A stochastic model for speciation by mating preferences

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CMAP

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In collaboration with C. Coron, M. Costa and C. Smadi







Motivations







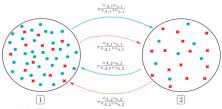
Pundamilia nyererei

What is the effect of a mating preference on the speciation?

- ▶ Homogamy: individuals with similar phenotypes have a higher reproductive success,
- ▶ Heterogamy: individuals with different phenotypes have a higher reproductive sucess.

Model description

- Two patches,
- Haploid population,
- two alleles: a and A,



- A-individuals
- * a-individuals
- migration of A-individuals from one patch to the other (rate indicated)
- migration of a-individuals from one patch to the other (rate indicated)

Dynamics of the population $Z_{A,1}^K$

$$Z_{A,1}^K = \frac{\sharp \{ \text{individuals of type } A \text{ in the patch } 1 \}}{K}$$

• Birth: $b \frac{\beta Z_{A,1}^K + Z_{a,1}^K}{Z_{A,1}^K + Z_{a,1}^K} Z_{A,1}^K$,

 \rightarrow **Homogamy**: two mating individuals that carry the same allele have a probability β **times larger** to give birth to a viable offspring.



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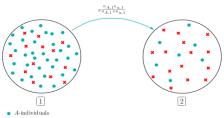
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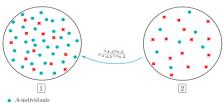


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- Migration from the patch 2: $p \frac{Z_{a,2}^K}{Z_{A,2}^K + Z_{a,2}^K} Z_{A,2}^K$

The total population is represented by

$$\mathbf{Z}_{t}^{K} = (Z_{A,1}^{K}, Z_{a,1}^{K}, Z_{A,2}^{K}, Z_{a,2}^{K})$$



Large population approximation

Assumptions:

$$\triangleright K \to +\infty$$

$\mathsf{Theorem}$

$$\lim_{K \to \infty} \sup_{s \le T} \| \mathbf{Z}^K(s) - \mathbf{z}^{(\mathbf{z}^0)}(s) \| = 0 \quad \text{in probability}, \tag{1}$$

where $\|.\|$ denotes the L^{∞} -Norm on \mathbb{R}^4 , and $\mathbf{z}^{(\mathbf{z}^0)}$ is the solution to the dynamical system

$$\begin{cases} \frac{d}{dt}z_{A,1}(t) = z_{A,1} \left[b \frac{\beta z_{A,1} + z_{a,1}}{z_{A,1} + z_{a,1}} - d - c(z_{A,1} + z_{a,1}) - p \frac{z_{a,1}}{z_{A,1} + z_{a,1}} \right] + p \frac{z_{A,2} z_{a,2}}{z_{A,2} + z_{a,2}} \\ \frac{d}{dt}z_{a,1}(t) = z_{a,1} \left[b \frac{\beta z_{a,1} + z_{A,1}}{z_{A,1} + z_{A,1}} - d - c(z_{A,1} + z_{a,1}) - p \frac{z_{A,1}}{z_{A,1} + z_{a,1}} \right] + p \frac{z_{A,2} z_{a,2}}{z_{A,2} + z_{a,2}} \\ \frac{d}{dt}z_{A,2}(t) = z_{A,2} \left[b \frac{\beta z_{A,2} + z_{a,2}}{z_{A,2} + z_{a,2}} - d - c(z_{A,2} + z_{a,2}) - p \frac{z_{a,2}}{z_{A,2} + z_{a,2}} \right] + p \frac{z_{A,1} z_{a,1}}{z_{A,1} + z_{a,1}} \\ \frac{d}{dt}z_{a,2}(t) = z_{a,2} \left[b \frac{\beta z_{a,2} + z_{A,2}}{z_{A,2} + z_{a,2}} - d - c(z_{A,2} + z_{a,2}) - p \frac{z_{A,2}}{z_{A,2} + z_{a,2}} \right] + p \frac{z_{A,1} z_{a,1}}{z_{A,1} + z_{a,1}}. \end{cases}$$

[Ethier, Kurtz 1986]

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Stationary states

There exists 4 stable stationary states:

2 equilibria for which each type remains present

$$(\zeta,0,0,\zeta), (0,\zeta,\zeta,0)$$

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where $\zeta = \frac{b\beta - d}{c}$.



Stationary states

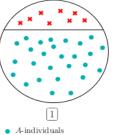
There exists 10 unstable stationary states:

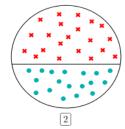
- the extinction state: (0,0,0,0),
- 4 equilibria with only one type in only one patch,
- the equilibrium with both types remaining in both patches with the same proportion of each type,
- 4 equilibria with both types remaining in both patches with different proportions of each type.

Long time behavior of (2)

Let

$$\mathcal{D} := \{ \mathbf{z} \in \mathbb{R}^4_+, z_{A,1} - z_{a,1} > 0, z_{a,2} - z_{A,2} > 0 \},$$





* a-individuals

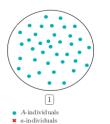
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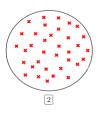
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Theorem

Let $p < p_0$. Then any solution to (2) which starts from \mathcal{D} converges to the equilibrium $(\zeta, 0, 0, \zeta)$.





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Theorem

Let $p < p_0$. Then if the initial condition of (2) lies in

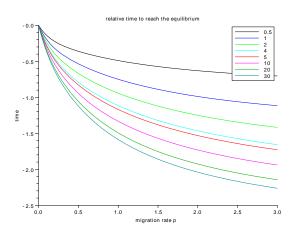
$$\mathcal{K}_p := \left\{ \textbf{z} \in \mathcal{D}, \ \{z_{A,1} + z_{a,1}, \ z_{A,2} + z_{a,2}\} \in \left[\frac{b(\beta+1) - 2d - p}{2c}, \frac{2b\beta - 2d + p}{2c} \right] \right\},$$

there exist two positive constants k_1 and k_2 , depending on the initial condition, such that for every $t \ge 0$,

$$\|\mathbf{z}(t) - (\zeta, 0, 0, \zeta)\| \le k_1 e^{-k_2 t}.$$

Influence of the migration parameter p

$$p\mapsto T_{\varepsilon}(p)-T_{\varepsilon}(0)$$

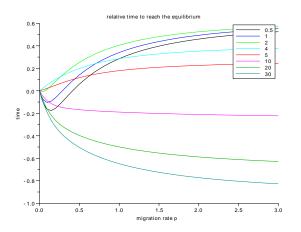


$$z_{a,1}(0) = z_{A,1}(0) - 0.1, \quad z_{A,2}(0) = 1, \quad z_{a,2}(0) = 30.$$

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Influence of the migration parameter p

$$p\mapsto T_{\varepsilon}(p)-T_{\varepsilon}(0)$$



$$z_{a,1}(0) = z_{A,1}(0) - 0.1, \quad z_{A,2}(0) = 15, \quad z_{a,2}(0) = 16.$$

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Time before speciation

If
$$(\mathbf{Z}^K(0))_{K\geq 1}$$
 converges to \mathbf{z}^0





Theorem

Let $\varepsilon > 0$ and $p < p_0$, the time before \mathbf{Z}^K reaches the set

$$\mathcal{B}_{\varepsilon} := [\zeta - \varepsilon, \zeta + \varepsilon] \times \{0\} \times \{0\} \times [\zeta - \varepsilon, \zeta + \varepsilon],$$

is

$$\frac{1}{b(\beta-1)}\log(K)$$





when K tends to $+\infty$.

Conclusion: Without any ecological difference, mating preferences are enough to entail a reproductive isolation.

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