

Supplementary Tables & Figures

Modelling sex-specific crossover patterning in Arabidopsis

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Table S1: Chromosome metrics and beam-film parameters.	2
Table S2. Beam-film parameters used for simulations shown in each figure.	3
Figure S1. Experimental and simulated crossover distributions.	6
Figure S2. CoC curves for simulated and experimental recombination data.	7
Figure S3. Examples of simulated data that did not fit experimental data.	8
Figure S4. Proportions of different double crossover (DCO) classes.	9

Table S1. Chromosome metrics and beam-film parameters

Chr	Sex	Mb	$\mu\text{m SC}$	Beam-film parameters															
				N [#]	B [#]	E [#]	Bs [#]	Be [#]	Bd [#]	Smax [^]	Bsmax [#]	A [#]	L _p [^]	L _{Mb} [*]	L _{sc} [*]	cL [^]	cR [^]	M [#]	T2prob [^]
1	M	30.4	49.2	64	1	0.6	0.475	0.5	0.01	8.5	1	1	0.65	19.8	32.0	0.8	1	1	0.005
2	M	19.7	31.9	41	1	0.6	0.175	0.225	0.01	7.5	1	1	0.85	16.7	27.1	0.3	0.9	1	0.0065
3	M	23.5	37.9	49	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	16.4	26.6	0.4	0.9	1	0.008
4	M	18.6	30.1	39	1	0.6	0.125	0.225	0.01	4	1	1	0.6	11.2	18.0	0.6	0.9	1	0.0055
5	M	27.0	43.6	56	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	21.6	34.9	1.1	0.9	1	0.0065
1	F	30.4	25.0	64	1	0.6	0.475	0.5	0.01	8	1	1	1	30.4	25.0	0.7	0.5	1	0.003
2	F	19.7	16.2	41	1	0.6	0.175	0.25	0.01	7	1	1	1.6	31.5	25.9	1.2	0.8	1	0.004
3	F	23.5	19.3	49	1	0.6	0.5	0.65	0.01	6	1	1	1	23.5	19.3	0.5	0.6	1	0.005
4	F	18.6	15.3	39	1	0.6	0.125	0.225	0.01	7	1	1	1.7	31.6	26.0	0.8	0.5	1	0.003
5	F	27.0	22.2	56	1	0.6	0.4	0.5	0.01	6.5	1	1	1	27.0	22.2	0.7	0.6	1	0.003
1	F _{DSB}	30.4	25.0	38	1	0.6	0.475	0.525	0.01	7	1	1	0.9	27.4	22.5	0.5	0.7	1	0.006
2	F _{DSB}	19.7	16.2	38	1	0.6	0.475	0.525	0.01	6.5	1	1	1.5	29.6	24.3	1	0.8	1	0.004
3	F _{DSB}	23.5	19.3	29	1	0.6	0.5	0.65	0.01	6	1	1	1	23.5	19.3	0.5	0.6	1	0.008
4	F _{DSB}	18.6	15.3	23	1	0.6	0.125	0.225	0.01	6	1	1	1.7	31.6	26.0	0.8	0.4	1	0.005
5	F _{DSB}	27.0	22.2	34	1	0.6	0.4	0.5	0.01	8.5	1	1	1.1	29.7	24.4	0.6	0.5	1	0.005

#Optimised parameter

^ Fixed parameter

* Calculated based on L_p

Table S2. Beam-film parameters used for simulations shown in each figure.

