

Numerical approach and computer simulations of population evolution with migration between locations

Maria Magdoń-Maksymowicz

Department of Mathematical Statistics

University of Agriculture in Krakow

Andrzej Maksymowicz

The School of Banking and Management in Krakow



... population evolution with migration between locations

Plan :

- Model
 - Specification of model parameters
- Results
 - Simulations and numerical calculations
- Conclusions



... population evolution with migration between locations

The model and model parameters

We consider two habitats (L_1, L_2) occupied by two species (A, B).
Model parameters (b, m, T) & p are:

birth rate	b	
bad mutations rate	m	admitted at birth
resistance to mutations	T	
migration rate between locations	p	

We used

in locations:	L_1	$m=0.01$	L_2	$m=0.04$
for species:	A	$(b, T)=(0.25, 1)$		
	B	$(b, T)=(0.20, 8)$		

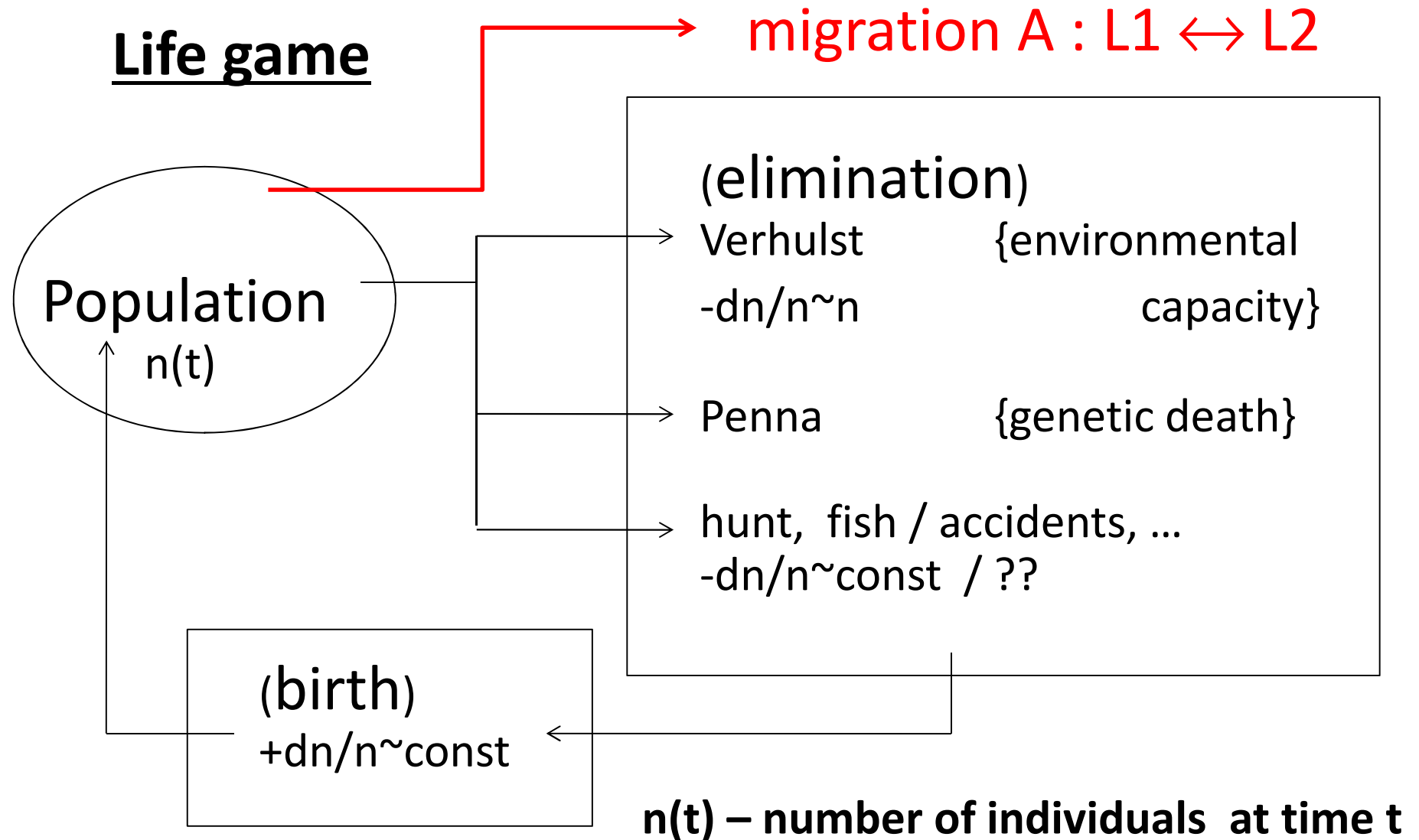
- On output, we get population $x = n/N$ normalized to limited environmental capacity N .
- The set of parameters were chosen, so that **for $p=0$:**
A wins competition in L_1 and B is the winner in location L_2 .

