# Ancestral recombination-selection graph and fixation probability 

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IMS 2010 Gothenburg
9-13 August

# Ancestral recombination-selection graph 

## and fixation probability

Application to the Hill-Robertson effect

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## Two-locus selection model



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## Two-locus selection model

| type | $\phi_{\phi} A$ | $\phi_{b} A$ | $\phi^{\bullet} a$ | $\phi_{b} a$ |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
| mortality | $1-s$ | $1-c s$ | $1-c s$ | 1 | $0<c, s<1$ |
| frequency | $\varepsilon$ | 0 | $x-\varepsilon$ | $1-x$ | probability $x$ |
|  | 0 | $\varepsilon$ | $x$ | $1-x-\varepsilon$ | probability $1-x$ |

## Linkage disequilibrium

$$
D=x_{A B}-x_{A} x_{B}=(\varepsilon-\varepsilon x)(x)+(-\varepsilon x)(1-x)=0
$$

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$$

## Epistasis

positive $A B$ fitter than expected $(1-s)<(1-c s)^{2}$
negative $A B$ less fit than expected $(1-s)>(1-c s)^{2}$
absent $A B$ as fit as expected $(1-s)=(1-c s)^{2}$

## Recombination

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Negative linkage disequilibrium $D<0$

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Negative linkage disequilibrium $D<0$


Negative epistasis
in an infinite population

## Recombination



Negative linkage disequilibrium $D<0$


Negative epistasis
in an infinite population

Random drift in a finite population

## Recombination makes more likely

## the fixation of beneficial mutants

## in finite populations

## Moran model for population of size $N$

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- One individual at random to be replaced by the offspring
- Replacement in all cases with probability $1-s=1-\frac{\sigma}{N}$
- Type-specific replacement with probability $s=\frac{\sigma}{N}$ and then with conditional probability

$$
\begin{cases}0 & \text { if } A B \\ 1-c & \text { if } A b \text { or } a B \\ 1 & \text { if } a b\end{cases}
$$

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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 (Kingman 1982)

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selection $S$ of each lineage at rate $\frac{\sigma}{2}$ (Krone and Neuhauser 1997)

## Probability of fixation of $A$

$$
x_{A}(0)+\sum_{\tau \geq 0} E\left[x_{A}(\tau+1)-x_{A}(\tau)\right]
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$$
x_{A}(0)+\frac{\sigma}{N^{2}} \sum_{\tau \geq 0} E\left[x_{A B}(\tau) x_{a b}(\tau)+c x_{A b}(\tau) x_{a b}(\tau)+(1-c) x_{A B}(\tau) x_{a B}(\tau)\right]
$$

## Calculation

$$
\frac{2}{N^{2}} \sum_{\tau \geq 0} E\left[x_{A B}(\tau) x_{a b}(\tau)\right] \rightarrow \int_{0}^{\infty} E\left[x_{A B}(t) x_{a b}(t)\right] d t
$$

$E\left[x_{A B}(t) x_{a b}(t)\right]=P(A B$ and $a b$ in this order at time $t)$
where $t$ is for time in units of $\frac{N^{2}}{2}$ time steps as $N \rightarrow \infty$

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$$
E\left(T_{2}\right) x_{A B}(0) x_{a b}(0)+
$$



$$
P\left(R_{2}\right) E\left(T_{3}\right) x_{A}(0) x_{B}(0) x_{a b}(0)+
$$



$$
(1-c) P\left(R_{2}\right) P\left(S_{3}\right) E\left(T_{4}\right) x_{A}(0) x_{B}(0) x_{a B}(0) x_{a b}(0)+
$$



$$
(1-c) P\left(R_{2}\right) P\left(S_{3}\right) P\left(C_{4}\right) E\left(T_{3}\right) x_{A}(0) x_{a B}(0) x_{a b}(0)+\cdots
$$



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

$$
\begin{aligned}
P(A \text { fixation }) \approx \varepsilon & +\frac{\varepsilon \sigma}{2}(c+x(1-2 c)) \\
& +\frac{\varepsilon \sigma^{2}}{12}\left(c^{2}+x(1-2 c)(1+2 c(1-x))\right) \\
& -\frac{\varepsilon \sigma^{3}}{24} x(1-x)(c+x(1-2 c))
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& -\frac{\varepsilon \sigma^{3}}{24} x(1-x)(c+x(1-2 c)) \\
& +\frac{\varepsilon \rho \sigma^{2}}{432} x(1-x)(1-2 c)(3-c)
\end{aligned}
$$

positive term in $\rho$

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

$$
\begin{aligned}
P(A \text { fixation }) \approx \varepsilon & +\frac{\varepsilon \sigma}{4}+\frac{\varepsilon \sigma^{2}}{48}-\frac{\varepsilon \sigma^{3}}{192} x(1-x) \\
& -\frac{\varepsilon \sigma^{4}}{11520}\left(1+15 x-29 x^{2}+14 x^{3}\right)
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& -\frac{\varepsilon \sigma^{4}}{11520}\left(1+15 x-29 x^{2}+14 x^{3}\right) \\
& +\frac{19 \varepsilon \rho \sigma^{3}}{3456} x(1-x)
\end{aligned}
$$

## positive term in $\rho$

## Final comments

- The analysis confirms the Hill-Robertson effect in favor of recombination in finite populations with positive or no epistasis


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- branching processes (Barton 1995)
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- The results are valid for a wide class of models (exchangeable in the realm of the Kingman coalescent) and can be extended to other classes (e.g. lambda coalescent)
- The same approach can be used to study factors of evolution in multilocus models


## Thanks!


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