Figure	Part	Details	Beam-film parameters													
			N	B	E	Bs	Be	Bd	Smax	Bsmax	A	L	cL	cR	M	T2prob
1	A-D	male	56	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
1	A-D	female	56	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
3	A-C	L - 0.4	60	1	0.6	0.45	0.55	0.01	7	1	1	0.4	0.8	0.8	1	0.004
3	A-C	L - 0.7	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.004
3	A-C	L - 1	60	1	0.6	0.45	0.55	0.01	7	1	1	1	0.8	0.8	1	0.004
3	A-C	L - 1.3	60	1	0.6	0.45	0.55	0.01	7	1	1	1.3	0.8	0.8	1	0.004
3	D-F	Smax - 3	60	1	0.6	0.45	0.55	0.01	3	1	1	0.7	0.8	0.8	1	0.004
3	D-F	Smax - 5	60	1	0.6	0.45	0.55	0.01	5	1	1	0.7	0.8	0.8	1	0.004
3	D-F	Smax - 7	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.004
3	D-F	Smax - 9	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
3	G-I	T2prob - 0	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0
3	G-I	T2prob - 0.002	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.002
3	G-I	T2prob - 0.004	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
3	G-I	T2prob - 0.006	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.006
4	A-B	L - 0.8	60	1	0.6	0.45	0.55	0.01	7	1	1	0.8	0.8	0.8	1	0.004
4	A-B	L - 0.9	60	1	0.6	0.45	0.55	0.01	7	1	1	0.9	0.8	0.8	1	0.004
4	A-B	L - 1	60	1	0.6	0.45	0.55	0.01	7	1	1	1	0.8	0.8	1	0.004
4	A-B	L - 1.1	60	1	0.6	0.45	0.55	0.01	7	1	1	1.1	0.8	0.8	1	0.004
4	C-D	CI	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0
4	C-D	CII	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.004
4	C-D	CI & CII	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.004
5	A-C	wt	30	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
5	A-C	wt	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
5	A-C	wt	90	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
5	A-C	wt	120	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.004
5	D-F	mut	30	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.2
5	D-F	mut	60	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.2
5	D-F	mut	90	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.2
5	D-F	mut	120	1	0.6	0.45	0.55	0.01	9	1	1	0.7	0.8	0.8	1	0.2
5	G	wt male - chr 1 - 125 DSBs	32	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.005
5	G	wt male - chr 2 - 125 DSBs	21	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.0065
5	G	wt male - chr 3 - 125 DSBs	25	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.008
5	G	wt male - chr 4 - 125 DSBs	19	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.0055
5	G	wt male - chr 5 - 125 DSBs	28	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
5	G	wt male - chr 1 - 250 DSBs	64	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.005
5	G	wt male - chr 2 - 250 DSBs	42	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.0065
5	G	wt male - chr 3 - 250 DSBs	49	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.008
5	G	wt male - chr 4 - 250 DSBs	39	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.0055
5	G	wt male - chr 5 - 250 DSBs	56	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
5	G	wt male - chr 1 - 375 DSBs	96	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.005
5	G	wt male - chr 2 - 375 DSBs	63	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.0065
5	G	wt male - chr 3 - 375 DSBs	75	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.008
5	G	wt male - chr 4 - 375 DSBs	57	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.0055
5	G	wt male - chr 5 - 375 DSBs	84	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
5	G	wt male - chr 1 - 500 DSBs	128	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.005
5	G	wt male - chr 2 - 500 DSBs	84	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.0065
5	G	wt male - chr 3 - 500 DSBs	98	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.008
5	G	wt male - chr 4 - 500 DSBs	78	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.0055
5	G	wt male - chr 5 - 500 DSBs	112	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
5	G	mut male - chr 1 - 125 DSBs	32	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.2
5	G	mut male - chr 2 - 125 DSBs	21	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.2
5	G	mut male - chr 3 - 125 DSBs	25	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.2
5	G	mut male - chr 4 - 125 DSBs	19	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.2
5	G	mut male - chr 5 - 125 DSBs	28	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.2

