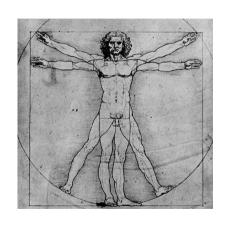
Computer simulations The genomes of recombinant inbred lines

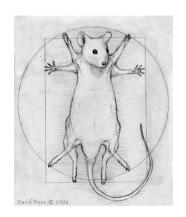
Karl Broman

Biostatistics & Medical Informatics, UW-Madison

kbroman.org
github.com/kbroman
@kwbroman
Course web: kbroman.org/AdvData

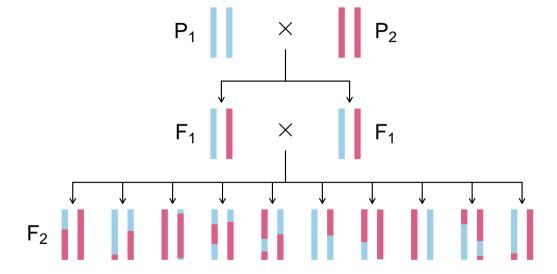




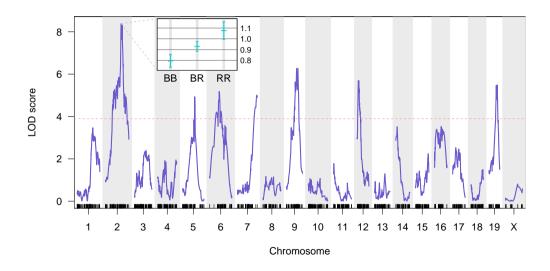


daviddeen.com

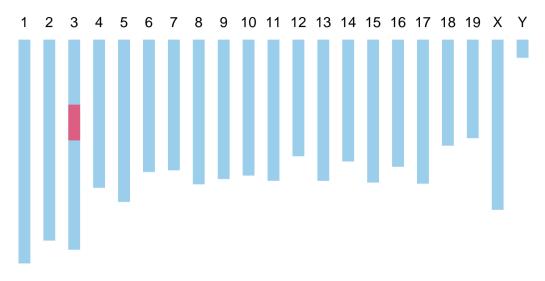
Intercross



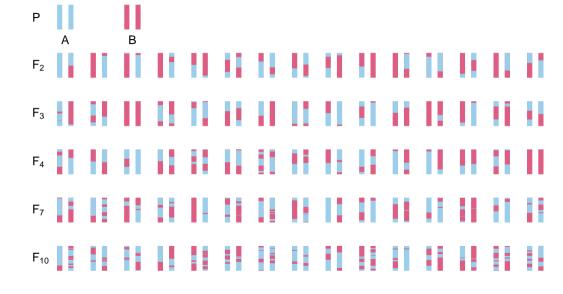
QTL mapping



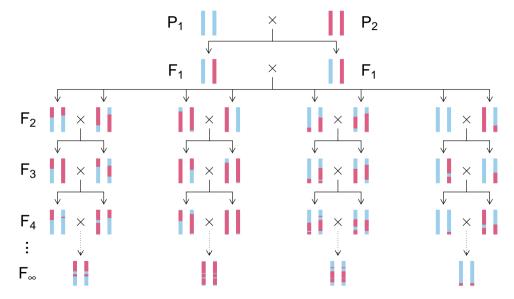
Congenic line



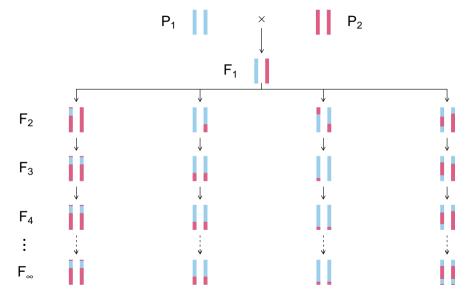
Advanced intercross lines



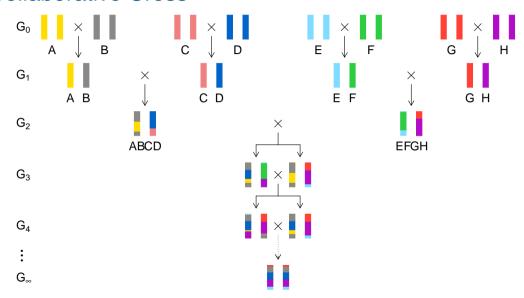
Recombinant inbred lines



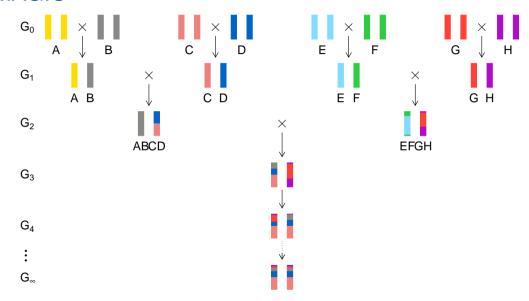
Recombinant inbred lines



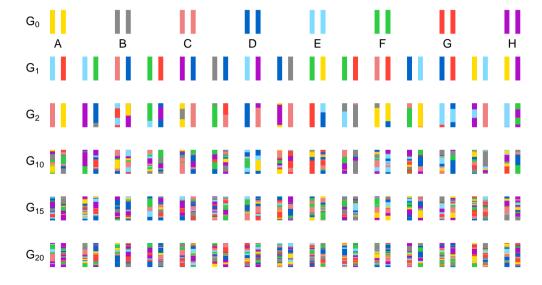
Collaborative Cross



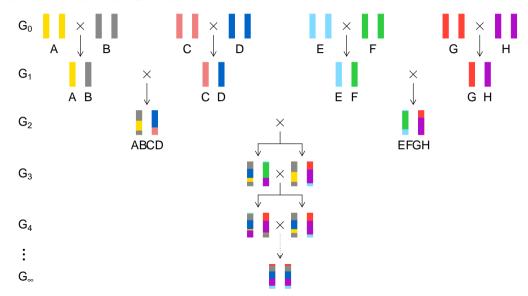
MAGIC

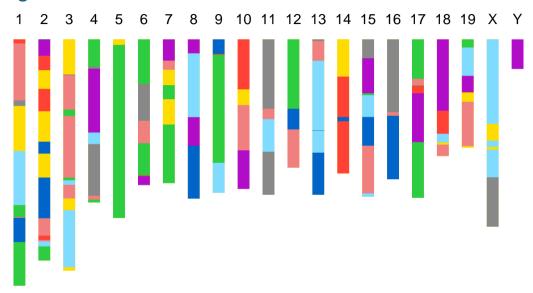


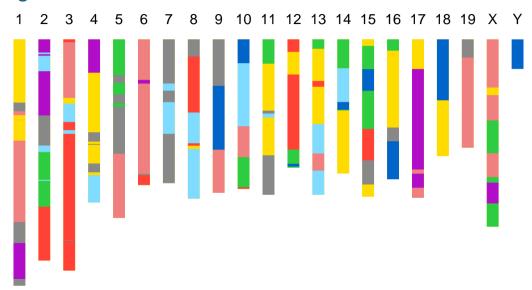
Heterogeneous stock

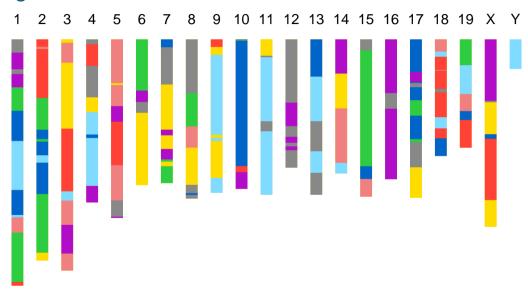


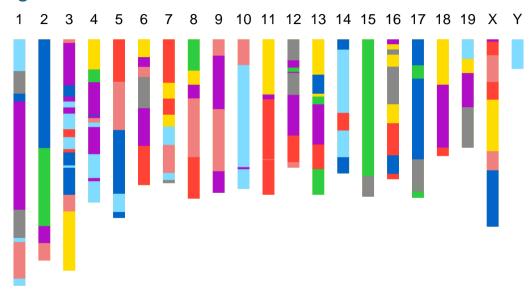
Collaborative Cross

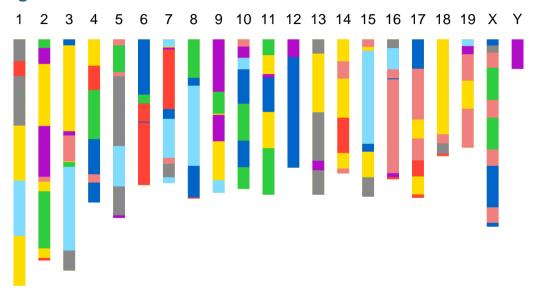








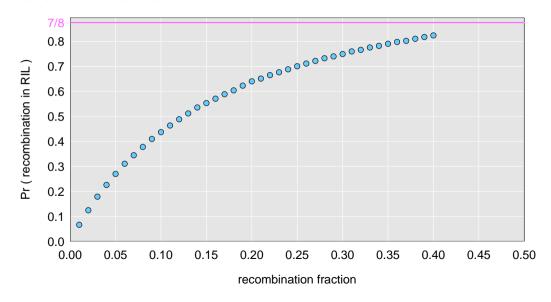




Recombination fraction

r is the "recombination fraction"

Simulation results



Haldane & Waddington 1931

