

Ancestral lines under selection and recombination

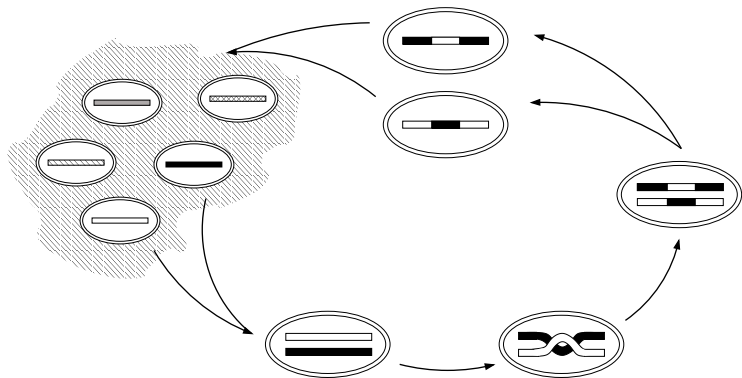
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joint work with Frederic Alberti

1. The selection-recombination equation (forward in time)
2. Its solution via genealogical thinking

Recombination



Sequences and selection

individual: sequence of n sites, $S = \{1, \dots, n\}$

types: $x := (x_1, \dots, x_n) \in \prod_{i \in S} X_i =: X$, $X_i = \{0, 1\}$

marginal types: $x_U := (x_i)_{i \in U}$, $U \subseteq S$

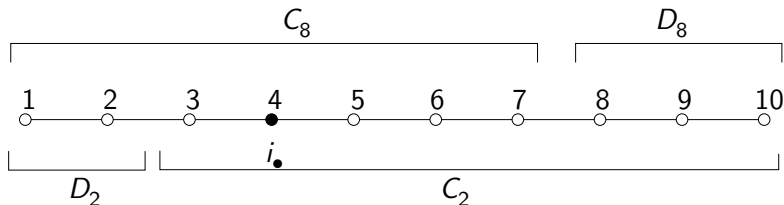
reproduction and selection:

- single selected site: $i_\bullet \in S$ (all others 'neutral')
- types x with $x_{i_\bullet} = 1$: reproduce at rate 1 ('bad, unfit')
- types x with $x_{i_\bullet} = 0$: reproduce at rate $1 + s$ ('good, fit')
- offspring replaces randomly chosen individual

Ordering the sites

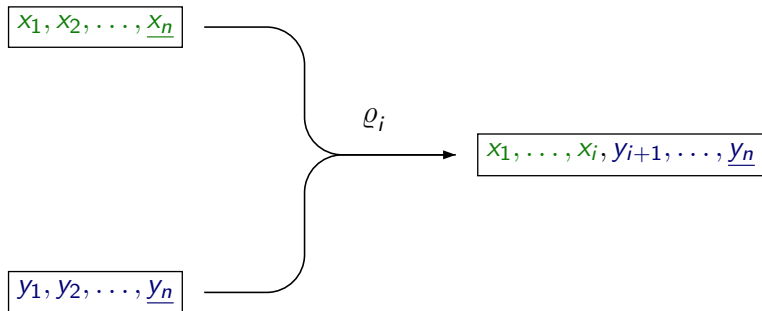
partial order on S :

- $i \preceq j$ if either $i_\bullet \leq i \leq j$ or $i_\bullet \geq i \geq j$
- $i \prec j$ if $i \preceq j$ and $i \neq j$
- $D_i := \{j \in S \mid i \preceq j\}$ (i -tail), $C_i := S \setminus D_i$ (i -head), $i \in S \setminus i_\bullet$



- $\rightsquigarrow i_\bullet \in C_i$
- (single-crossover) recombination between C_i and D_i (partitions S into C_i and D_i) at rate ρ_i