5	G	mut male - chr 1 - 250 DSBs	64	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.2
5	G	mut male - chr 2 - 250 DSBs	42	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.2
5	G	mut male - chr 3 - 250 DSBs	49	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.2
5	G	mut male - chr 4 - 250 DSBs	39	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.2
5	G	mut male - chr 5 - 250 DSBs	56	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.2
5	G	mut male - chr 1 - 375 DSBs	96	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.2
5	G	mut male - chr 2 - 375 DSBs	63	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.2
5	G	mut male - chr 3 - 375 DSBs	75	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.2
5	G	mut male - chr 4 - 375 DSBs	57	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.2
5	G	mut male - chr 5 - 375 DSBs	84	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.2
5	G	mut male - chr 1 - 500 DSBs	128	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.2
5	G	mut male - chr 2 - 500 DSBs	84	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.2
5	G	mut male - chr 3 - 500 DSBs	98	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.2
5	G	mut male - chr 4 - 500 DSBs	78	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.2
5	G	mut male - chr 5 - 500 DSBs	112	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.2
5	G	wt female - chr 1 - 125 DSBs	32	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.003
5	G	DSBs	21	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.004
5	G	DSBs	25	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.005
5	G	DSBs	19	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.003
5	G	DSBs	28	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
5	G	DSBs	64	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.003
5	G	DSBs	42	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.004
5	G	DSBs	49	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.005
5	G	DSBs	39	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.003
5	G	DSBs	56	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
5	G	wt female - chr 1 - 375 DSBs	96	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.003
5	G	DSBs	63	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.004
5	G	DSBs	75	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.005
5	G	DSBs	57	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.003
5	G	DSBs	84	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
5	G	DSBs	57	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.003
5	G	DSBs	128	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.003
5	G	DSBs	84	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.004
5	G	DSBs	98	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.005
5	G	DSBs	78	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.003
5	G	DSBs	112	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
5	G	mut female - chr 1 - 125 DSBs	32	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.2
5	G	DSBs	21	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.2
5	G	DSBs	25	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.2
5	G	DSBs	19	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.2
5	G	DSBs	28	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.2
5	G	DSBs	64	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.2
5	G	DSBs	42	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.2
5	G	DSBs	49	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.2

		mut female - chr 4 - 250														
5	G	DSBs	39	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.2
5	G	mut female - chr 5 - 250	56	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.2
5	G	DSBs	96	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.2
5	G	mut female - chr 1 - 375	63	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.2
5	G	DSBs	75	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.2
5	G	mut female - chr 2 - 375	57	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.2
5	G	DSBs	84	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.2
5	G	mut female - chr 3 - 375	128	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.2
5	G	DSBs	84	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.2
5	G	mut female - chr 1 - 500	98	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.2
5	G	DSBs	78	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.2
5	G	mut female - chr 2 - 500	112	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.2
S1 & S2	chr1	male	64	1	0.6	0.48	0.5	0.01	8.5	1	1	0.65	0.8	1	1	0.005
S1 & S2	chr1	female	64	1	0.6	0.48	0.5	0.01	8	1	1	1	0.7	0.5	1	0.003
S1 & S2	chr2	male	41	1	0.6	0.18	0.23	0.01	7.5	1	1	0.85	0.3	0.9	1	0.0065
S1 & S2	chr2	female	41	1	0.6	0.18	0.25	0.01	7	1	1	1.6	1.2	0.8	1	0.004
S1 & S2	chr3	male	49	1	0.6	0.5	0.65	0.01	5.5	1	1	0.7	0.4	0.9	1	0.008
S1 & S2	chr3	female	49	1	0.6	0.5	0.65	0.01	6	1	1	1	0.5	0.6	1	0.005
S1 & S2	chr4	male	39	1	0.6	0.13	0.23	0.01	4	1	1	0.6	0.6	0.9	1	0.0055
S1 & S2	chr4	female	39	1	0.6	0.13	0.23	0.01	7	1	1	1.7	0.8	0.5	1	0.003
S1 & S2	chr5	male	56	1	0.6	0.4	0.5	0.01	9.5	1	1	0.8	1.1	0.9	1	0.0065
S1 & S2	chr5	female	56	1	0.6	0.4	0.5	0.01	6.5	1	1	1	0.7	0.6	1	0.003
S3	chr5	low	56	1	0.6	0.4	0.5	0.01	3	1	1	1.7	0.1	0.1	1	0.002
S3	chr5	high	56	1	0.6	0.4	0.5	0.01	10	1	1	0.4	1.3	1.3	1	0.006
S4		CI - all	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0
S4		CII - T2Prob 0.001	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.001
S4		CI & CII - T2Prob 0.001	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.001
S4		CII - T2Prob 0.005	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.005
S4		CI & CII - T2Prob 0.005	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.005
S4		CII - T2Prob 0.01	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.01
S4		CI & CII - T2Prob 0.01	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.01
S4		CII - T2Prob 0.025	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.025
S4		CI & CII - T2Prob 0.025	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.025
S4		CII - T2Prob 0.125	60	1	0.6	0.45	0.55	0.01	0	1	1	0.7	0.8	0.8	1	0.125
S4		CI & CII - T2Prob 0.125	60	1	0.6	0.45	0.55	0.01	7	1	1	0.7	0.8	0.8	1	0.125

