

Selection and mutation in a shifting and fluctuating environment

Sepideh Mirrahimi

CNRS, Institut de mathématiques de Toulouse

Joint work with Susely Figueroa Iglesias (IMT)

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Motivating example 1: The influence of fluctuating temperature on bacteria

Bacteria evolved in **fluctuating temperature** (daily variation between 24°C and 38°C, mean 31°C), **outperforms** the strain that evolved in **constant environments** (31°C):

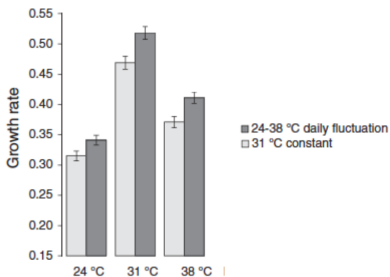


Figure from: Ketola et al. 2013

Motivating example 2: Earth's temperature changes (increase and oscillations)

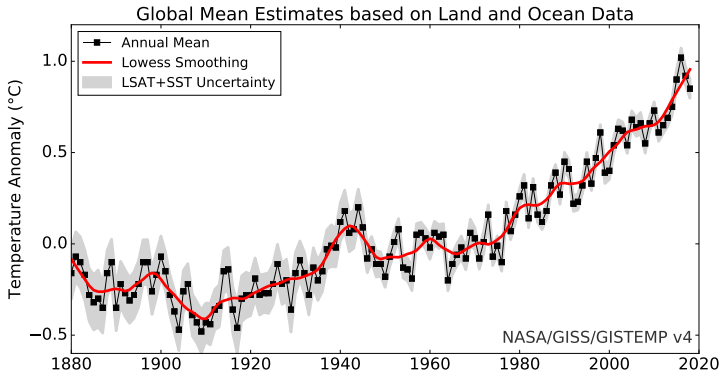


Figure from: data.giss.nasa.gov

Selection-mutation in a shifting and fluctuating environment

$$\left\{ \begin{array}{l} \frac{\partial}{\partial t} m - \underbrace{\sigma \frac{\partial^2}{\partial z^2} m}_{\text{mutations}} = m \left(\underbrace{R(e(t), z - ct)}_{\text{growth rate}} - \underbrace{\kappa M}_{\text{competition}} \right), \\ M(t) = \int_{\mathbb{R}} m(t, y) dy, \quad m(t = 0, \cdot) = m_0(\cdot). \end{array} \right.$$

- z : phenotypic trait
- $m(t, z)$: density of trait z
- σ : mutation effective size
- $e(t)$: environment state
- $R(e, z)$: growth rate
- c : speed of the linear change
- $M(t)$: size of the population
- κ : intensity of the competition

A change of notation and questions

Density in the moving framework: $n(t, z) = m(t, z + ct)$:

$$\begin{cases} \frac{\partial}{\partial t} n - c \frac{\partial}{\partial z} n - \sigma \frac{\partial^2}{\partial z^2} n = n(R(e(t), z) - \kappa N), \\ N(t) = \int_{\mathbb{R}} n(t, y) dy. \end{cases}$$

Objective: to characterize the phenotypic density n .

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Objective: to characterize the phenotypic density n .

Questions:

- Critical speed of environment change above which the population gets extinct?
- Impact of the oscillations on the critical speed and the phenotypic distribution?
- In which situation, a population evolved in a periodic environment (without linear change) would outperform a population evolved in a constant environment?

Some references

- **Time-varying environments:**

Lynch et al (1991), Lynch and Lande (1993), Burger and Lynch (1995), Lande and Shannon (1996)

(assumptions: quadratic stabilizing selection, Gaussian phenotypic distribution, the environment change acts only on the optimum)

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- **The Hamilton-Jacobi approach:**

First papers: Diekmann et al (2005), Barles–Perthame (2007–2008)

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- **The Hamilton-Jacobi approach:**

First papers: Diekmann et al (2005), Barles–Perthame (2007–2008)

- More quantitative results:

M.–Roquejoffre (2015-2016), M.–Gandon (2017-2019)

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Assumptions

Notation:

$$\bar{R}(z) = \frac{1}{T} \int_0^T R(e(t), z) dt.$$

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Assumptions:

- e T -periodic with respect to t
- There exists a unique $z_m \in \mathbb{R}^d$ such that

$$\max_{z \in \mathbb{R}^d} \bar{R}(z) = \bar{R}(z_m) > 0$$

- There exists a unique $\bar{z} < z_m$ such that

$$\bar{R}(\bar{z}) + \frac{c^2}{4\sigma} = \bar{R}(z_m).$$

Critical speed for survival

$$\lambda_\sigma : \text{principal eigenvalue} \quad \left\{ \begin{array}{l} \frac{\partial}{\partial t} p - \sigma \frac{\partial^2}{\partial z^2} p - R(e, z)p = \lambda_\sigma p, \\ p(t, z) = p(t + T, z), \quad p > 0. \end{array} \right.$$

$$\text{Define} \quad c_\sigma = \begin{cases} 2\sqrt{-\sigma\lambda_\sigma} & \text{if } \lambda_\sigma < 0, \\ 0 & \text{otherwise.} \end{cases}$$

