nerazdvajanje kromosoma

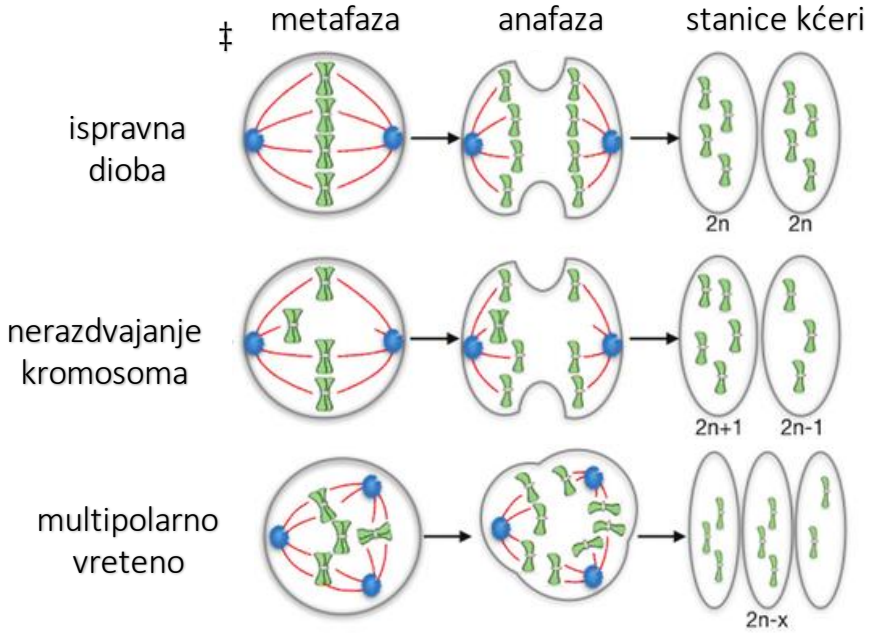
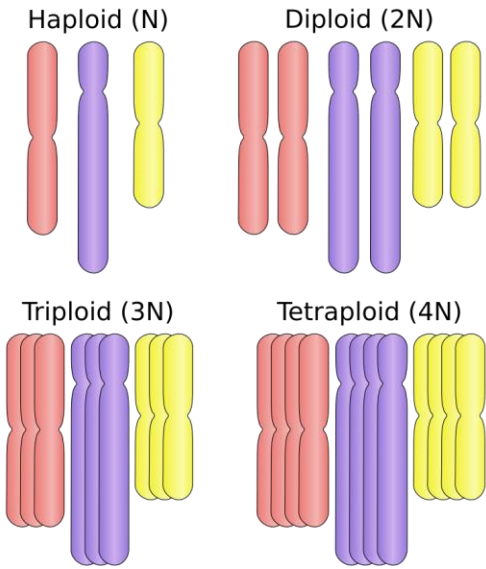
multipolarno vreteno



† Tjio, J.H., Levan, A. (1956). The Chromosome Number of Man. In: Persaud, T.V.N. (eds) Problems of Birth Defects. Springer, Dordrecht.

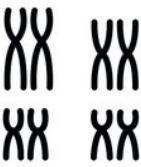
‡ Fusco, Pina & Esposito, Maria & Tonini, Gian Paolo. (2018). Chromosome instability in neuroblastoma (Review). Oncology Letters. 16.

Dioba stanica



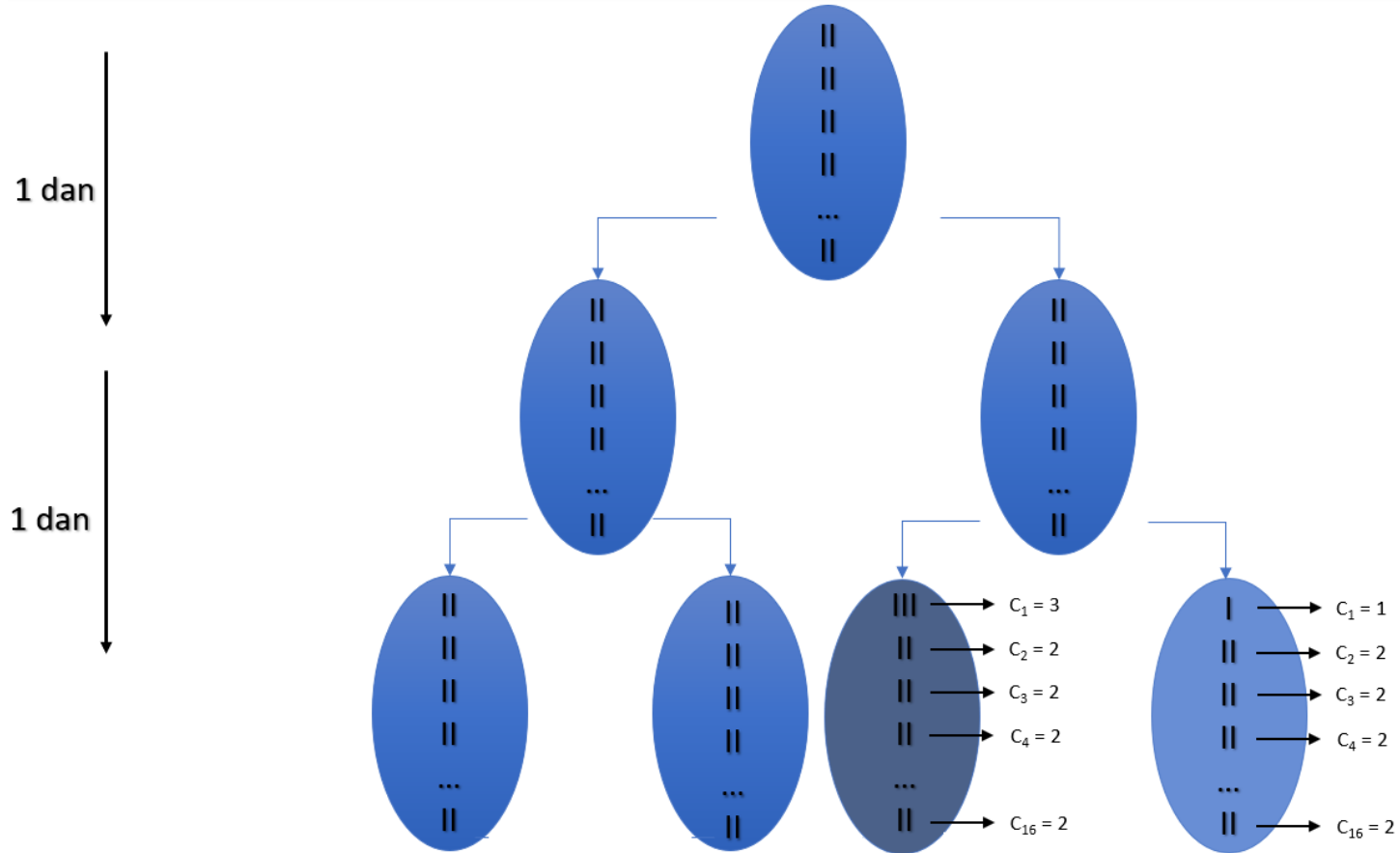
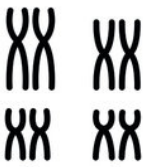
‡ Fusco, Pina & Esposito, Maria & Tonini, Gian Paolo. (2018). Chromosome instability in neuroblastoma (Review). Oncology Letters. 16.