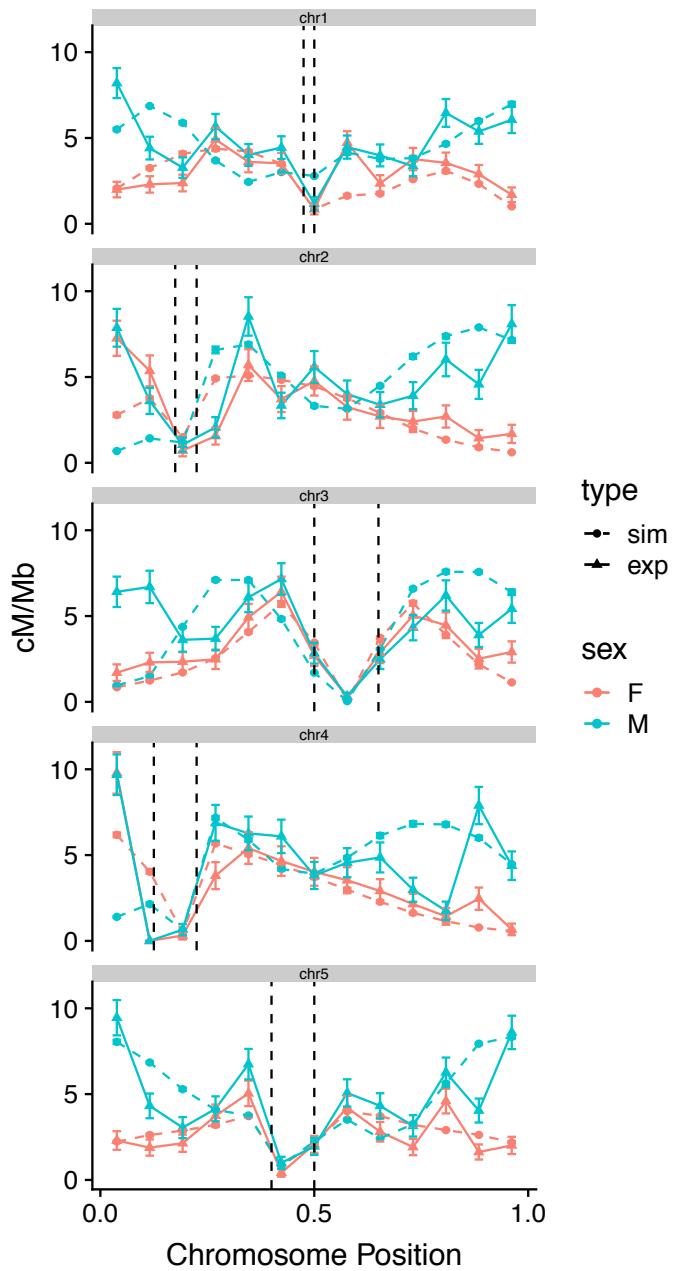


Figure S1. Experimental and simulated crossover distributions. Each analysis includes experimental (solid lines) and simulated (dashed lines) data for male (blue) and female (orange). Dashed lines represent the limits of the centromeric region over which precursor (DSB) number is markedly reduced both biologically (38) and during simulations. Male and female simulations shown, assume 250 DSBs genome-wide. Chromosomes were divided into 13 equal-sized adjacent intervals for analysis.