INBREEDING AND LINKAGE*

J. B. S. HALDANE AND C. H. WADDINGTON

John Innes Horticultural Institution, London, England

Received August 9, 1930

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Result for selfing

Then
$$c_n + \lambda d_n \equiv c_n + \frac{1}{4}(1 - 2x)d_n + \frac{1}{2}\lambda(1 - 2x)d_n$$

$$\therefore \ \lambda = \frac{1 - 2x}{2 + 4x} \cdot$$

Then since $d_{\infty} = 0$, and $c_1 = 0$, $d_1 = 2$,

$$c_{\infty} = c_{\infty} + \lambda d_{\infty} = c_1 + \lambda d_1 = \frac{1 - 2x}{1 + 2x}$$

Put $y = D_{\infty}$ (the final proportion of crossover zygotes)

$$\therefore C_{\infty} + D_{\infty} = 1, C_{\infty} - D_{\infty} = c_{\infty} \therefore y = \frac{1}{2}(1 - c_{\infty}).$$

$$\therefore y = \frac{2x}{1 + 2x}.$$
(1.3)

Result for sib-mating

Omitting some rather tedious algebra, the solution of these equations is:

$$\zeta = \frac{q}{2 - 3q}, \quad \theta = \frac{2q}{2 - 3q}, \quad \kappa = \frac{1}{2 - 3q},$$

$$\lambda = \frac{1 - 2q}{2 - 3q}, \quad \mu = \frac{1 - 2q}{2 - 3q}, \quad \nu = \frac{2q}{2 - 3q}$$

as may easily be verified.

$$c_{\infty} = c_{n} + 2e_{n} + \frac{1}{1 + 6x} [(1 - 2x)(d_{n} + 2f_{n} + 2j_{n} + \frac{1}{2}k_{n}) + 2g_{n} + 4x(h_{n} + i_{n})]$$
(3.4)

and $y = \frac{1}{2}(1 - c_{\infty})$.