Sequences and recombination

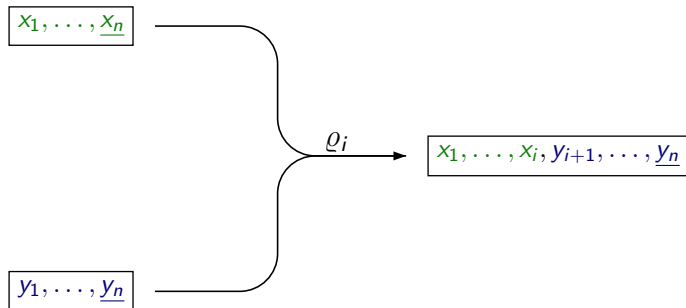


Selection-recombination equation

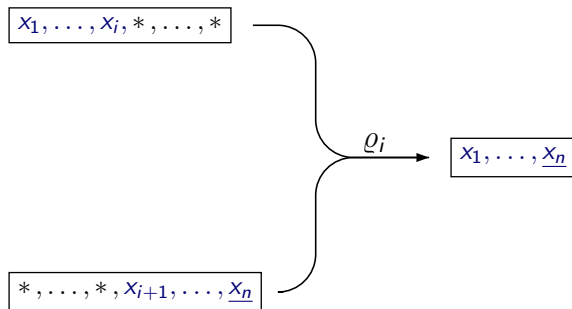
- $\omega_t = (\omega_t(x))_{x \in X}$ probability measure on X
($\omega_t(x)$ proportion of individuals of type x at time t , $x \in X$)
- solves **selection-recombination equation (SRE)**

$$\dot{\omega}_t(x) = s[\delta_{x_{i_\bullet, 0}} - \omega_t(*, 0, *)] \omega_t(x) + \sum_{i \in S \setminus i_\bullet} \varrho_i [\omega_t(x_{C_i}, *) \omega_t(*, x_{D_i}) - \omega_t(x)], \quad x \in X$$

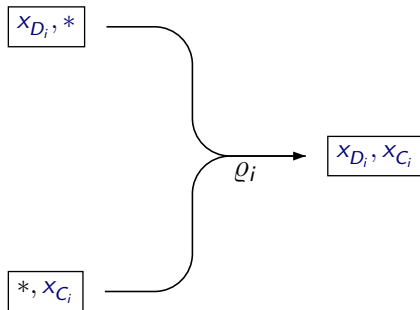
Recombination



Recombination



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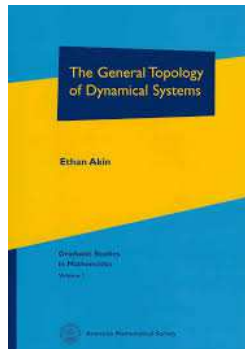
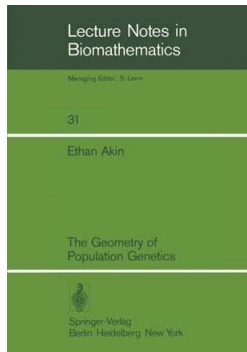
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Research context

- **hitchhiking**: neutral x_i ($i \neq i_\bullet$) selectively favoured if linked with $x_{i_\bullet} = 0$
- **Akin 1979**: “The differential equations which model the action of selection and recombination are nonlinear equations which are impossible to solve explicitly.”



Research context

- **hitchhiking**: neutral x_i ($i \neq i_\bullet$) selectively favoured if linked with $x_{i_\bullet} = 0$
- **Akin 1979**: the “selection-recombination equation is a large, nonlinear system of differential equations that are impossible to solve explicitly”
- **Stephan, Song, Langley 2006**: approximate solution ($n = 3$, wild approximations, technical result, numerical evaluation only, not generalisable)

Recombinators

- canonical projection: for $\emptyset \neq U \subseteq S$,

$$\pi_U : X \rightarrow \prod_{i \in U} X_i = X_U, \quad \pi_U(x) = (x_i)_{i \in U} = x_U$$