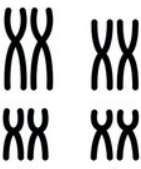
Teorijski model za evoluciju kariotipa



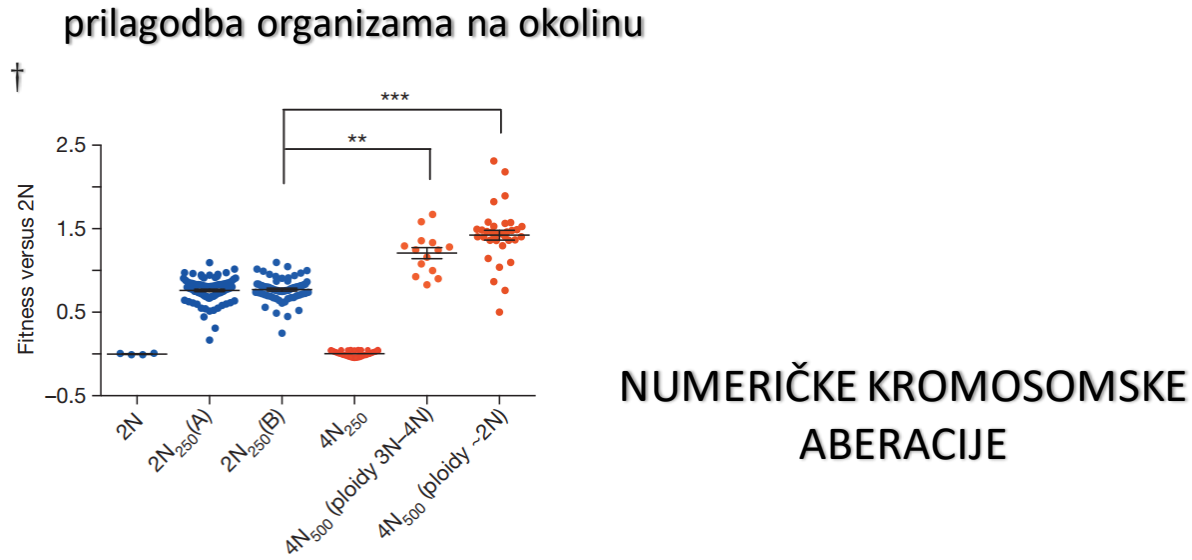
NUMERIČKE KROMOSOMSKE
ABERACIJE

Teorijski model za evoluciju kariotipa

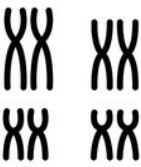




Teorijski model za evoluciju kariotipa



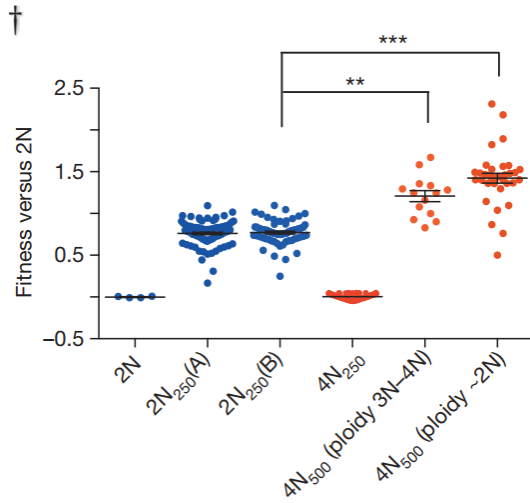
† Selmecki, A., Maruvka, Y., Richmond, P. et al. Polyploidy can drive rapid adaptation in yeast. Nature 519, 349–352 (2015).



Teorijski model za evoluciju kariotipa

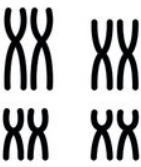
prilagodba organizama na okolinu

tumorska i predtumorska tkiva



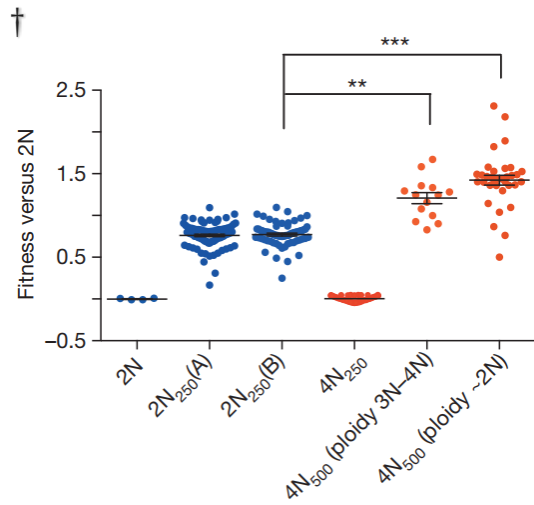
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Teorijski model za evoluciju kariotipa

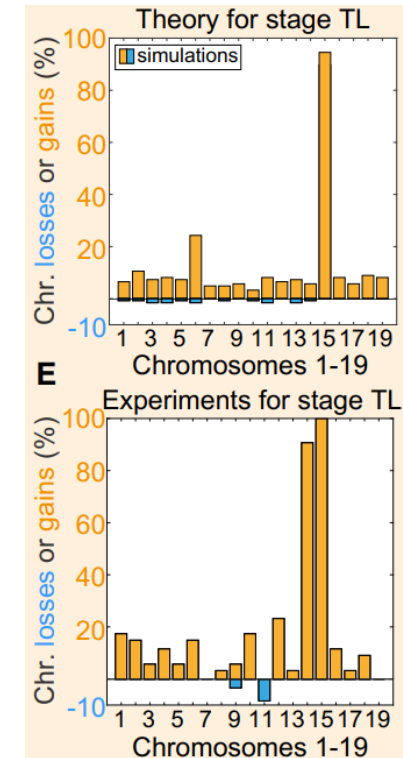
prilagodba organizama na okolinu



NUMERIČKE KROMOSOMSKE ABERACIJE

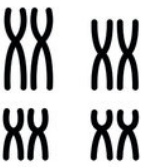
tumorska i predtumorska tkiva

‡



† Selmecki, A., Maruvka, Y., Richmond, P. et al. Polyploidy can drive rapid adaptation in yeast. *Nature* 519, 349–352 (2015).

‡ Ban I, Tomašić L, Trakala M, Tolić IM, Pavin N. Proliferative advantage of specific aneuploid cells drives evolution of tumor karyotypes. *Biophys J.* 2023 Feb 21;122(4):632-645.

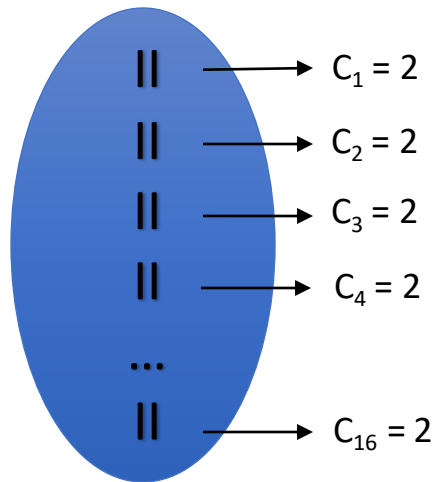


Teorijski model za evoluciju kariotipa

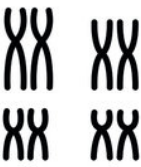
$$\vec{K} \equiv (c_1, c_2, \dots, c_n)$$



velik broj mogućih kombinacija
kariotipa u stanici



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Teorijski model za evoluciju kariotipa