In the case considered, $d_0 = 1$, $c_{\infty} = \zeta d_0 = 1 - 2x/1 + 6x$. Hence the proportion of crossover zygotes, y = 4x/1 + 6x (3.5).

Result for sib-mating

Omitting some rather tedious algebra, the solution of these equations is:

$$\zeta = \frac{q}{2 - 3q}, \quad \theta = \frac{2q}{2 - 3q}, \quad \kappa = \frac{1}{2 - 3q},$$

$$\lambda = \frac{1 - 2q}{2 - 3q}, \quad \mu = \frac{1 - 2q}{2 - 3q}, \quad \nu = \frac{2q}{2 - 3q}$$

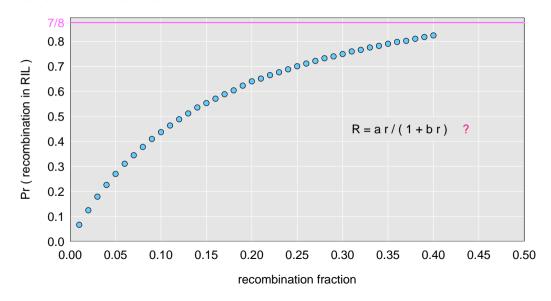
as may easily be verified.

$$\therefore c_{\infty} = c_{n} + 2e_{n} + \frac{1}{1 + 6x} [(1 - 2x)(d_{n} + 2f_{n} + 2j_{n} + \frac{1}{2}k_{n}) + 2g_{n} + 4x(h_{n} + i_{n})]$$
(3.4)

and $y = \frac{1}{2}(1 - c_{\infty})$.

In the case considered, $d_0 = 1$, $c_\infty = \zeta d_0 = 1 - 2x/1 + 6x$. Hence the proportion of crossover zygotes, y = 4x/1 + 6x (3.5).

Simulation results



Non-linear regression

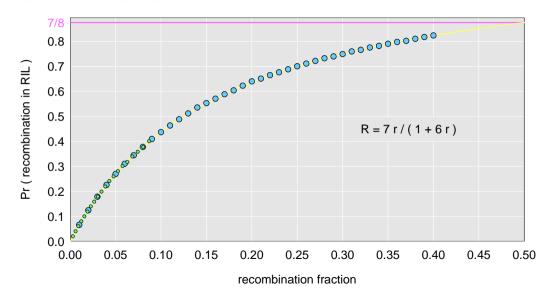
Non-linear regression

```
out <- nls(R \sim a*r/(1 + b*r),
          data = data.frame(r=r, R=R),
          start = list(a=4, b=6)
summary(out)
 Estimate Std. Error
a 7.016 0.011
b
 6.023
               0.016
```

Non-linear regression

					More	data	
	Estimate	Std.	Error		Estimate	Std.	Error
a	7.016		0.011	a	7.003		0.008
b	6.023		0.016	b	6.005		0.012

Simulation results



Markov chain

▶ Sequence of random variables $\{X_0, X_1, X_2, ...\}$ satisfying

$$Pr(X_{n+1}\mid X_0,X_1,\ldots,X_n)=Pr(X_{n+1}\mid X_n)$$

- ▶ Transition probabilities $P_{ij} = Pr(X_{n+1} = j \mid X_n = i)$
- ► Here, X_n = "parental type" at generation n.
- ▶ We are interested in absorption probabilities

$$\pi_j = \Pr(\mathsf{X}_\mathsf{n} \to \mathsf{j} \mid \mathsf{X}_\mathsf{0})$$

Absorption probabilities

Consider the case of absorption into the state
$$\begin{bmatrix} A & A \\ A & A \end{bmatrix}$$
 (write this AA|AA)

Let h_i = probability, starting at i, of being absorbed into AA|AA.

Then $h_{AA|AA} = 1$ and $h_{AB|AB} = 0$.

Condition on the first step: $h_i = \sum_k P_{ik} h_k$

For selfing, this gives a system of 3 linear equations.

Equations for selfing

Cn AABB and aabb.

D. AAbb and aaBB.

 E_n AABb, AaBB, Aabb, and aaBb.

Fn AB.ab.

Gn Ab.aB.

We assume $2C_n+2D_n+4E_n+F_n+G_n=2$, so that $C_1=D_1=E_1=G_1=0$, and $F_1=2$. Clearly $E_{\infty}=F_{\infty}=G_{\infty}=0$, and D_{∞} is the final proportion of crossover zygotes. Then considering the results of selfing each generation, we have:

$$\begin{array}{l} C_{n+1} = C_n + \frac{1}{2} E_n + \frac{1}{4} (1 - \beta - \delta + \beta \delta) F_n + \frac{1}{4} \beta \delta G_n \\ D_{n+1} = D_n + \frac{1}{2} E_n + \frac{1}{4} \beta \delta F_n + \frac{1}{4} (1 - \beta - \delta + \beta \delta) G_n \\ E_{n+1} = \frac{1}{2} E_n + \frac{1}{4} (\beta + \delta - 2\beta \delta) (F_n + G_n) \\ F_{n+1} = \frac{1}{2} (1 - \beta - \delta + \beta \delta) F_n + \frac{1}{2} \beta \delta G_n \\ G_{n+1} = \frac{1}{2} \beta \delta F_n + \frac{1}{2} (1 - \beta - \delta + \beta \delta) G_n \end{array}$$