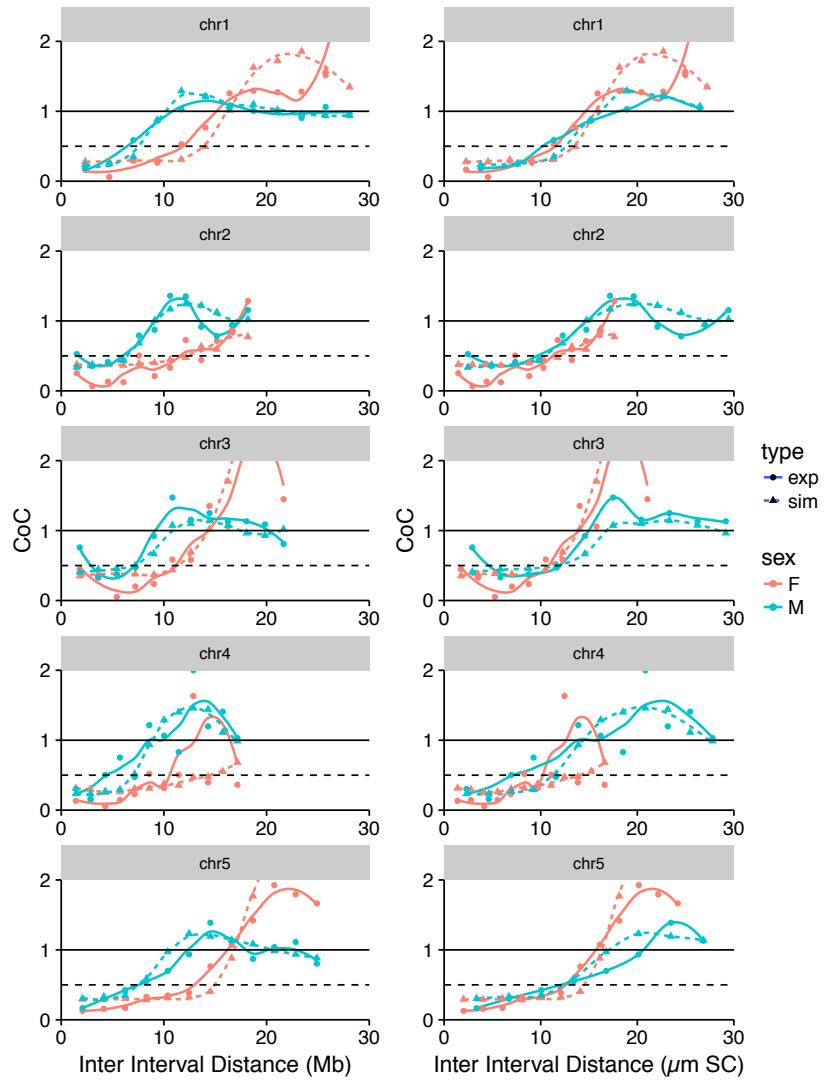


Figure S2. CoC curves for simulated and experimental recombination data. Each analysis includes experimental (solid lines) and simulated (dashed lines) data for male (blue) and female (orange). CoC curves with inter-interval distance measured in either Mb or $\mu\text{m SC}$ are shown. Male curves are shifted to the right relative to female curves when inter-interval distance is measured in Mb, but are similar when inter-interval distance is measured in $\mu\text{m SC}$. Male and female simulations shown assume 250 DSBs genome-wide. Chromosomes were divided into 13 equal-sized adjacent intervals for analysis