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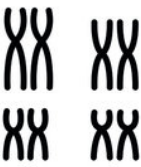
velik broj mogućih kombinacija
kariotipa u stanici



$$L = 6$$

$$n = 16$$

$$2.8 \times 10^{12}$$



Teorijski model za evoluciju kariotipa

$$\vec{K} \equiv (c_1, c_2, \dots, c_n)$$

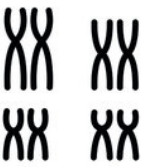


velik broj mogućih kombinacija
kariotipa u stanici

velik broj stanica u populaciji

$$\begin{array}{r} L = 6 \\ n = 16 \\ \hline 2.8 \times 10^{12} \end{array}$$

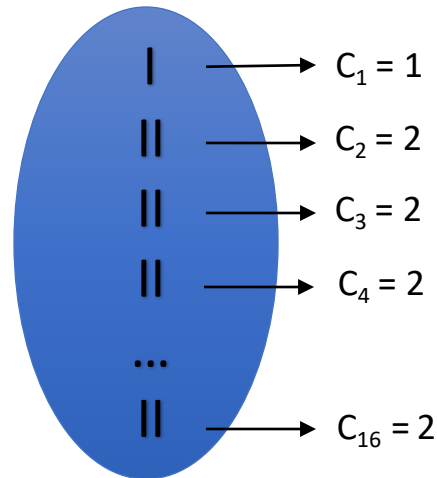
$$2^g$$

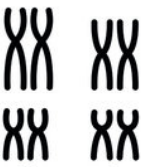


Koncept makrokariotipa

$$\vec{K} \equiv (c_1, c_2, \dots, c_n)$$

$$\vec{K} = (1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2)$$

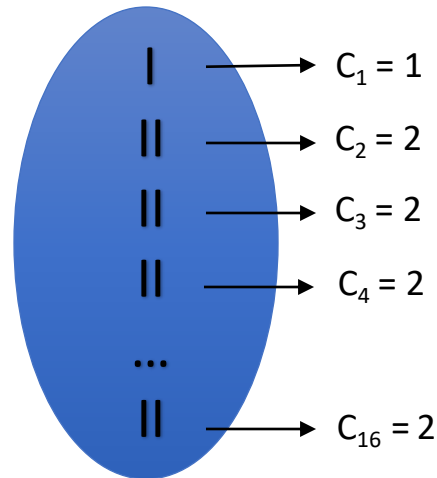




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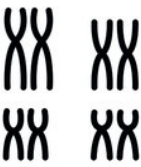
$$\vec{K} = (1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2)$$



$$\vec{M}(\vec{K}) \equiv (x_1, x_2, \dots, x_L)$$

$$\vec{M}(\vec{K}) = (1, 15)$$

$$x_1 + x_2 + \dots + x_L = n.$$



Koncept makrokariotipa

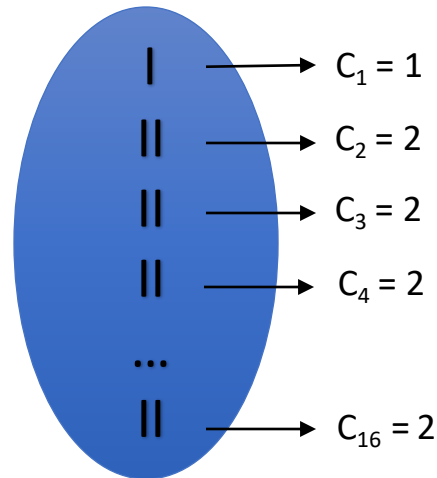
$$\vec{K} \equiv (c_1, c_2, \dots, c_n)$$

$$\vec{K} = (1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2)$$

$$L = 6$$

$$n = 16$$

$$4.6 \times 10^6$$



$$\vec{M}(\vec{K}) \equiv (x_1, x_2, \dots, x_L)$$

$$x_1 + x_2 + \dots + x_L = n.$$

$$\vec{M}(\vec{K}) = (1, 15)$$

Stohastički pristup evoluciji populacije stanica



Parametar	Oznaka	Definicija
Missegregacija kromosoma s 1 kopijom	p_{m_1}	$1 \times x_1 \times p_0$
Missegregacija kromosoma s 2 kopije	p_{m_2}	$2 \times x_2 \times p_0$
Ispravna stanična dioba	p_d	$1 - p_m - p_v$
Životni vijek stanice	t_0	$1 + 0.005 \times (x_1)^{1.5}$

$$p_0 = 0.0025$$

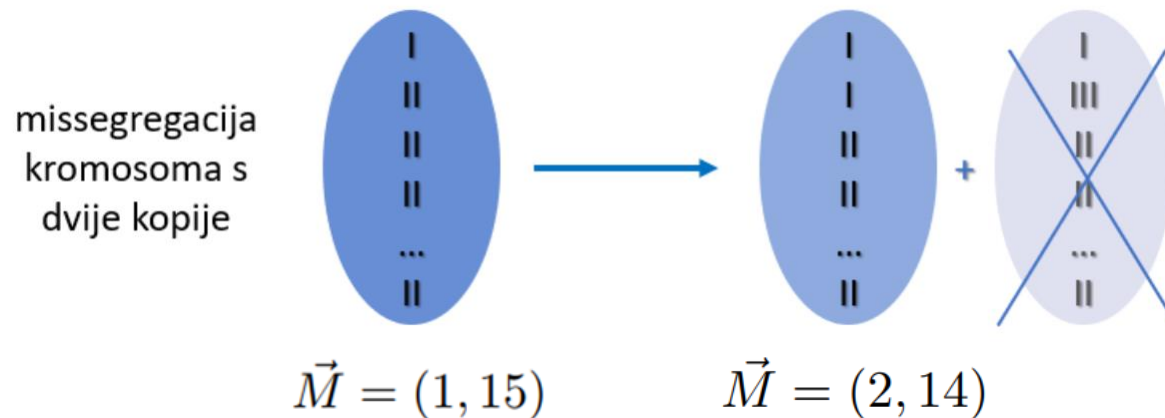
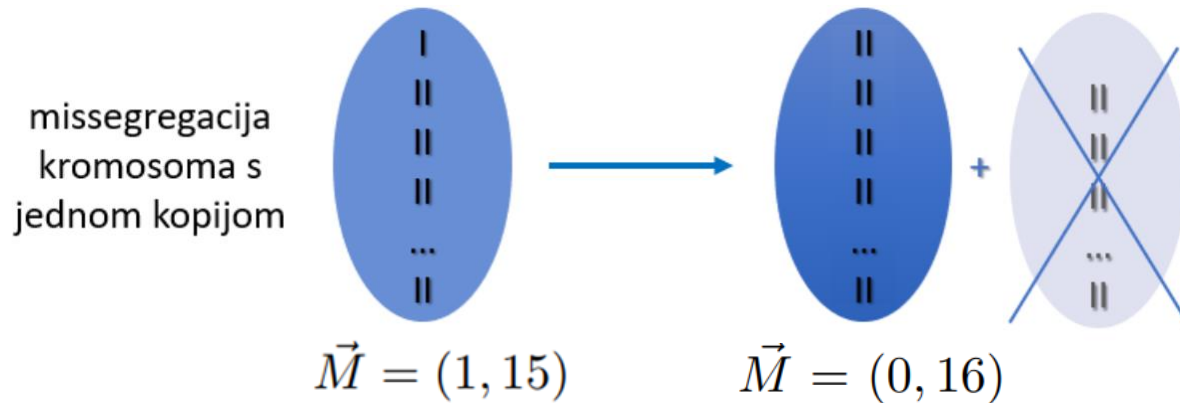


