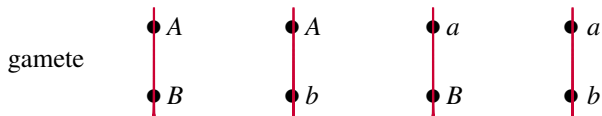


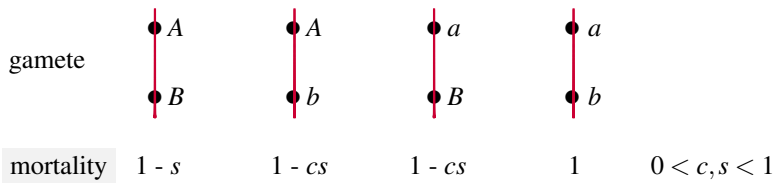
Probability of fixation of a new mutant and
the ancestral recombination-selection graph:
Application to the Hill-Robertson effect

Sabin Lessard and Amir Kermany
Université de Montréal

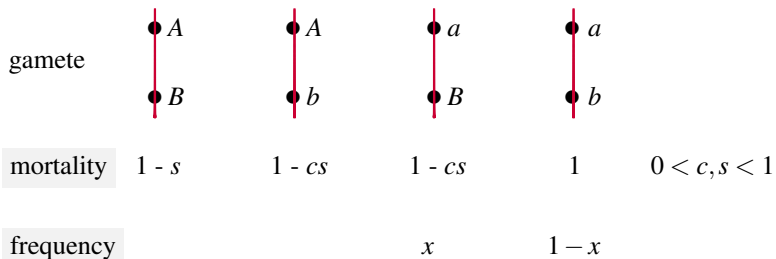
Two-locus selection model







Two-locus selection model



Two-locus selection model



Two-locus selection model

gamete					
mortality	$1 - s$	$1 - cs$	$1 - cs$	1	$0 < c, s < 1$
frequency	ϵ	0	$x - \epsilon$	$1 - x$	probability x
	0	ϵ	x	$1 - x - \epsilon$	probability $1 - x$

Linkage disequilibrium

$$D = x_{AB} - x_A x_B = (\epsilon - \epsilon x)x + (-\epsilon x)(1 - x) = 0$$

Linkage disequilibrium

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Epistasis

positive if $1 - s < (1 - cs)^2$

negative if $1 - s > (1 - cs)^2$

Recombination

Recombination



Negative linkage disequilibrium

Recombination



Negative linkage disequilibrium



Negative epistasis
in an infinite population



Random drift
in a finite population
(Hill and Robertson 1966)

Moran model for population of size N

- ▶ At each time step, one offspring produced by two individuals at random
- ▶ Recombinant offspring with probability $r = \frac{\rho}{N}$

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and then with conditional probability

$$\left\{ \begin{array}{ll} 0 & \text{if } AB \\ 1 - c & \text{if } Ab \text{ or } aB \\ 1 & \text{if } ab \end{array} \right.$$

Ancestral recombination-selection graph (ARSG)

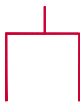
Backwards in time with $\frac{N^2}{2}$ time steps as unit of time as $N \rightarrow \infty$



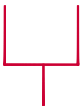
coalescence C of each pair of lineages at rate 1

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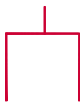
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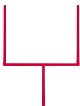
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Ancestral recombination-selection graph (ARSG)

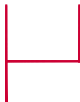
Backwards in time with $\frac{N^2}{2}$ time steps as unit of time as $N \rightarrow \infty$



coalescence C of each pair of lineages at rate 1



recombination R of each lineage at rate $\frac{\rho}{2}$



selection S of each lineage at rate $\frac{\sigma}{2}$

Probability of fixation of A

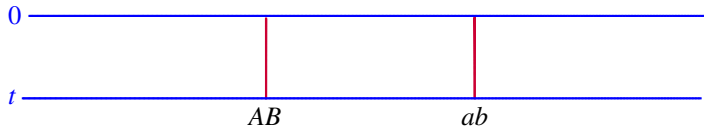
$$x_A(0) + \frac{\sigma}{N^2} \sum_{\tau \geq 0} E[x_{AB}(\tau)x_{ab}(\tau) + cx_{Ab}(\tau)x_{ab}(\tau) + (1-c)x_{AB}(\tau)x_{aB}(\tau)]$$

Calculation

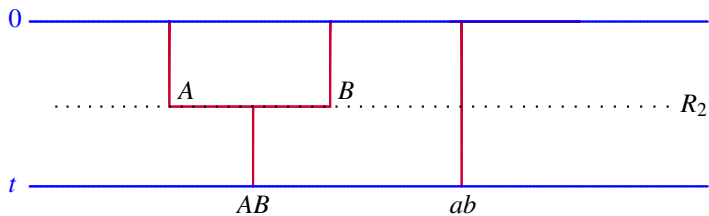
$$\frac{2}{N^2} \sum_{\tau \geq 0} E[x_{AB}(\tau)x_{ab}(\tau)] \rightarrow \int_0^{\infty} E[x_{AB}(t)x_{ab}(t)] dt$$