or C_{n+1} , D_{n+1} , and F_{n+1} , G_{n+1} ,

$$d_n$$
 (1.2)

all values of n.
2x)dn

$$=\frac{1-2x}{1+2x}.$$

Put $y = D_{\infty}$ (the final proportion of crossover zygotes)

$$\therefore C_{\infty} + D_{\infty} = 1, C_{\infty} - D_{\infty} = c_{\infty} \therefore y = \frac{1}{2}(1 - c_{\infty}).$$

$$\therefore y = \frac{2x}{1 + 2x}.$$
(1.3)

Equations for sib-mating

Typical mating	Number of types				
$AABB \times AABB$	2	$C_{n+1} = C_n + H + U + \frac{1}{8}(\beta^2 + \delta^2)$			
$AAbb{\times}AAbb$	2		$\{(\alpha^2 + \gamma^2)M + \frac{1}{4}(\beta^2 + \delta^2)P + \frac{1}{16}\beta^2\delta^2W + \frac{1}{16}(\alpha^2\delta^2 + \delta^2)P + \frac{1}{16}(\alpha^2$		
$AABB \times aabb$	2	$E_{n+1} = \frac{1}{16}\alpha^2 \gamma^2 W$	$+\frac{1}{16}(\alpha^2\delta^2+\beta^2\gamma^2)X+\frac{1}{16}\beta$	βδ²Y.	
$AAbb \times aaBB$	2	$F_{n+1} = \frac{1}{16} \beta^2 \delta^2 W$	$+\frac{1}{16}(\alpha^2 5^2 + \beta^2 \gamma^2)X + \frac{1}{16}\alpha$	$^{2}\gamma^{2}Y$.	
$AABB \times AAbb$	8	$G_{n+1} = \frac{1}{16}(\alpha\beta +$	$\gamma \delta)(U+V) + \frac{1}{16}\alpha\beta\gamma\delta(W-V)$	+2X+Y).	
$AABB \times AABb$	8	$H_{n+1} = \frac{1}{2}H_{1}$	01 00 130 110 11	(* 1 2 2)	1.1.1.1.1
$AAbb \times AABb$	8	$U + \frac{1}{16}($ $(\alpha \delta + \beta \cdot $ $I_{n+1} = \frac{1}{2}I +$	Typical mating $AABB \times Ab.aB$	Number of types	$N_{n+1} = \frac{1}{4}R + \frac{1}{4}(\alpha\beta + \gamma\delta)(U+V) + \frac{1}{4}\alpha\beta\gamma\delta(W+2X+Y).$
		$U + \frac{1}{16}($ $(\alpha \delta + \beta -$	$AAbb \times AB.ab$	4	$P_{\alpha+1} = \frac{1}{2}S + \frac{1}{2}(\alpha\beta + \gamma\delta)(U+V) + \frac{1}{2}\alpha\beta\gamma\delta(W+2X+Y).$
$AABB \times Aabb$	8	$J_{n+1} = \frac{1}{16} (\alpha \delta - \beta \delta) (\alpha \delta - \beta \delta)$	$AABb \times AABb$	4	$Q_{a+1} = 2G + \frac{1}{2}(H + I + J + K) + \frac{1}{4}(\alpha^2 + \gamma^2)(L + M) + \frac{1}{2}(\beta^2 + \delta^2)$ $(N + P) + \frac{1}{4}Q + \frac{1}{6}(R + S + T) + \frac{1}{4}(\alpha^2 + \alpha\beta + \beta^2 + \gamma^2 + \gamma^5 + \delta^2)$ $(U + V) + \frac{1}{4}(\alpha\delta + \beta\gamma)^2(W + Y) + \frac{1}{6}(\alpha\gamma + \beta\delta)^3X.$
$AAbb \times AaBB$	8	$K_{n+1} = \frac{1}{16}$ $\beta \delta)(\alpha \delta - \frac{1}{16})$	$AABb{\times}AaBB$	4	$(U+V)+\frac{1}{3}(\alpha\alpha+\beta\gamma)^2(W+1)+\frac{1}{3}(\alpha\gamma+\beta\beta)^{2}\lambda^{2}$ $R_{n+1}=\frac{1}{2}(\beta^{2}+\delta^{2})L+\frac{1}{2}(\alpha^{2}+\gamma^{2})N+\frac{1}{8}R+\frac{1}{8}(\beta+\delta)U+\frac{1}{8}(\alpha+\gamma)V+\frac{1}{3}(\alpha\delta+\beta\gamma)^{2}(W+Y)+\frac{1}{8}(\alpha\gamma+\beta\beta)^{3}X.$
$AABB \times AB.ab$	4	$L_{n+1} = \frac{1}{4}(\alpha + \alpha^2 \gamma^2 W - \alpha^2 \gamma^$	$AABb{\times}Aabb$	4	$S_{n+1} = \frac{1}{4}(\beta^2 + \delta^3)M + \frac{1}{4}(\alpha^2 + \gamma^2)P + \frac{1}{8}S + \frac{1}{8}(\alpha + \gamma)U + \frac{1}{8}(\beta + \delta)V + \frac{1}{16}$ $(a\delta + \beta\gamma)^2(W + Y) + \frac{1}{8}(\alpha\gamma + \beta\delta)^2X.$
$AAbb \times Ab.aB$	4	$M_{n+1} = \frac{1}{4}$	m	1.0	$T_{n+1} = \frac{1}{6}(\alpha\beta + \gamma\delta)(U+V) + \frac{1}{16}(\alpha\delta + \beta\gamma)^{3}(W+Y) + \frac{1}{6}(\alpha\gamma + \beta\delta)^{3}X.$
		β°82W-1	$AABb \times aaBb$		$U_{n+1} = \frac{1}{2}I + \frac{1}{2}(\alpha\beta + \gamma\delta)(U + V) + \frac{1}{2}(\alpha\delta + \beta\gamma)(W + V) + \frac{1}{8}(\alpha\gamma + \beta\delta)U + \frac{1}{8}(\beta\gamma + \delta)$
			$AABb \times AB.ab$	8	$V + \frac{1}{4}\alpha\gamma(\beta\gamma + \alpha\delta)W + \frac{1}{4}(\alpha\gamma + \beta\delta)(\alpha\delta + \beta\gamma)X + \frac{1}{4}\beta\delta(\beta\gamma + \alpha\delta)Y$.
			$AABb \times Ab.aB$	8	$V_{n+1} = \frac{1}{2}K + \frac{1}{4}(\alpha\beta + \gamma\delta)(M+P) + \frac{1}{8}(R+T) + \frac{1}{8}(\beta + \delta)U + \frac{1}{8}(\alpha + \gamma)$ $V + \frac{1}{8}\beta\delta(\beta\gamma + \alpha\delta)W + \frac{1}{8}(\alpha\gamma + \beta\delta)(\alpha\delta + \beta\gamma)X + \frac{1}{8}\alpha\gamma(\beta\gamma + \alpha\delta)Y.$
			$AB.ab \times AB.ab$	1	$W_{n+1} = 2(E+J) + \frac{1}{2}(\alpha^2 + \gamma^2)L + \frac{1}{2}(\beta^2 + \delta^2)N + \frac{1}{4}(S+T) + \frac{1}{2}(\alpha^2 + \gamma^2)$ $U + \frac{1}{2}(\beta^2 + \delta^2)V + \frac{1}{4}\alpha^2\gamma^2W + \frac{1}{2}(\alpha^2\delta^2 + \beta^2\gamma^2)X + \frac{1}{4}\beta^2\delta^2Y.$
			$AB.ab \times Ab.aB$	2	$X_{\alpha+1} = \frac{1}{2}T + \frac{1}{2}(\alpha\beta + \gamma\delta)(U+V) + \frac{1}{2}\alpha\beta\gamma\delta(W+2X+Y).$
			$Ab.aB \times Ab.aB$	1	$Y_{n+1} = 2(F + K) + \frac{1}{2}(\alpha^2 + \gamma^2)M + \frac{1}{2}(\beta^2 + \delta^2)P + \frac{1}{2}(R + T) + \frac{1}{2}(\beta^2 + \delta^2)U + \frac{1}{2}(\alpha^2 + \gamma^2)V + \frac{1}{2}\beta^2\delta^2W + \frac{1}{2}(\alpha^2\delta^2 + \beta^2\gamma^2)X + \frac{1}{4}\alpha^2\gamma^2Y.$