vrijeme početka života stanice

Stohastički pristup evoluciji populacije stanica



$$L = 2$$





Stohastički pristup evoluciji populacije stanica



NEDOSTATAK

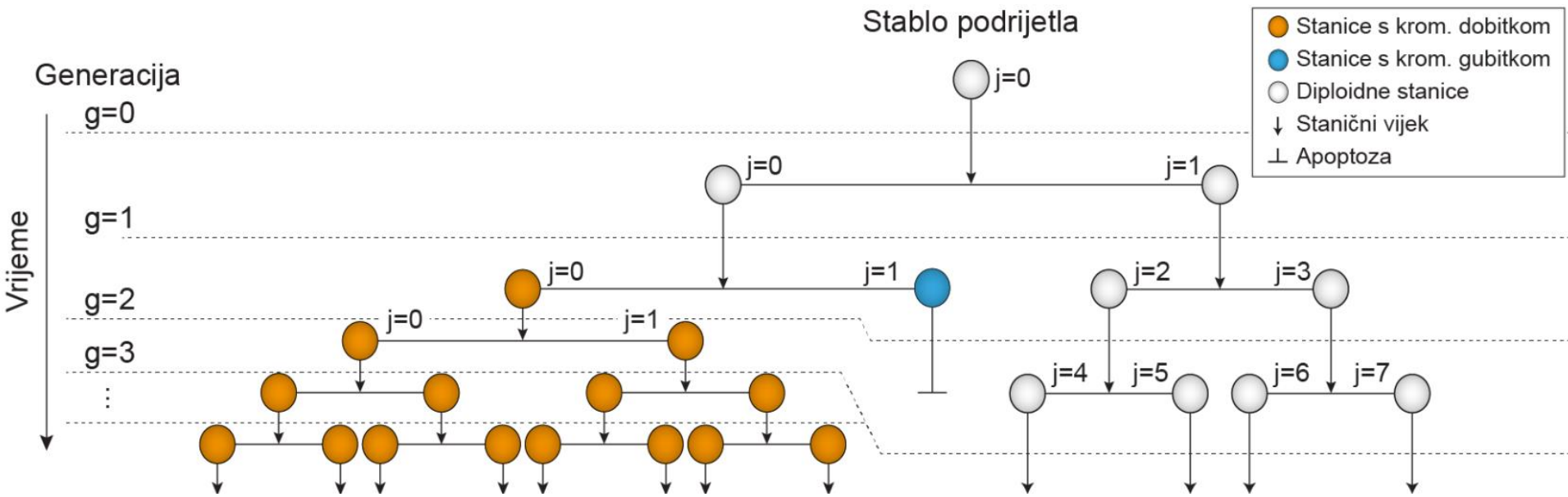


ovisnost o
ishodu



nemogućnost praćenja
velikog broja stanica

Pristup srednjeg polja

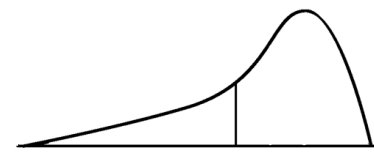


Pristup srednjeg polja



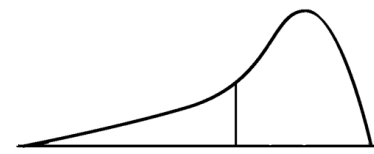
$$\frac{dP_{g,j}(\vec{K})}{dt} = k_d(\vec{K})P_{g-1, \lfloor \frac{j}{2} \rfloor}(\vec{K}) +$$

Pristup srednjeg polja



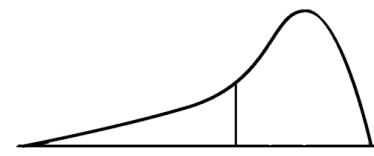
$$\begin{aligned} \frac{dP_{g,j}(\vec{K})}{dt} &= k_d(\vec{K})P_{g-1, \lfloor \frac{j}{2} \rfloor}(\vec{K}) + \\ &+ \frac{1}{2} \sum_{i=1}^n k_{m_i}(\vec{K} + \vec{e}_i)P_{g-1, \lfloor \frac{j}{2} \rfloor}(\vec{K} + \vec{e}_i) + \end{aligned}$$

Pristup srednjeg polja



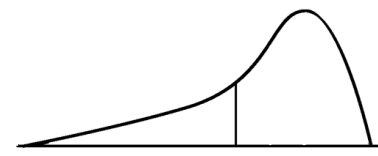
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Pristup srednjeg polja



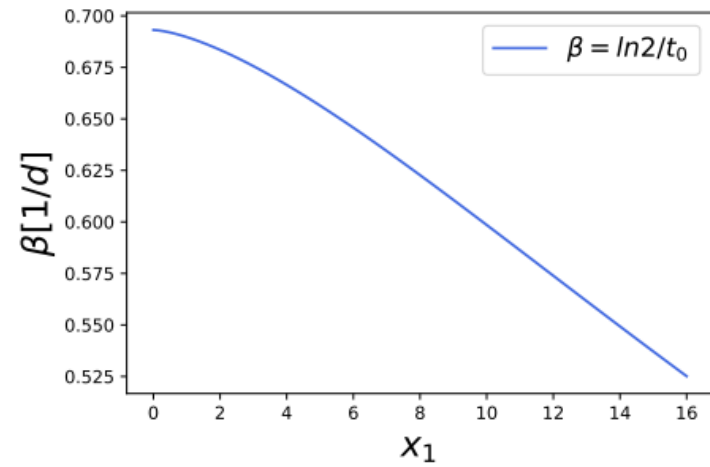
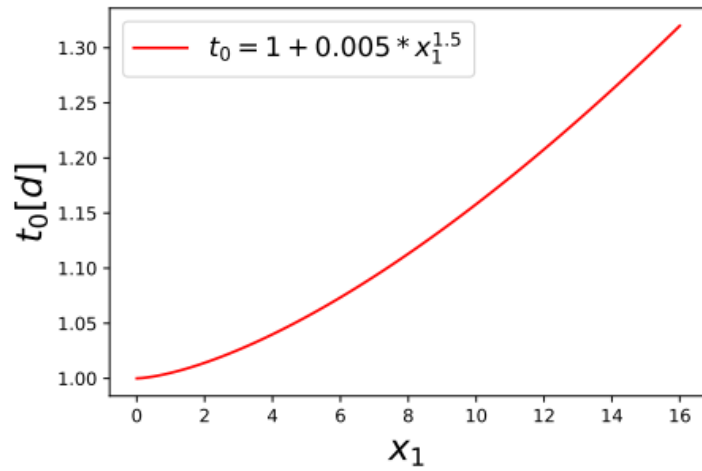
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Pristup srednjeg polja



$$\begin{aligned} \frac{d\tilde{N}(\vec{M})}{dt} = & \left[\tilde{k}_d(\vec{M}) - \tilde{k}_m(\vec{M}) - \tilde{k}_a(\vec{M}) \right] \tilde{N}(\vec{M}) \quad + \\ & + \sum_{l=2}^L \tilde{k}_{m_l}(\vec{M} - \vec{e}_{l-1} + \vec{e}_l) \tilde{N}(\vec{M} - \vec{e}_{l-1} + \vec{e}_l) \quad + \\ & + \sum_{l=1}^{L-1} \tilde{k}_{m_l}(\vec{M} + \vec{e}_l - \vec{e}_{l+1}) \tilde{N}(\vec{M} + \vec{e}_l - \vec{e}_{l+1}) \end{aligned}$$