Calculation

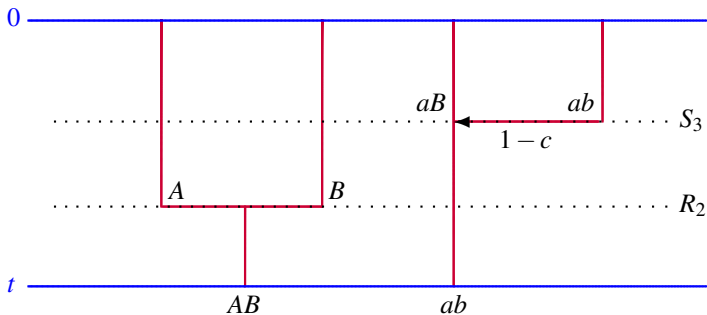
$$\begin{aligned} \frac{2}{N^2} \sum_{\tau \geq 0} E[x_{AB}(\tau)x_{ab}(\tau)] &\rightarrow \int_0^\infty E[x_{AB}(t)x_{ab}(t)] dt \\ &\approx E(T_2)x_{AB}(0)x_{ab}(0) + \end{aligned}$$



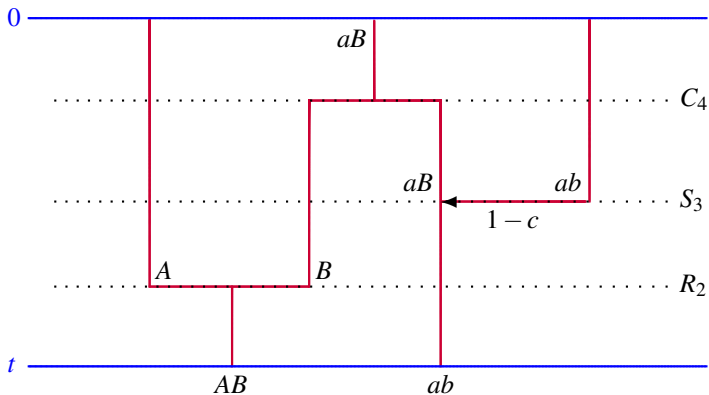
$$P(R_2)E(T_3)x_A(0)x_B(0)x_{ab}(0)+$$



$$(1 - c)P(R_2)P(S_3)E(T_4)x_A(0)x_B(0)x_{aB}(0)x_{ab}(0) +$$



$$(1 - c)P(R_2)P(S_3)P(C_4)E(T_3)x_A(0)x_{aB}(0)x_{ab}(0) +$$



Result with positive epistasis ($c < \frac{1}{2}$)

$$P(A \text{ fixation}) \approx \varepsilon + \frac{\varepsilon\sigma}{2}(c + x(1 - 2c)) \\ + \frac{\varepsilon\sigma^2}{12}(c^2 + x(1 - 2c)(1 + 2c(1 - x)))$$

Result with positive epistasis ($c < \frac{1}{2}$)

$$\begin{aligned} P(\text{A fixation}) \approx \varepsilon &+ \frac{\varepsilon\sigma}{2}(c + x(1 - 2c)) \\ &+ \frac{\varepsilon\sigma^2}{12}(c^2 + x(1 - 2c)(1 + 2c(1 - x))) \\ &+ \frac{\varepsilon\rho\sigma^2}{432}x(1 - x)(1 - 2c)(3 - c) \end{aligned}$$

Result with no epistasis ($c = \frac{1}{2}$)

$$P(A \text{ fixation}) \approx \varepsilon + \frac{\varepsilon\sigma}{4} + \frac{\varepsilon\sigma^2}{48} - \frac{\varepsilon\sigma^3}{192}x(1-x)$$

Result with no epistasis ($c = \frac{1}{2}$)

$$\begin{aligned} P(A \text{ fixation}) \approx \varepsilon &+ \frac{\varepsilon\sigma}{4} + \frac{\varepsilon\sigma^2}{48} \\ &- \frac{\varepsilon\sigma^3}{192}x(1-x) \\ &+ \frac{19\varepsilon\rho\sigma^3}{3456}x(1-x) \end{aligned}$$

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- ▶ The analysis confirms the Hill-Robertson effect in favor of recombination

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- ▶ The analysis confirms the Hill-Robertson effect in favor of recombination
- ▶ The approach allows us to get approximations of any order with respect to σ and ρ , and this with relative ease
- ▶ The results are valid for a wide class of models and can be extended to other classes

Thanks!