Result for sib-mating

Omitting some rather tedious algebra, the solution of these equations is:

$$\zeta = \frac{q}{2 - 3q}, \quad \theta = \frac{2q}{2 - 3q}, \quad \kappa = \frac{1}{2 - 3q},$$

$$\lambda = \frac{1 - 2q}{2 - 3q}, \quad \mu = \frac{1 - 2q}{2 - 3q}, \quad \nu = \frac{2q}{2 - 3q}$$

as may easily be verified.

$$c_{\infty} = c_{n} + 2e_{n} + \frac{1}{1 + 6x} [(1 - 2x)(d_{n} + 2f_{n} + 2j_{n} + \frac{1}{2}k_{n}) + 2g_{n} + 4x(h_{n} + i_{n})]$$
(3.4)

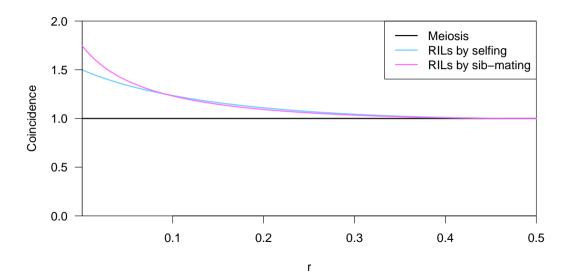
and $y = \frac{1}{2}(1 - c_{\infty})$.

In the case considered, $d_0 = 1$, $c_\infty = \zeta d_0 = 1 - 2x/1 + 6x$. Hence the proportion of crossover zygotes, y = 4x/1 + 6x (3.5).