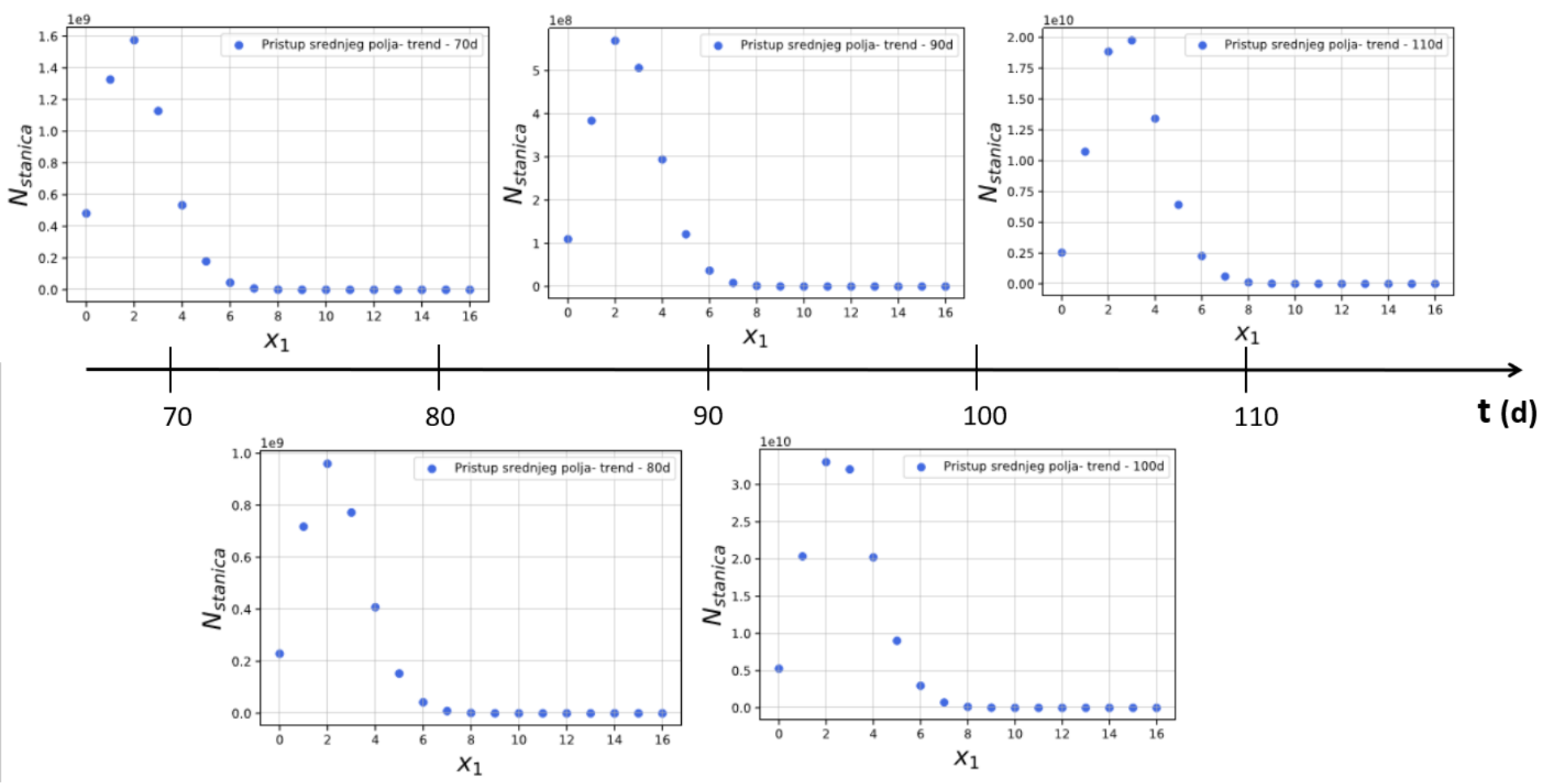
Rezultati





Rezultati

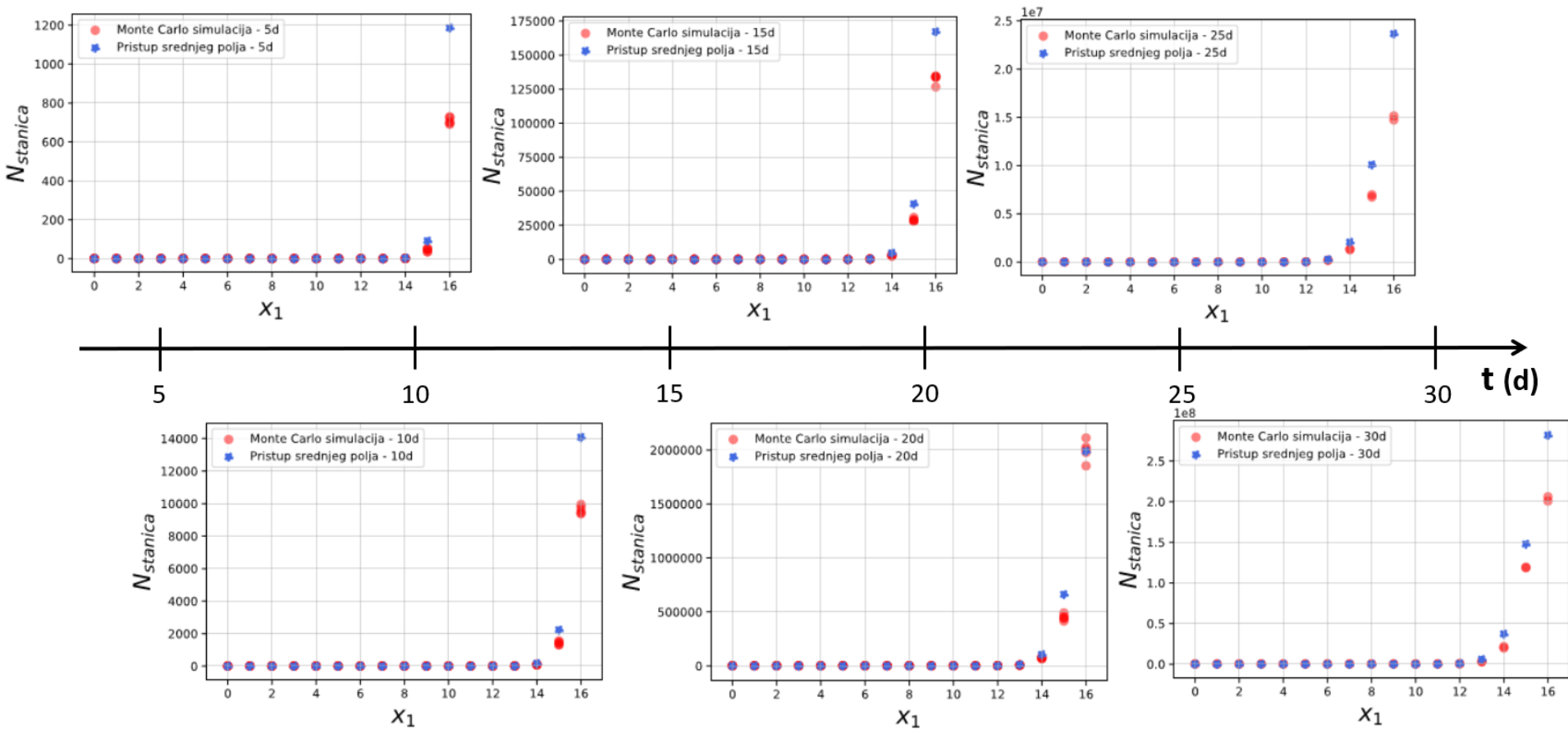
početno diploidna populacija



Rezultati



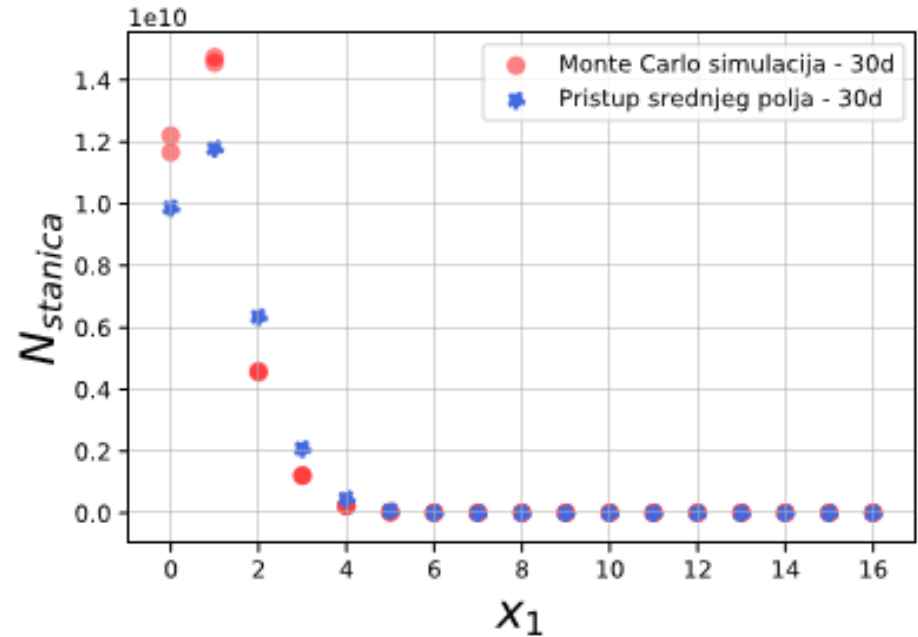
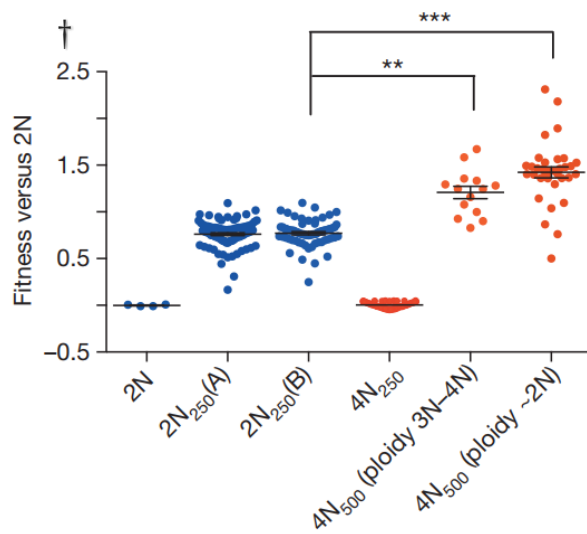
početno haploidna populacija



Zaključak



prilagodba organizama na okolinu



L=2 (II) → L=6 (IIIIII)

† Selmecki, A., Maruvka, Y., Richmond, P. et al. Polyploidy can drive rapid adaptation in yeast. Nature 519, 349–352 (2015).