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Proposition (Figueroa Iglesias and M., SIMA 2018, Preprint 2019)

- (i) $c \geq c_\sigma$: $N(t) \rightarrow 0$ as $t \rightarrow \infty$.
 (ii) $c < c_\sigma$: $n(t, \cdot)$ converges to a the unique positive solution to

$$\begin{cases} \frac{\partial}{\partial t} n_p - c \frac{\partial}{\partial z} n_p - \sigma \frac{\partial^2}{\partial z^2} n_p = n_p (R(e, z) - \kappa N_p), \\ N_p(t) = \int_{\mathbb{R}} n_p(t, y) dy, \quad n_p(t + T, z) = n_p(t, z). \end{cases}$$

An asymptotic expansion for the critical speed

Regime of small mutations:

$$\sigma = \varepsilon^2.$$

Theorem (Figueroa Iglesias and M., Preprint 2019)

$$c_\varepsilon = 2\varepsilon\sqrt{\bar{R}(z_m)} - \varepsilon^2\sqrt{-\bar{R}''(z_m)} + o(\varepsilon^2).$$

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Define the renormalized speeds

$$c = \tilde{c}\varepsilon, \quad c_\varepsilon = \tilde{c}_\varepsilon\varepsilon.$$

Objective: to characterize the phenotypic density $n_{\varepsilon,p}$ when $\tilde{c} < \tilde{c}_\varepsilon$:

$$\begin{cases} \frac{\partial}{\partial t} n_{\varepsilon,p} - \varepsilon\tilde{c}\frac{\partial}{\partial z} n_{\varepsilon,p} - \varepsilon^2\frac{\partial^2}{\partial z^2} n_{\varepsilon,p} = n_{\varepsilon,p}[R(e(t), z) - \kappa N_{\varepsilon,p}(t)], \\ N_{\varepsilon,p}(t) = \int_{\mathbb{R}} n_{\varepsilon,p}(t, y) dy, \quad n_{\varepsilon,p}(0, z) = n_{\varepsilon,p}(T, z). \end{cases}$$

The Hamilton-Jacobi approach

The **Hopf-Cole** transformation:

$$n_{\varepsilon,p}(t, z) = \frac{1}{(2\pi\varepsilon)^{\frac{d}{2}}} \exp\left(\frac{u_{\varepsilon,p}(t, z)}{\varepsilon}\right).$$

An expected asymptotic expansion:

$$u_{\varepsilon,p}(t, z) = u + \varepsilon v + O(\varepsilon^2).$$

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Theorem (Figuroa Iglesias and M., SIMA 2018, Preprint 2019)

As $\varepsilon \rightarrow 0$, $u_{\varepsilon,p}$ converges to u which is uniquely and explicitly determined. In particular, it solves

$$\begin{cases} -\left|\frac{\partial}{\partial z} u(z) + \frac{\tilde{c}}{2}\right|^2 = \bar{R}(z) - \kappa \bar{N} - \frac{\tilde{c}^2}{4}, \\ \max u(z) = u(\bar{z}) = 0. \end{cases}$$

The population follows the optimum with a constant lag

Theorem (Figueroa Iglesias and M., SIMA 2018, Preprint 2019)

Assume that $\tilde{c} < \lim_{\varepsilon \rightarrow 0} \tilde{c}_\varepsilon$. Then, as $\varepsilon \rightarrow 0$,

$$n_{\varepsilon,p}(t, z) \longrightarrow N_p(t) \delta(z - \bar{z}).$$

with $N_p(t)$ the unique periodic solution of

$$\frac{dN_p}{dt} = N_p [R(e, \bar{z}) - N_p].$$

In the original problem before the translation

$$m_{\varepsilon,p}(t, x) \longrightarrow \varrho(t) \delta(z - \bar{z} - ct).$$

Recall: \bar{z} the unique point such that $\bar{R}(\bar{z}) + \frac{c^2}{4\sigma} = \bar{R}(z_m)$ and $\bar{z} < z_m$.

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A constant environment: Non-Gaussian distribution

$$R(e, z) = R_0(z) = 3 - (z + 0.5)^2 (0.2 + (z - 0.5)^2), \quad \varepsilon = 0.1$$

Blue dots:
numerical computation of
equilibrium

Black line:
our approximation

Vertical dotted line:
mean phenotypic trait

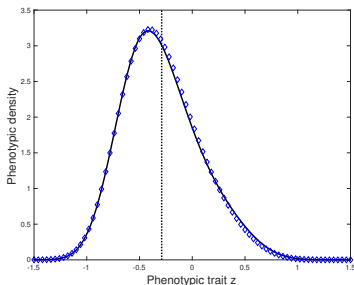


Figure from Mirrahimi–Gandon 2019

Some notations

Average **size of the population** over a period of time:

$$\bar{N}_{\epsilon,p} = \frac{1}{T} \int_0^T N_{\epsilon,p}(t) dt$$

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Mean **phenotypic trait**:

$$\mu_{\epsilon,p}(t) = \frac{1}{N_{\epsilon,p}(t)} \int_{\mathbb{R}} z n_{\epsilon,p}(t, z) dz$$

