From Fisher to Feynman: nonequilibrium statistics of molecular evolution

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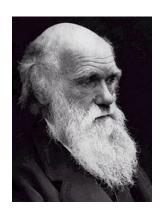
Thanks

Ville Mustonen

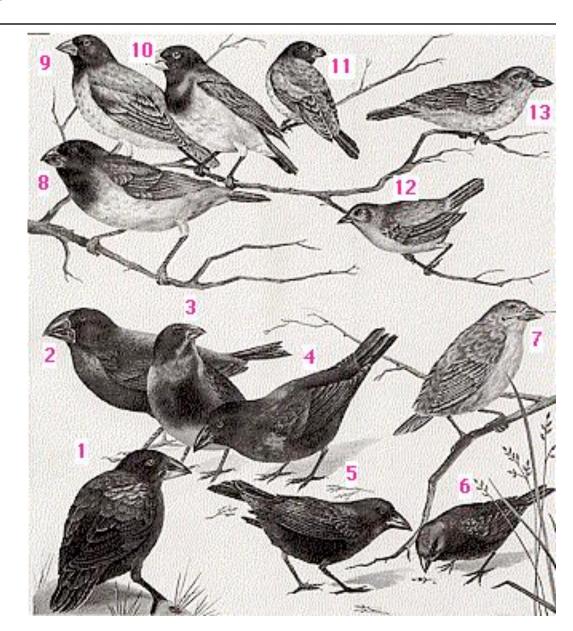
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Justin Kinney (Princeton)
Luca Peliti (Naples)

SFB TR 12 SFB 680 European Research Training Network STIPCO

Darwinian evolution and adaptation



- Adaptative evolution of phenotypes in a population occurs due to natural variation and natural selection.
- Adaptive evolution is an ongoing process, because selection pressures keep changing.



Determinants of molecular evolution

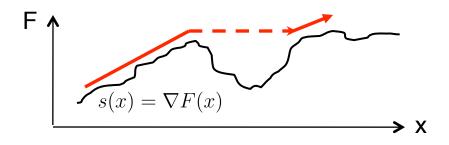
■ Equation of motion for the population fractions (frequencies) $x = (x_1, ...x_n)$ of phenotypes or genotypes in a population:

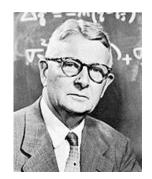
• Equation of motion for the frequency distribution:

$$\frac{\partial}{\partial t}P(x,t) = \nabla \left[\frac{1}{N}\nabla g(x) - s(x)g(x) - mx\right]P(x,t)$$

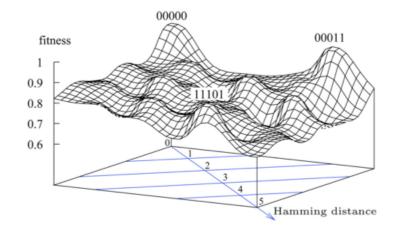
Adaptative evolution

- Evolution in a fitness landscape(S. Wright 1932):
 - interplay of selection and genetic drift
 - longer time intervals: mutations





■ Example: fitness landscape in the fungus *Aspergillus niger*



[A. de Visser, SC. Park, J. Krug 09]

Adaptative evolution

- Fundamental Theorem of Natural Selection (R.A. Fisher 1930):
 - deterministic evolution under time-independent selection alone

$$\frac{d}{dt}F(t) = s^2(x(t))$$

- evolution under time-dependent selection: nonequilibrium

$$\frac{d}{dt}\Phi(t) = s^2(x(t),t)$$
 | fitness flux

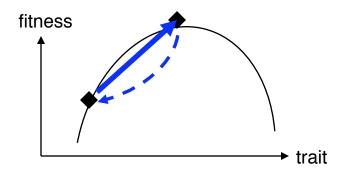
■ Is there an entropy principle of biological evolution? (Schrödinger, What Is Life 1943)

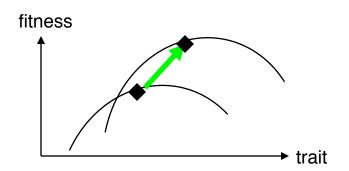




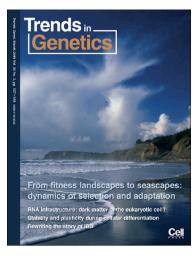
Adaptive evolution

- A comprehensive **theory of molecular evolution** must include:
 - 1. **stochastic forces** (mutations and genetic drift)
 - 2. evolutionary histories, correlations between genomic changes: distinguish *compensatory evolution* from *adaptation:*





3. time-dependent selection: nonequilibrium **fitness seascapes**



1. Fitness flux theorem	

Population histories and fitness flux

A population history is a sequence of frequency measurements

$$\mathbf{x} = (x_0, \dots, x_n)$$
 at times (t_0, \dots, t_n) .