HVALA!



Tehnikalije



	0.00- 0.01 d	0.01- 0.02 d	0.02- 0.03 d		0.99- 1.00 d	1.00- 1.01 d	1.01- 1.02 d		1.99- 2.00 d	2.00- 2.01 d	2.01- 2.02 d		2.99- 3.00 d	3.00- 3.01 d	3.01- 3.02 d
$X_1=0$	1	0	0		0	0	0		0	0	0		0	0	0
$X_1=1$	0	0	0		0	0	0		0	0	0		0	0	0
$X_1=2$	0	0	0		0	0	0		0	0	0		0	0	0
$X_1=3$	0	0	0	...	0	0	0	...	0	0	0	...	0	0	0
$X_1=4$	0	0	0		0	0	0		0	0	0		0	0	0
$X_1=5$	0	0	0		0	0	0		0	0	0		0	0	0
...						
$X_1=15$	0	0	0		0	0	0		0	0	0		0	0	0
$X_1=16$	0	0	0		0	0	0		0	0	0		0	0	0

Tehnikalije



NEDOSTATAK



ovisnost o
ishodu

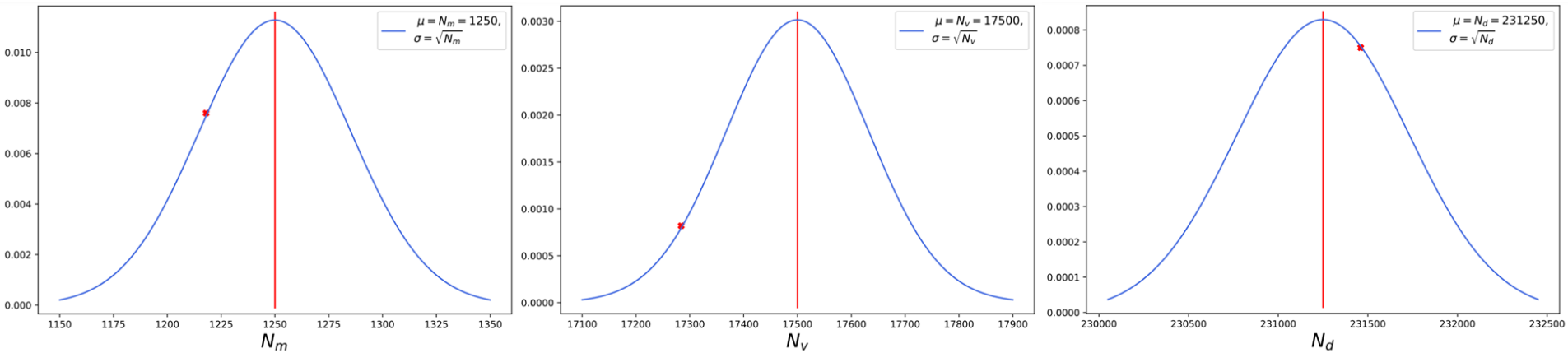


~
nemogućnost praćenja
velikog broja stanica

Tehnikalije

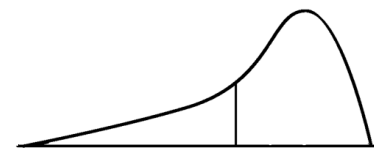


$$N_0 = 250000$$
$$\vec{M} = (2, 14)$$



$$N_m = N_0 \times p_{m_1} = 1250 \quad N_v = N_0 \times p_{m_2} = 17500 \quad N_d = N_0 \times p_d = 231250$$

Tehnikalije



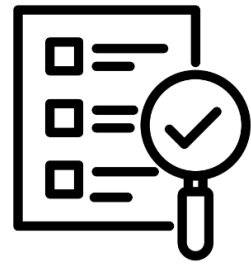
$$\frac{dP_{g,j}(\vec{K})}{dt} = k_d(\vec{K})$$

$$+ \frac{1}{2}$$

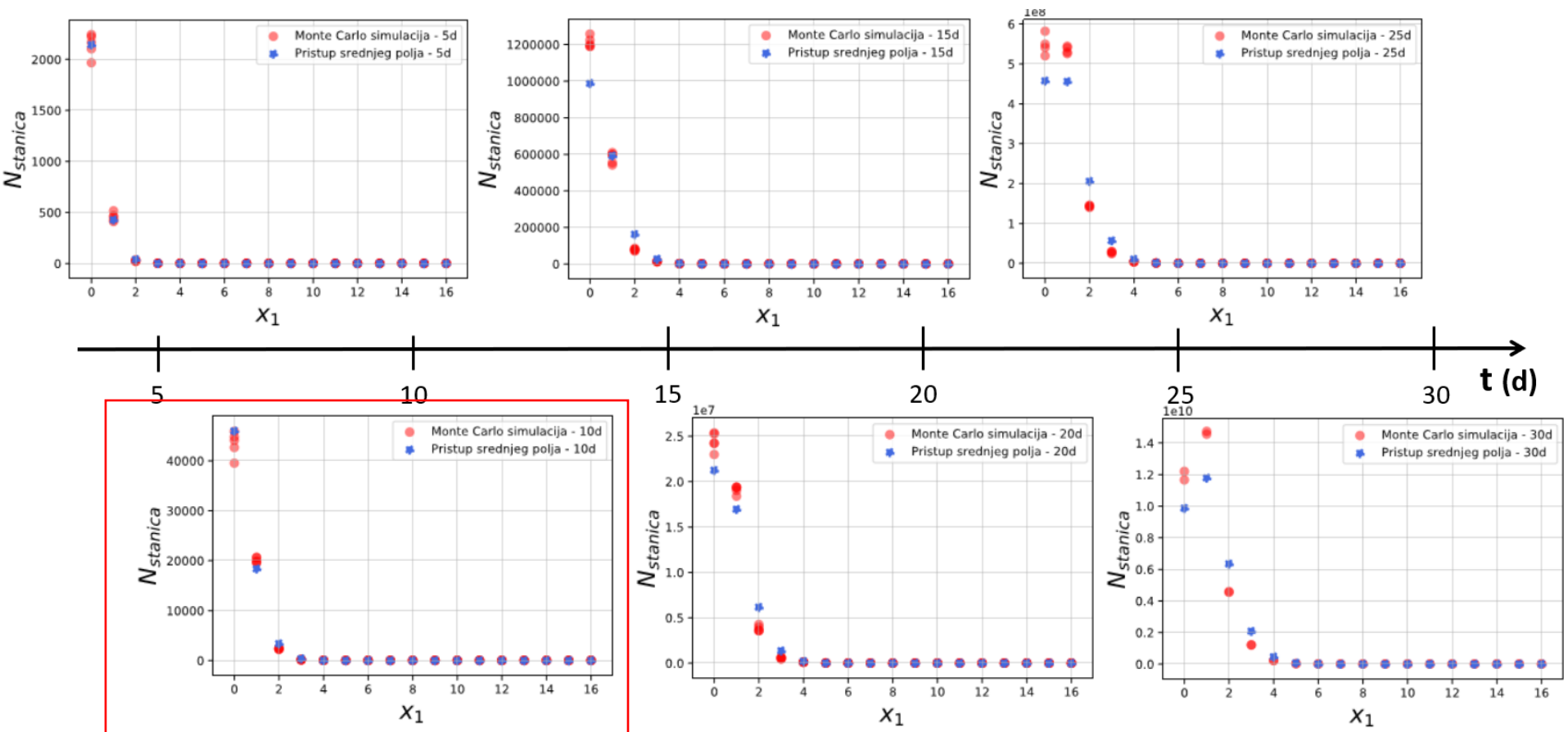
$$N(\vec{K}) \equiv \sum_{g=0}^{\infty} \sum_{j=0}^{2^g-1} P_{g,j}(\vec{K})$$

$$\tilde{N}(\vec{M}) = \frac{1}{x_1!x_2!\dots x_L!} \times \sum_{perm.} N(\vec{K})$$