... population evolution with migration between locations

The model and model parameters – (2)

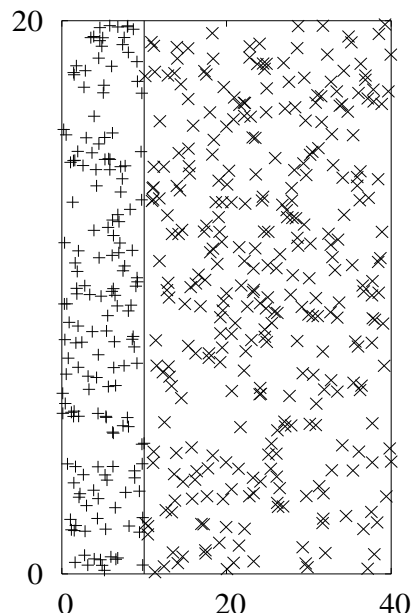
- We make species A migrants, with probability p of a transfer, and we concentrate on two basic schemes:
 - ‘one way ticket’ migration from habitat 1 to 2
 - ‘return visa’ migration from 1 to 2 or back from 2 to 1
- Limiting case $p=0$, within the simplest reference logistic model (with death toll due to the limited environmental capacity only) predicts population $x^*=n/N=b/(1+b)$.
- In simulations we apply Penna model that also accounts for genetic death if number of ‘bad mutations’ reaches threshold value T . Then population $x < x^*$ is expected in each of the isolated locations.
- As we mentioned, in each locations only one species takes over, here A wins in 1 while B wins in habitat 2.
- More exciting is the case with migration, $p>0$, between habitats.

Scheme of numerical calculations

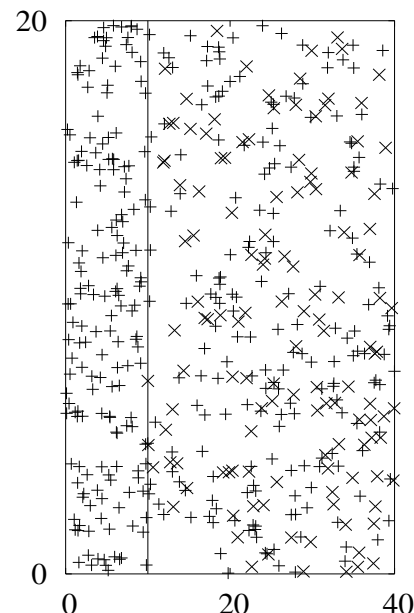


... population evolution with migration between locations

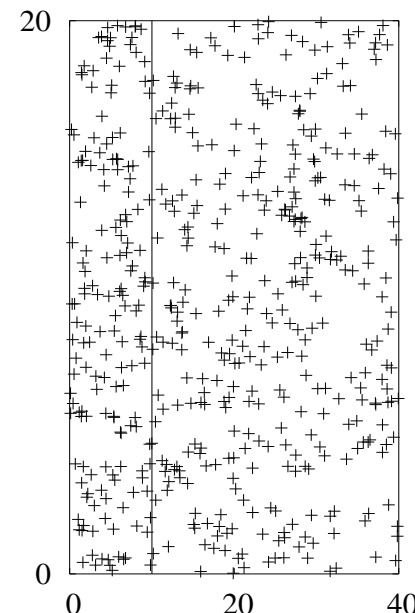
Results for 'one way ticket' migration species A from habitat L_1 to L_2
from left scheme: $p = 0$, moderate p ($p < p_1$), high p , panic ($p > p_2$)



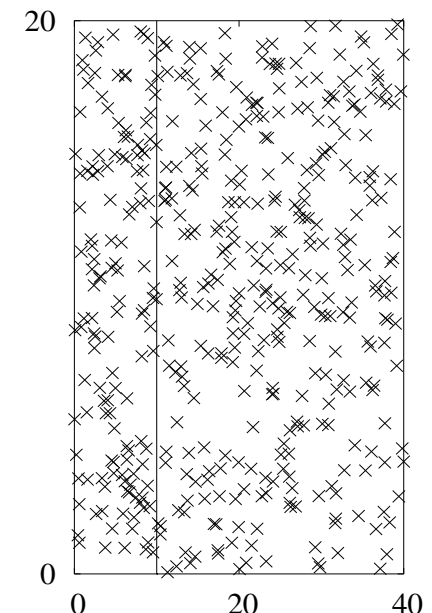
population splits: 10+01 ($p=0$)



coexistence: 10+11 ($p=0.0008$)



emigrants wins: 10+10 ($p=0.012$)



natives wins: 01+01 ($p=0.03$)

Maria Magdoń-Maksymowicz and Andrzej Maksymowicz

... population evolution with migration

Conclusions

- With no migration, one species takes over and wins.
- For 'one way ticket' migration scheme:
 - population of emigrants is a linear function of p , for $0 < p < p_1 = 0.001$ – type A occupies location L_1 and we get a mixed population in L_2
 - we get saturation for $p_1 < p < p_2 = 0.02$, type A wins in both habitats L_1 and L_2
 - catastrophic exodus-like reverse effect for $p > p_2$, with total extinction of species A, species B wins in both locations L_1 and L_2 .
- For 'return visa' scheme: population of emigrants reaches some kind of a dynamic equilibrium in habitat L_2 while in L_1 species A wins.

THANK YOU FOR ATTENTION