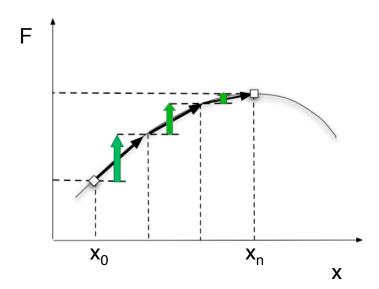
■ The **fitness flux** of a population history is the cumulative **selective effect of frequency changes**:

$$\Phi(\mathbf{x}) \equiv \sum_{i=1}^{n} \Delta x_i \, s(x_i, t_i).$$

Flux in a fitness landscape:

$$s(x) = \nabla F(x)$$

$$\Phi(\mathbf{x}) = \sum_{i=1}^{n} \Delta x_i \nabla F(x_i)$$
$$= F(x_n) - F(x_0).$$



Population histories and fitness flux

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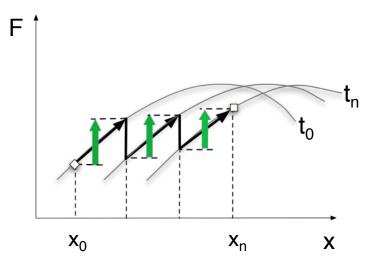
$$\Phi(\mathbf{x}) \equiv \sum_{i=1}^{n} \Delta x_i \, s(x_i, t_i).$$

• Flux in a **fitness seascape**:

$$s(x,t) = \nabla F(x,t)$$

$$\Phi(\mathbf{x}) = \sum_{i=1}^{n} \Delta x_i \nabla F(x_i, t_i)$$

$$\neq F(x_n, t_n) - F(x_0, t_0).$$



Evolutionary equilibrium and fitness

• If the **neutral** process under mutations and genetic drift has an **equilibrium frequency distribution** $P_0(x)$, the process in an **arbitrary fitness landscape** also has an equilibrium

$$P_{\rm eq}(x) = P_0(x) e^{NF(x)} .$$

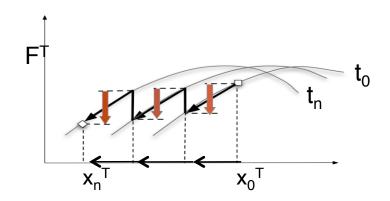
Hence, fitness measures the information of population states:

$$N\langle F
angle_{
m eq} = \int dx \, P_{
m eq}(x) \log rac{P_{
m eq}(x)}{P_0(x)} \equiv H(P_{
m eq}|P_0)$$
 . KL entropy

[J.Berg, S.Willmann, M.L., BMC Evol. Biol. 2004, V. Mustonen, M.L., PNAS 2010, in press]

Fitness flux and time reversal

Each population history x has a reverse history x^T, in which all frequency transitions have opposite fitness effects:

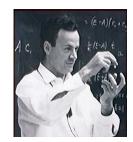


■ The probabilities of forward and reverse history are related:

$$\mathcal{P}(\mathbf{x}^T) = \mathcal{P}(\mathbf{x}) \; \mathrm{e}^{-N\Phi(\mathbf{x})} \; + \; \Delta \mathcal{H}(\mathbf{x})$$
 fitness flux
$$\begin{vmatrix} \mathrm{entropy} \; \mathrm{difference} \\ \mathrm{of} \; \mathrm{initial} \; \mathrm{conditions} \end{vmatrix}$$

$$\mathcal{H}(x,t) \equiv \log[P(x,t)/P_0(x)]$$

$$\Delta \mathcal{H}(\mathbf{x}) = \mathcal{H}(x_n,t_n) - \mathcal{H}(x_0,t_0)$$



Hence, fitness flux measures the information of the evolution process:

$$N\langle\Phi\rangle = H(\mathcal{P}|\mathcal{P}^T) - H(P(t_n)|P_0) + H(P(t_0)|P_0)$$

Fitness flux theorem

■ **Theorem:** For an evolutionary process with mutations, genetic drift and selection given by an arbitrary fitness seascape,

$$\langle e^{-N\Phi + \Delta \mathcal{H}} \rangle = 1$$
.

\(\cdots\): average over population histories,

$$\Delta \mathcal{H}(\mathbf{x}) = \mathcal{H}(x_n, t_n) - \mathcal{H}(x_0, t_0)$$

.

■ Corollary: Φ increases almost universally,

$$\langle \Phi \rangle \geq \Delta H$$
.

$$\Delta H = \langle \Delta \mathcal{H} \rangle = H(P(t_n)|P_0) - H(P(t_0)|P_0)$$
 entropy difference between initial and final state.