Tehnikalije



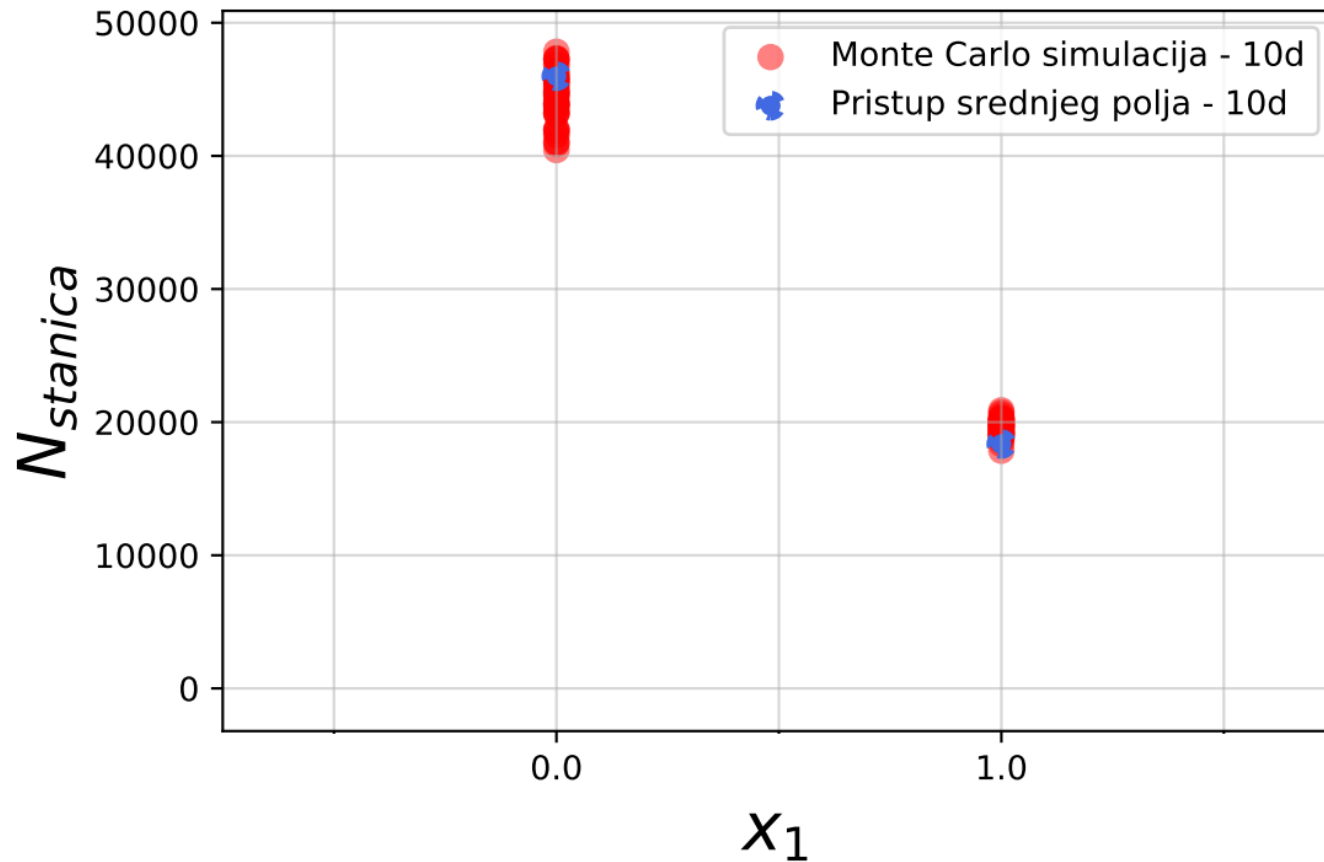
početno diploidna populacija



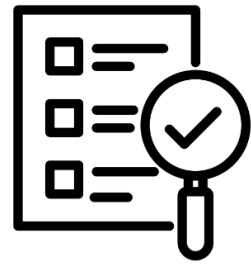
Tehnikalije



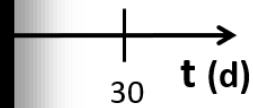
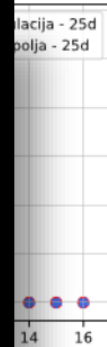
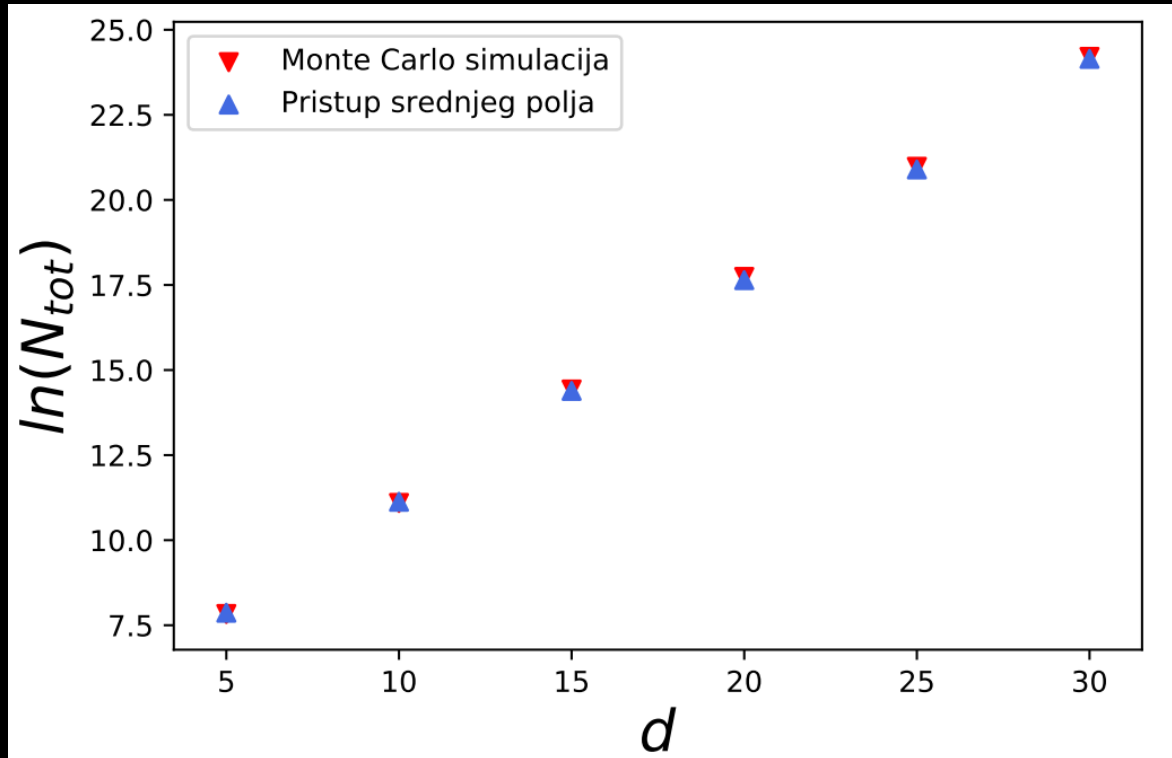
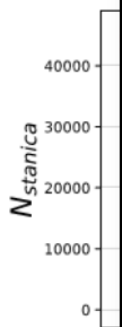
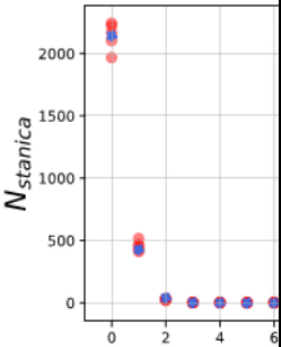
početno diploidna populacija, 50 ishoda