- marginal measure wrt sites in U : for $\nu \in \mathcal{P}(X)$,

$$\pi_U.\nu = \nu \circ \pi_U^{-1} =: \nu_U$$

type distribution of sites in U

for $x_U \in X_U$: $\nu_U(x_U) = \nu(x_U, *)$

- recombinator: for $i \in S \setminus i_\bullet$,

$$\mathcal{P}(X) \longrightarrow \mathcal{P}(X)$$

$$R_i(\nu) := \nu_{C_i} \otimes \nu_{D_i}$$

$$(R_i(\nu))(x) = \nu(x_{C_i}, *) \cdot \nu(*, x_{D_i})$$

distribution of sequences pieced together from random i -heads and i -tails

Selection-recombination equation

$$\dot{\omega}_t(x) = s[\delta_{x_{i_\bullet,0}} - \omega_t(*, 0, *)] \omega_t(x) + \sum_{i \in S \setminus i_\bullet} \varrho_i [\omega_t(x_{C_i}, *) \omega_t(*, x_{D_i}) - \omega_t(x)], \quad x \in X$$

or

$$\dot{\omega}_t = s(F - f(\omega_t))\omega_t + \sum_{i \in S \setminus i_\bullet} \varrho_i (R_i - \mathbb{1})\omega_t = \Psi_{\text{sel}}(\omega_t) + \Psi_{\text{rec}}(\omega_t)$$

with

$$F\omega(x) = \delta_{x_{i_\bullet,0}}\omega(x)$$

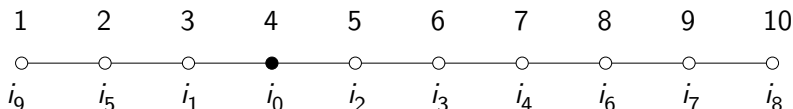
and

$$f(\omega) := \omega_{\{i_\bullet\}}(0)$$

(proportion of fit individuals = proportion of individuals with 0 at selected site)

Recursive solution

- $(i_k)_{0 \leq k < n}$ non-decreasing permutation of S (sensu \Leftarrow)
- $i_0 = i_\bullet$
- $C^{(k)} = C_{i_k}, D^{(k)} = D_{i_k}, \varrho^{(k)} := \varrho_{i_k} \quad (k > 0)$



hierarchy of solutions:

- $(\omega_t^{(0)})_{t \geq 0}$ solution of SRE with $\varrho^{(\ell)} = 0$ for all ℓ
(pure selection equation)
- $(\omega_t^{(k)})_{t \geq 0}$ solution of **SRE truncated at k**
(with $\varrho^{(\ell)} = 0$ for all $\ell > k$), $1 \leq k < n$

Recursive solution

Theorem

The solution of the pure selection equation is given by

$$\omega_t^{(0)} = \frac{e^{stF} \omega_0}{e^{st} f(\omega_0) + 1 - f(\omega_0)} =: \varphi_t(\omega_0),$$

and the solutions $\omega_t^{(k)}$ can be computed via the recursion

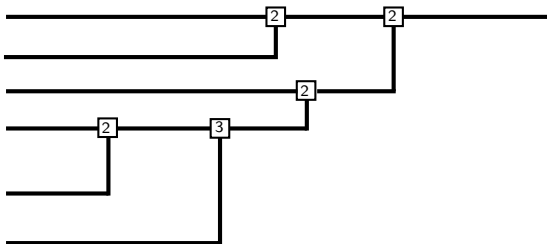
$$\omega_t^{(k)} = e^{-\varrho^{(k)} t} \omega_t^{(k-1)} + \omega_{C^{(k)},t}^{(k-1)} \otimes \int_0^t \varrho^{(k)} e^{-\varrho^{(k)} \tau} \omega_{D^{(k)},\tau}^{(k-1)} d\tau.$$

analytical proof ✓

(genealogical) meaning?

Ancestral graph

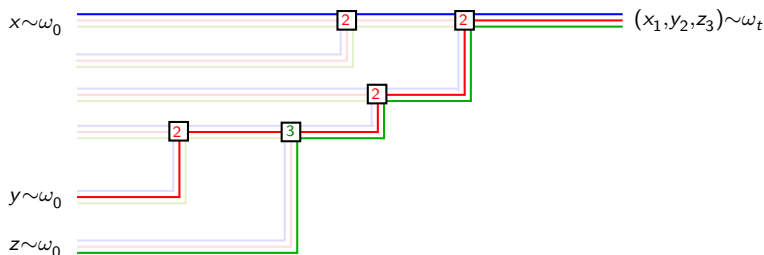
i -splitting events on every line at rate ρ_i , $i \in S \setminus i_\bullet$ (backward!)
 $S = \{1, 2, 3\}$, $i_\bullet = 1$:



Ancestral graph

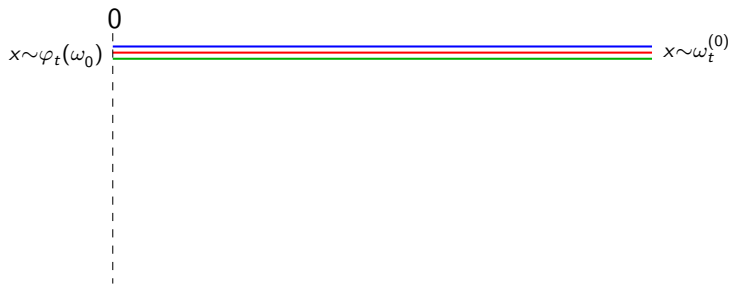
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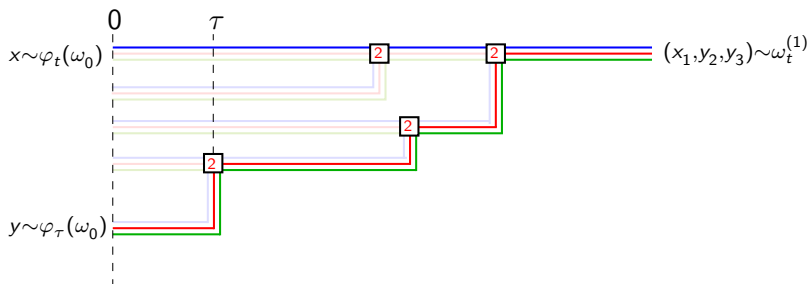
$s = 0$: partitioning of sequence across ancestors
sampling from ω_0

Ancestral graph, $s > 0$



$$\omega_t^{(0)} = \varphi_t(\omega_0)$$

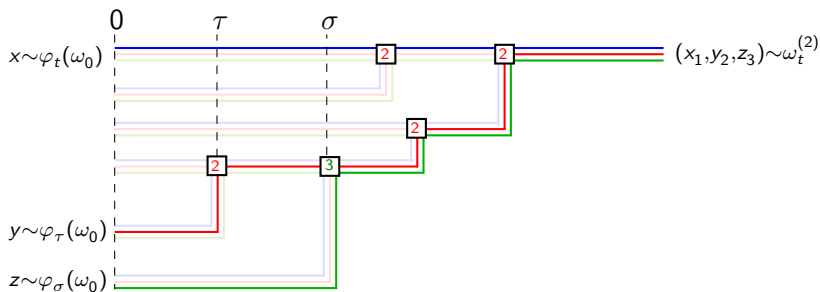
Ancestral graph, $s > 0$



decomposition according to **leftmost** 2-event on ancestral line of tail (since $\Psi_{\text{sel}}(\nu_{\{i_\bullet\}} \otimes \nu_{\{S \setminus i_\bullet\}}) = \Psi_{\text{sel}}(\nu_{\{i_\bullet\}}) \otimes \nu_{\{S \setminus i_\bullet\}}$)

$$\begin{aligned} \rightsquigarrow \omega_t^{(1)} &= e^{-\varrho_2 t} \varphi_t(\omega_0) + \varphi_{\{1\},t}(\omega_0) \otimes \int_0^t \varrho_2 e^{-\varrho_2 \tau} \varphi_{\{2,3\},\tau}(\omega_0) d\tau \\ &= e^{-\varrho^{(1)} t} \omega_t^{(0)} + \omega_{C^{(1)},t}^{(0)} \otimes \int_0^t \varrho^{(1)} e^{-\varrho^{(1)} \tau} \omega_{D^{(1)},\tau}^{(0)} d\tau \end{aligned}$$

Ancestral graph, $s > 0$



decomposition according to **leftmost** k -event on ancestral line of $D^{(k)}$

$$\rightsquigarrow \omega_t^{(k)} = e^{-\varrho^{(k)}t} \omega_t^{(k-1)} + \omega_{C^{(k-1)},t}^{(k-1)} \otimes \int_0^t \varrho^{(k)} e^{-\varrho^{(k)}\sigma} \omega_{D^{(k-1)},\sigma}^{(k-1)} d\sigma$$