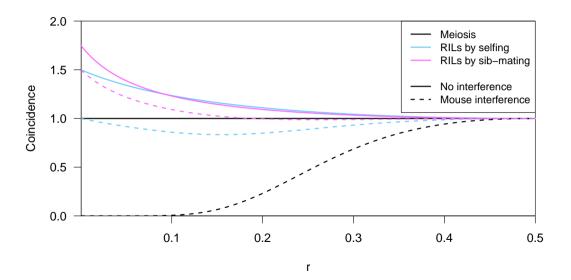
3-point coincidence

- r_{ij} = recombination fraction for interval (i, j) Assume $r_{12} = r_{23} = r$.
- Coincidence = c = Pr(double recombinant)/r² = Pr(rec'n in 23 | rec'n in 12)/Pr(rec'n in 23)
- No interference = 1 Positive interference < 1 Negative interference > 1
- Generally c is a function of r

Coincidence



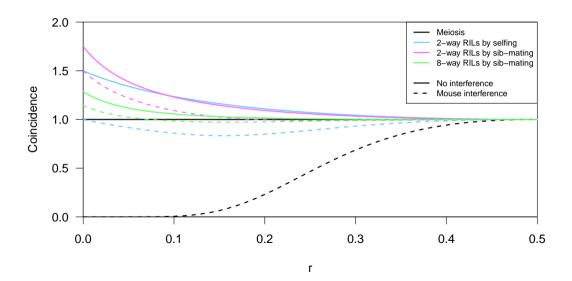
Coincidence



Coincidence in 8-way RILs

- ► The trick that allowed us to get the coincidence for 2-way RILs doesn't work for 8-way RILs.
- ► It's sufficient to consider 4-way RILs.
- ► Calculations for 3 points in 4-way RILs is still astoundingly complex.
 - 2 points in 2-way RILs by sib mating:
 55 parental types → 22 states by symmetry
 - 3 points in 4-way RILs by sib mating:
 2,164,240 parental types → 137,488 states by symmetry
- Even counting the states was difficult.

Coincidence

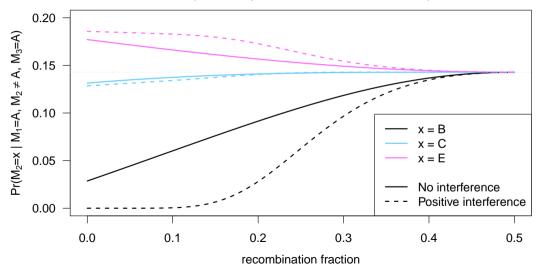


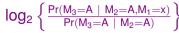
The formula

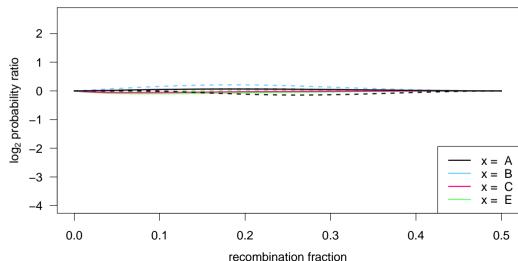
$$C = \frac{(1+6r)[280+1208r-848r^2+5c(7-28r-368r^2+344r^3)-2c^2(49-324r+452r^2)r^2-16c^3(1-2r)r^4]}{49(1+12r-12cr^2)[5+10r-4(2+c)r^2+8cr^3]}$$

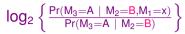
3-point symmetry

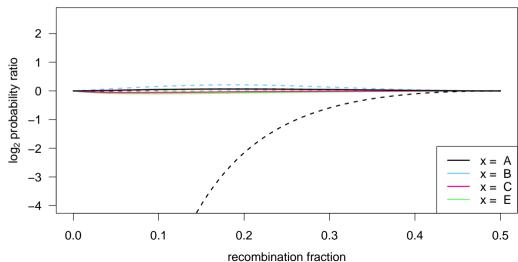
$$Pr(M_2 = x \mid M_1 = A, M_2 \neq A, M_3 = A)$$

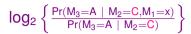


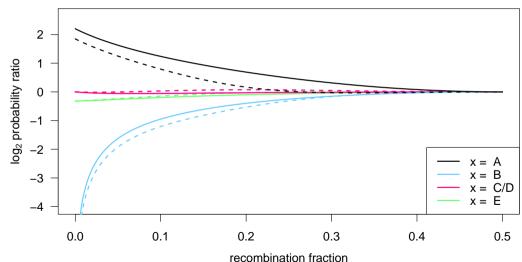




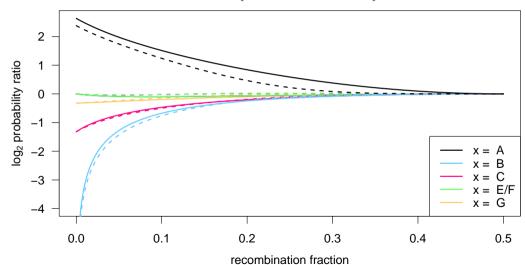








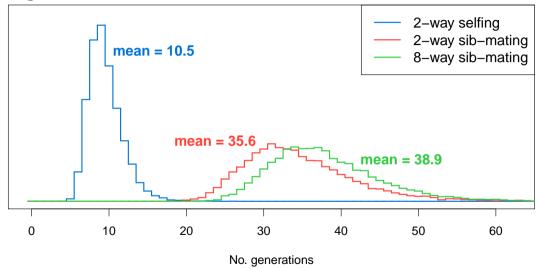
$$log_{2}\left\{ \frac{Pr(M_{3}=A\mid M_{2}=E,M_{1}=x)}{Pr(M_{3}=A\mid M_{2}=E)}\right\}$$



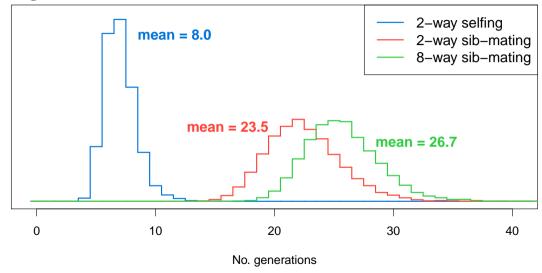
Whole genome simulations

- ► 2-way selfing, 2-way sib-mating, 8-way sib-mating
- ► Mouse-like genome, 1665 cM
- Strong positive crossover interference
- Inbreed to complex fixation
- ► 10,000 simulation replicates