Tehnikalije



početno diploidna populacija

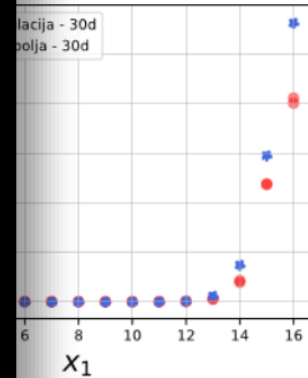
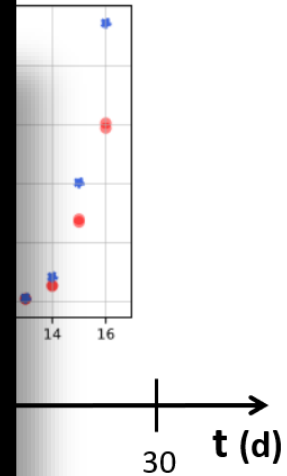
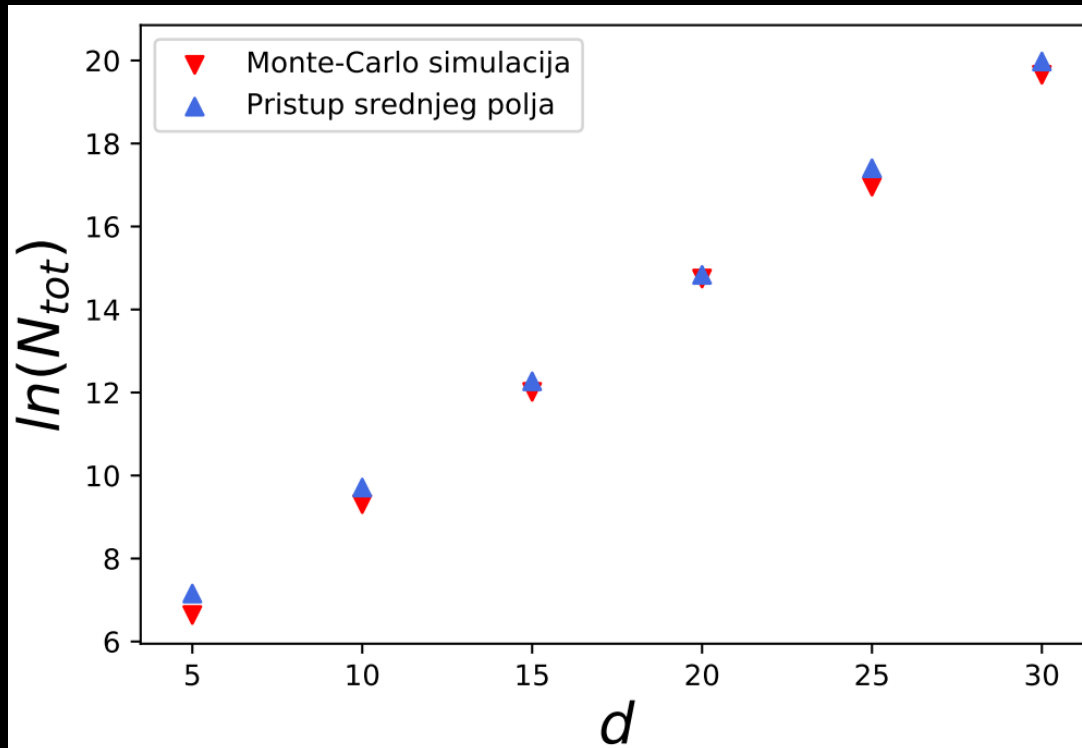
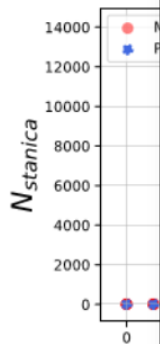
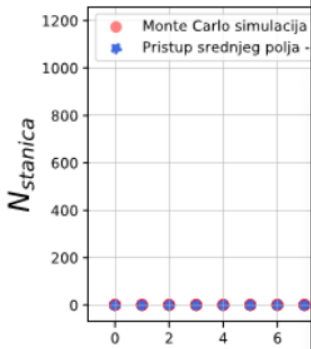


X₁

Tehnikalije



početno haploidna populacija

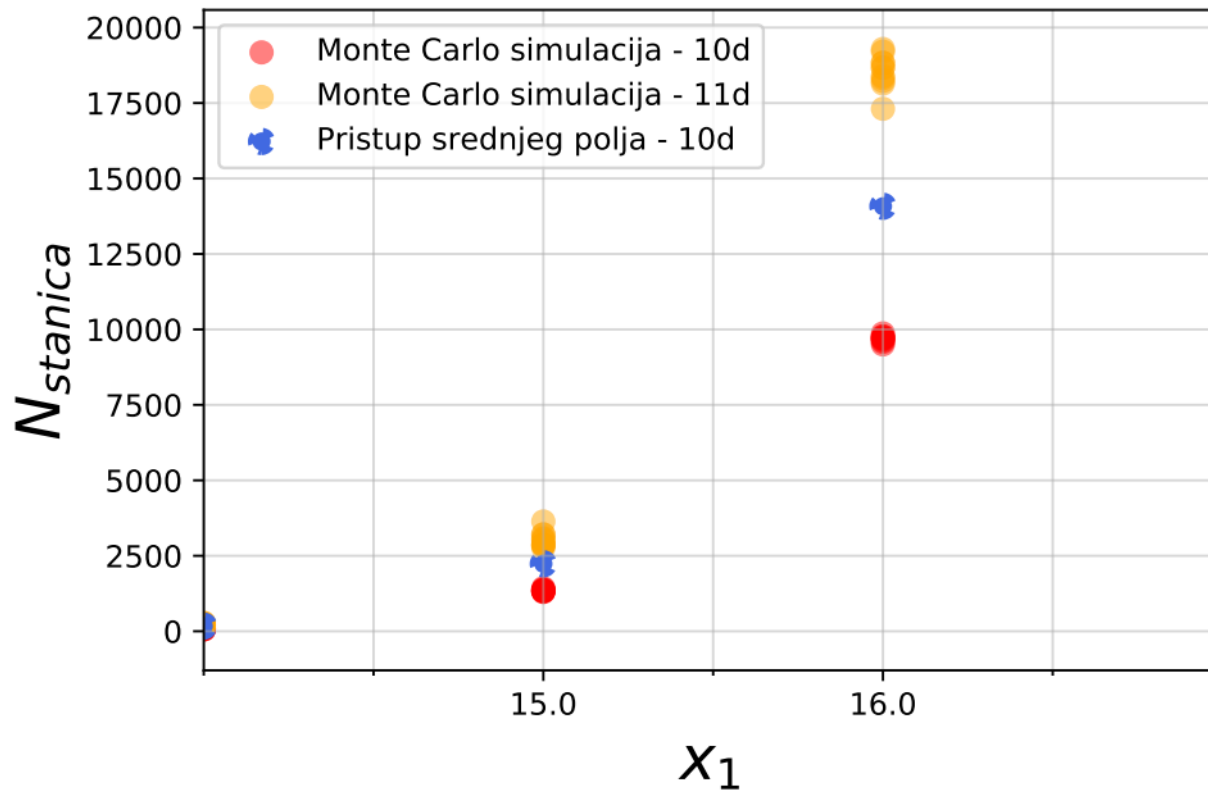


Tehnikalije



početno haploidna populacija

Broj stanica $N_{stanica}$ s obzirom na broj kromosoma s jednom kopijom x_1



Tehnikalije



10(10d)&10(11d) avg

Broj stanica $N_{stanica}$ s obzirom na broj kromosoma s jednom kopijom x_1