[V. Mustonen, M.L., PNAS 2010, in press]

Modes of fitness evolution

• Equilibrium:

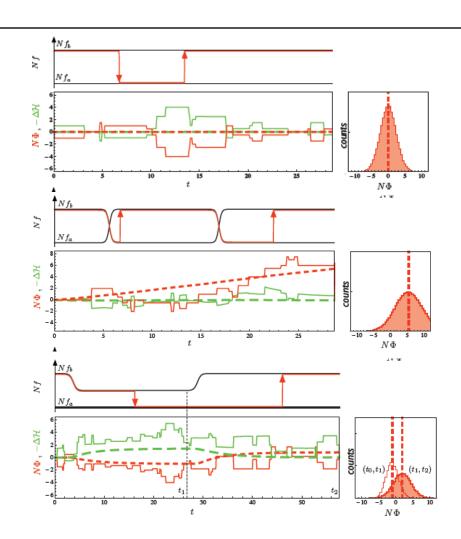
$$\langle \Phi \rangle = 0$$

Stationary nonequilibrium:

$$\langle \Phi \rangle > 0$$

Demographic nonequilibrium:

- $\langle \Phi \rangle < 0$ declining pop. size
- $\langle \Phi \rangle > 0$ recovery



• Strong-selection limit: Fundamental theorem of natural selection

$$\frac{d}{dt}\Phi(t) = s^2(x(t), t)$$

2.	Evolution	and	entropy

Biological evolution

■ (-) energy

$$-E(x,t)$$

thermodynamic equilibrium

$$P_{\rm eq}(x) = C \, \mathrm{e}^{-\beta E(x)}$$

heat flux

$$Q(\mathbf{x}) = \sum_{i=1}^{n} \Delta x_i (-\nabla E)(x_i, t_i)$$

local entropy

$$S(x,t) = -\log P(x,t)$$

fluctuation theorem

$$\langle e^{-\beta Q - \Delta S} \rangle = 1$$

$$\beta \langle Q \rangle + \Delta S = \Delta S_{tot} \ge 0$$

[Seifert 05, cf. Jarzynski 97, Crook 99].

fitness

evolutionary equilibrium

$$P_{\rm eq}(x) = P_0(x) e^{NF(x)}$$

fitness flux

$$\Phi(\mathbf{x}) = \sum_{i=1}^{n} \Delta x_i \, \nabla F(x_i, t_i)$$

local entropy

$$\mathcal{H}(x,t) = \log \frac{P(x,t)}{P_0(x)}$$

fitness flux theorem

$$\left\langle e^{-N\Phi+\Delta\mathcal{H}}\right\rangle = 1$$

 $N\langle\Phi\rangle - \Delta H > 0$

Thermodynamics

Biological evolution



Second Law

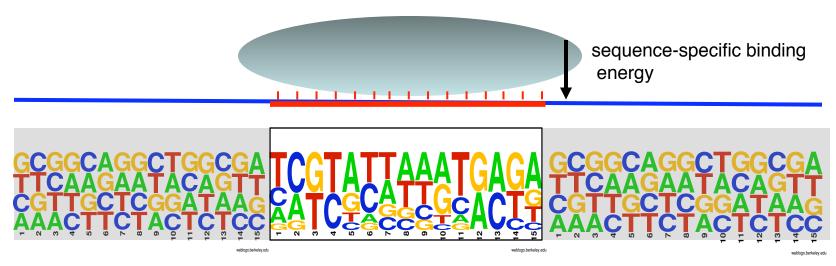


Adaptation can decrease entropy!



Genomic information

Transcription factors bind to DNA target sites.



- Target sites have a more **specific sequence** than background DNA.
- Information gain (entropy loss) in the adaptive formation of a new site (in bacteria or yeast):

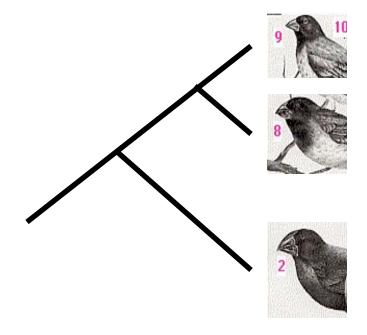
$$\Delta H \approx 15$$
 bytes.

[Mustonen, M.L., PNAS 2005, Mustonen, Kinney, Callen, M.L., PNAS 2008]

• Information content of the entire genome?

3. Irreversibility of evolution

(The length of time's arrow)



Adaptation and fitness flux in flies

- Cross-species data and polymorphisms within species are used to infer rate and average selection coefficient of substitutions.
- Adaptation is quantified by a positive fitness flux:

 Φ = (rate) x (average selection coefficient of substitutions).

- This indicates a stationary nonequilibrium process:
 - Selection coefficients at individual sites fluctuate at nearly the rate of neutral substitutions.
 - There is a surplus of beneficial over deleterious substitutions.

[Mustonen and M.L, PNAS 2007]

Amount of genome-wide adaptation?

Conclusions

- Adaptive evolution is a stochastic nonequilibrium process quantified by fitness flux Φ.
- Drosophila genomes:
 evidence for adaptive evolution driven by fitness seascapes.
- Fitness flux theorem:
 Increase of Φ is a nearly universal evolutionary principle.