Variance of the phenotypic distribution:

$$v_{\epsilon,p}(t) = \frac{1}{N_{\epsilon,p}(t)} \int_{\mathbb{R}} (z - \mu_{\epsilon,p}(t))^2 n_{\epsilon,p}(t, z) dz$$

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Mean fitness in an environment with constant state \bar{e} :

$$F_{\varepsilon,p}(\bar{e}) = \int_{\mathbb{R}} R(\bar{e}, z) \frac{1}{T} \int_0^T \frac{n_{\varepsilon,p}(t, z)}{N_{\varepsilon,p}(t)} dt dz$$

A constant environment: approximation of the moments

$R(e, z) = R_0(z)$ with a unique maximum at θ_0 : **optimal trait**

$$R_0(z) = r - s(z - \theta_0)^2 + a(z - \theta_0)^3 + O(z - \theta_0)^4$$

r: maximal growth rate, **s**: selection pressure,

a: asymmetry parameter

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average **population size**, **mean phenotypic trait** and **variance**:

$$N_{\varepsilon,0} = r - \varepsilon\sqrt{s} + o(\varepsilon), \quad \mu_{\varepsilon,0} = \theta_0 + \frac{a\varepsilon}{s\sqrt{s}} + o(\varepsilon), \quad v_{\varepsilon,0} = \frac{\varepsilon}{\sqrt{s}} + o(\varepsilon),$$

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and the **mean fitness** in such constant environment:

$$F_{\varepsilon,0} = r - \varepsilon\sqrt{s} + o(\varepsilon).$$

Case 1: Fluctuating optimal trait (no shift $c = 0$)

$$R(e, z) = r - s(z - \theta(e))^2, \quad \theta(e) = e, \quad e(t) = d \sin(2\pi t/b).$$

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average population size, mean phenotypic trait and variance:

$$\bar{N}_{\epsilon, p} \approx r - \frac{sd^2}{2} - \epsilon\sqrt{s}, \quad \mu_{\epsilon, p}(t) \approx \frac{\epsilon db\sqrt{s}}{\pi} \sin\left(\frac{2\pi}{b}(t - b/4)\right),$$

$$v_{\epsilon, p}(t) \approx \epsilon/\sqrt{s}.$$

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Comparison between the **mean fitness** of this population in an environment with constant state ($\bar{e} = 0$) and the mean fitness of a population evolved in the constant environment $\bar{e} = 0$:

$$F_{\epsilon, c}(\bar{e}) \approx F_{\epsilon, p}(\bar{e}) \approx r - \epsilon\sqrt{s}.$$

Case 2: Fluctuating selection pressure (no shift $c = 0$)

$$R(e, z) = r - s(e)z^2 + O(z^4), \quad \bar{s} = \frac{1}{T} \int_0^T s(e(\tau)) d\tau.$$

average population size, mean phenotypic trait and variance:

$$\bar{N}_{\varepsilon, p} \approx r - \varepsilon \sqrt{\bar{s}}, \quad \mu_{\varepsilon, p}(t) \approx 0, \quad v_{\varepsilon, p}(t) \approx \frac{\varepsilon}{\sqrt{\bar{s}}}.$$

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and the **mean fitness** in an environment with constant state \bar{e}

$$F_{\varepsilon, c}(\bar{e}) = r - \varepsilon\sqrt{s(\bar{e})} < F_{\varepsilon, p}(\bar{e}) \approx r - \varepsilon\frac{s(\bar{e})}{\sqrt{\bar{s}}}, \quad \text{if } \bar{s} > s(\bar{e}).$$

The population evolved in the periodic environment may outperform the population evolved in a constant environment.

Case 3: Fluctuating selection pressure with shift ($c > 0$)

Same growth rate (with varying selection pressure):

Critical speed of survival:

Periodic $s(e)$: $\tilde{c}_{\epsilon,p} \approx 2\sqrt{r} - \epsilon \frac{\sqrt{\bar{s}}}{\sqrt{r}}$

Constant $s = s(\bar{e})$: $\tilde{c}_{\epsilon,c} \approx 2\sqrt{r} - \epsilon \frac{\sqrt{s(\bar{e})}}{\sqrt{r}}$

$$\text{if } \bar{s} < s(\bar{e}) \Rightarrow \tilde{c}_{\epsilon,c} < \tilde{c}_{\epsilon,p}$$

\Rightarrow oscillations help the population to follow the environment shift

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$$\text{if } \bar{s} < s(\bar{e}) \Rightarrow \tilde{c}_{\epsilon,c} < \tilde{c}_{\epsilon,p}$$

\Rightarrow oscillations help the population to follow the environment shift

Note: opposite condition to have a more performant population in absence of linear change \Rightarrow what is beneficial in a (in average) constant environment is disadvantageous in a changing environment

Thank you for your attention !