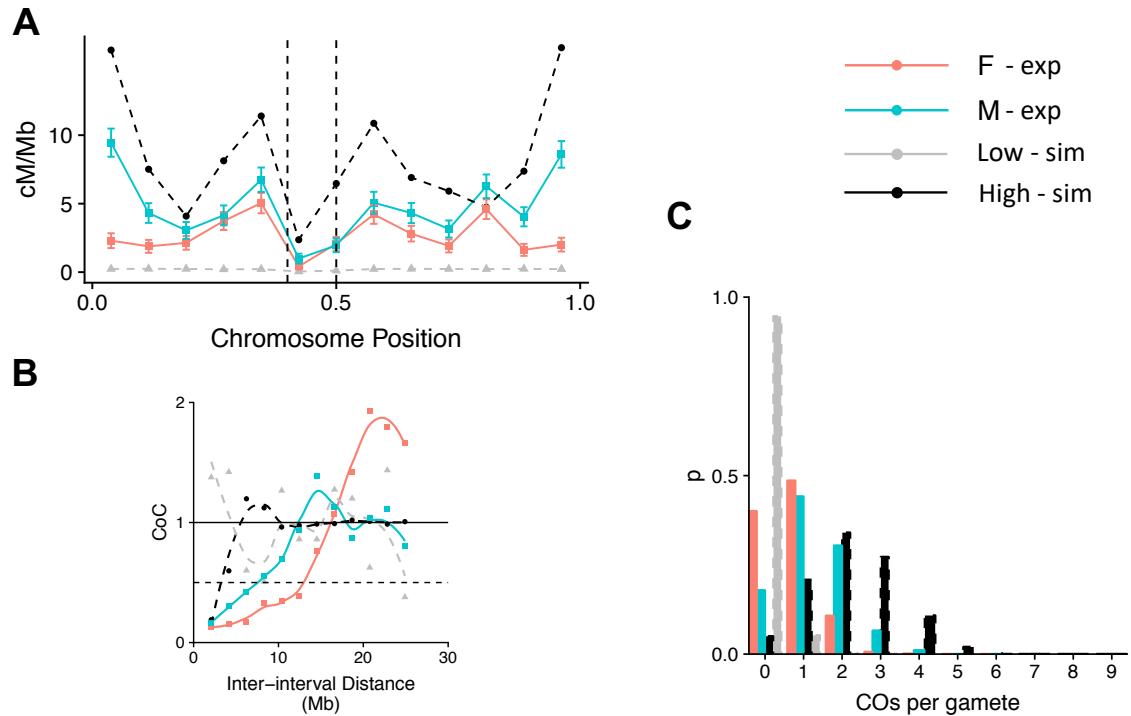


Figure S3. Examples of simulated data that did not fit experimental data. Each analysis shows experimental (solid lines) for male (blue) and female (orange) and simulated (dashed lines) data for high (black) and low (grey) recombining parameter sets. **A** Crossover distributions for *Arabidopsis* chromosome 5. Dashed lines represent the limits of the centromeric region over which precursor (DSB) number is markedly reduced both biologically (38) and during simulations. Error bars indicate 95% confidence intervals. **B** CoC curves for chromosome 5 with inter-interval distance (IID, the distance between a pair of genetic intervals) measured in Mb. **C** Event distribution for chromosome 5. Simulations shown assume 250 DSBs genome-wide. Chromosomes were divided into 13 equal-sized adjacent intervals for analysis.

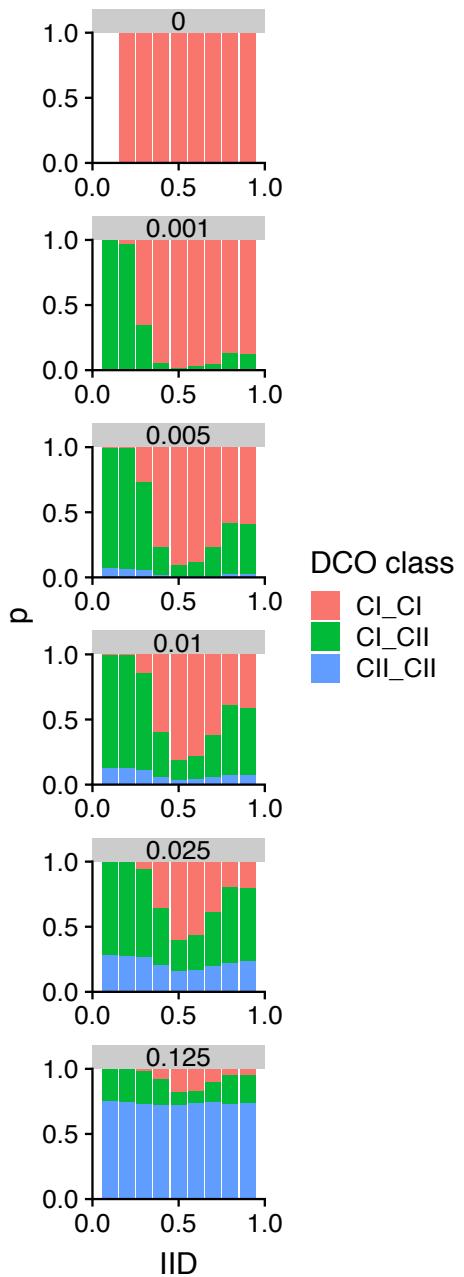


Figure S4. Proportions of different double crossover (DCO) classes. Charts show the proportions of DCOs formed between two class I crossovers (CI_CI), two class II crossovers (CII_CII), or a class I and a class II CO (CI_CII) for different IIDs and different values of T2Prob (grey bars). Total proportion of class II crossovers are as follows: T2Prob = 0, 0% class II COs; T2Prob = 0.001, 3% class II COs; T2Prob = 0.005, 13% class II COs; T2Prob = 0.01, 23% class II COs; T2Prob = 0.025, 43% class II COs; T2Prob = 0.125, 81% class II COs.