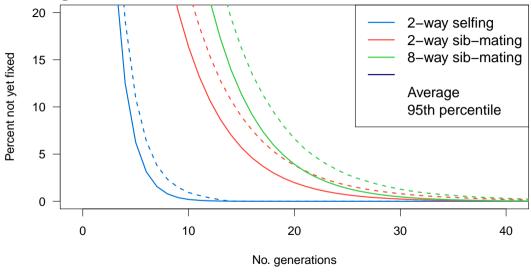
No. generations to fixation



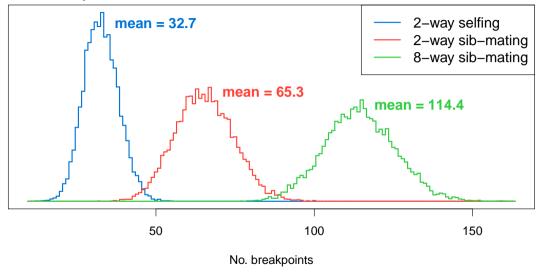
No. generations to 99% fixation



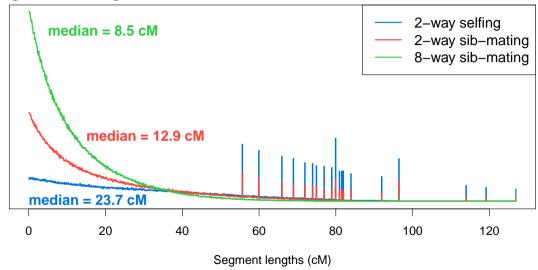
Percent genome not fixed



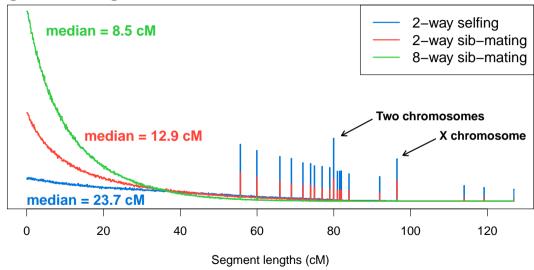
No. breakpoints



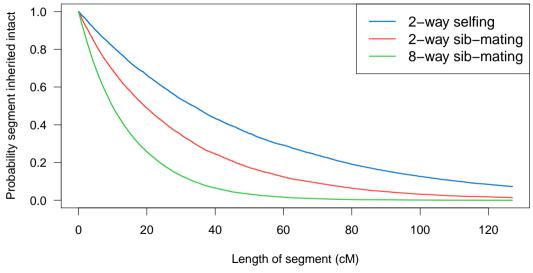
Segment lengths



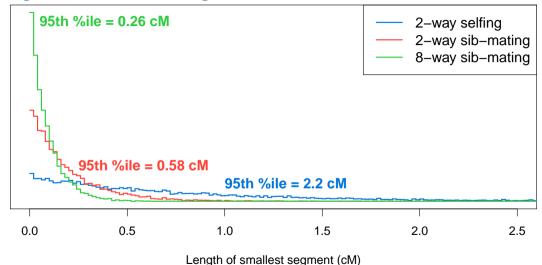
Segment lengths



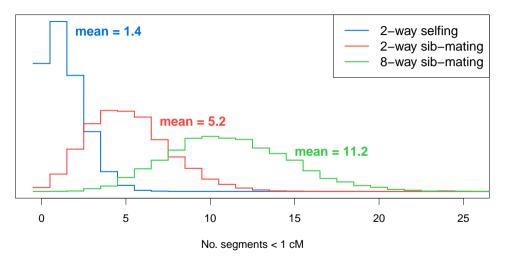
Probability a segment is inherited intact



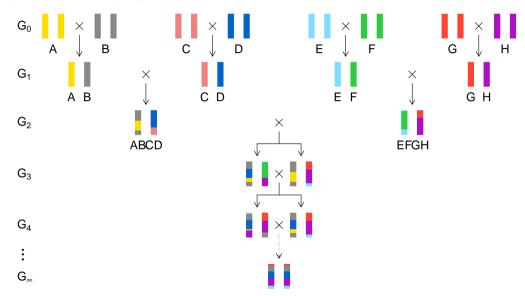
Length of smallest segment



No. segments < 1 cM



Collaborative Cross



The PreCC

What happens at G_2F_k ?

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Pr(g_1 = i) as a function of k
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 $Pr(g_1=i,g_2=j)$ as a function of k and the recombination fraction

Crazy table

Table 4 Two-locus haplotype probabilities at generation F_k in the formation of four-way RIL by sibling mating

Chr.	Individual	Prototype	No. states	Probability of each
A	Random	AA	4	$\frac{1}{4(1+6r)} - \left[\frac{6r^2 - 7r - 3rs}{4(1+6r)s}\right] \left(\frac{1 - 2r + s}{4}\right)^k + \left[\frac{6r^2 - 7r + 3rs}{4(1+6r)s}\right] \left(\frac{1 - 2r - s}{4}\right)^k$
		AB	4	$\frac{r}{2(1+6r)} + \left[\frac{10r^2 - r - rs}{4(1+6r)s}\right] \left(\frac{1 - 2r + s}{4}\right)^k - \left[\frac{10r^2 - r + rs}{4(1+6r)s}\right] \left(\frac{1 - 2r - s}{4}\right)^k$
		AC	8	$\frac{r}{2(1+6r)} - \left[\frac{2r^2 + 3r + rs}{4(1+6r)s}\right] \left(\frac{1 - 2r + s}{4}\right)^k + \left[\frac{2r^2 + 3r - rs}{4(1+6r)s}\right] \left(\frac{1 - 2r - s}{4}\right)^k$
×	Female	AA	2	$\frac{1}{3(1+4r)} + \frac{1}{6(1+r)} \left(-\frac{1}{2}\right)^k - \left[\frac{4r^3 - (4r^2 + 3r)t + 3r^2 - 5r}{4(4r^2 + 5r + 1)t}\right] \left(\frac{1-r+t}{4}\right)^k + \left[\frac{4r^3 + (4r^2 + 3r)t + 3r^2 - 5r}{4(4r^2 + 5r + 1)t}\right] \left(\frac{1-r+t}{4}\right)^k$
		AB	2	$\frac{2r}{3(1+4r)} + \frac{r}{3(1+r)} \left(-\frac{1}{2}\right)^k + \left[\frac{2r^3 + 6r^2 - (2r^2 + r)t}{2(4r^2 + 5r + 1)t}\right] \left(\frac{1-r+t}{4}\right)^k - \left[\frac{2r^3 + 6r^2 + (2r^2 + r)t}{2(4r^2 + 5r + 1)t}\right] \left(\frac{1-r-t}{4}\right)^k$
		AC	4	$\frac{2r}{3(1+4r)} - \frac{r}{6(1+r)} \left(-\frac{1}{2}\right)^k - \left[\frac{9r^2 + 5r + rt}{4(4r^2 + 5r + 1)t}\right] \left(\frac{1-r+t}{4}\right)^k + \left[\frac{9r^2 + 5r - rt}{4(4r^2 + 5r + 1)t}\right] \left(\frac{1-r-t}{4}\right)^k$
		СС	1	$\frac{1}{3(1+4r)} - \frac{1}{3(1+r)} \left(-\frac{1}{2}\right)^k + \left[\frac{9r^2 + 5r + rt}{2(4r^2 + 5r + 1)t}\right] \left(\frac{1-r+t}{4}\right)^k - \left[\frac{9r^2 + 5r - rt}{2(4r^2 + 5r + 1)t}\right] \left(\frac{1-r-t}{4}\right)^k$
х	Male	AA	2	$\frac{1}{3(1+4r)} - \frac{1}{3(1+r)} \left(-\frac{1}{2}\right)^k + \left[\frac{r^2 - (8r^3 + r^2 - 3r)t - 10r^2 + 5r}{2(4r^4 - 35r^3 - 29r^2 + 15r + 5)}\right] \left(\frac{1-r + t}{4}\right)^k + \left[\frac{r^3 + (8r^3 + r^2 - 3r)t - 10r^2 + 5r}{2(4r^4 - 35r^3 - 29r^2 + 15r + 5)}\right] \left(\frac{1-r - t}{4}\right)^k$
		AB	2	$\frac{2r}{3(1+4r)} - \frac{2r}{3(1+r)} \left(-\frac{1}{2}\right)^k + \left[\frac{r^4 + (5r^3 - r)t - 10r^3 + 5r^2}{4r^4 - 35r^3 - 29r^2 + 15r + 5}\right] \left(\frac{1 - r + t}{4}\right)^k + \left[\frac{r^4 - (5r^3 - r)t - 10r^3 + 5r^2}{4r^4 - 35r^3 - 29r^2 + 15r + 5}\right] \left(\frac{1 - r - t}{4}\right)^k$
		AC	4	$\frac{2r}{3(1+4r)} + \frac{r}{3(1+r)} \left(-\frac{1}{2}\right)^k - \left[\frac{2r^4 + (2r^3 - r^2 + r)t - 19r^3 + 5r}{2(4r^4 - 35r^3 - 29r^2 + 15r + 5)}\right] \left(\frac{1-r + t}{4}\right)^k - \left[\frac{2r^4 - (2r^3 - r^2 + r)t - 19r^3 + 5r}{2(4r^4 - 35r^3 - 29r^2 + 15r + 5)}\right] \left(\frac{1-r - t}{4}\right)^k$
		СС	1	$\frac{1}{3(1+4r)} + \frac{2}{3(1+r)} \left(-\frac{1}{2}\right)^k + \left[\frac{2r^4 + (2r^3 - r^2 + r)t - 19r^3 + 5r}{4r^4 - 35r^3 - 29r^2 + 15r + 5}\right] \left(\frac{1-r+t}{4}\right)^k + \left[\frac{2r^4 - (2r^3 - r^2 + r)t - 19r^3 + 5r}{4r^4 - 35r^3 - 29r^2 + 15r + 5}\right] \left(\frac{1-r-t}{4}\right)^k$

 $s = \sqrt{4r^2 - 12r + 5}$ and $t = \sqrt{r^2 - 10r + 5}$; the autosomal haplotype probabilities are valid for $r < \frac{1}{2}$.

Lesson

Computer simulations are hugely valuable.

Uses of simulations

- ► Study probabilities
- ► Estimate power/sample size
- Evaluate performance of a method
- ► Evaluate sensitivity/robustness of a method

Relative advantages?

- ► Simulations
- ► Numerical calculations
- Analytic calculations

References

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