

785 **Supp. Table 1** - Sex differences in recombination landscapes summarized across taxa.

Species	Sex chromosomes	Greater overall recombination	Greater recombination at telomeres	Greater recombination at centromeres / chromosome centers (*)	Notes	Sources
Eutherian Mammals						
<i>Bos taurus</i> (cattle)	XY	M	M	N.S.		Ma et al. (2015), Wang et al. (2016b)
<i>Canis familiaris</i> (domestic dog)	XY	F	M	F		Wong et al. (2010), Campbell et al. (2016)
<i>Cervus elaphus</i> (red deer)	XY	F	M	F	Male recombination marginally elevated near telomeres.	Johnston et al. (2017)
<i>Homo sapiens</i> (human)	XY	F	M	F		Broman et al. (1998), Kong et al. (2002)
<i>Mus musculus</i> (house mouse)	XY	F	M	F	No recombination near centromeres in males. Elevated near centromeres in females.	Shifman et al. (2006), Paigen et al. (2008), Cox et al. (2009)
<i>Ovis aries</i> (Soay sheep)	XY	M	M	N.S.	Elevated near telomeres in males unless near centromere (i.e., acrocentric chromosomes). Reduced near telomere in females.	Johnston et al. (2016)
<i>Pan troglodytes</i> (chimpanzee)	XY	F	M	F	Elevated near telomeres in males. Relatively uniform in females.	Venn et al. (2014)
<i>Sus scrofa</i> (domestic pig)	XY	F	F	N.S.	Elevated near telomeres regardless of centromere position in both sexes, somewhat more strongly in females.	Tortereau et al. (2012)

Marsupials						
<i>Monodelphis domestica</i> (gray short-tailed opossum)	XY	M	F	M*	Chiasmata cluster near telomeres in females. Majority are interstitial in males.	Hayman et al. (1988)
<i>Sminthopsis crassicaudata</i> (fat-tailed dunnart)	XY	M	F	M*	Chiasmata cluster near telomeres in females. Majority are interstitial in males.	Bennett et al. (1986)
Birds						
<i>Anser anser</i> (gray goose)	ZW	F	M	N.S.	Subtelomeric peaks more pronounced in males.	Torgasheva and Borodin (2017)
<i>Columba livia</i> (rock dove)	ZW	N.S.	N.S.	N.S.	Distribution of recombination nodules similar between sexes.	Pigozzi and Solari (1999)
<i>Coturnix japonica</i> (Japanese quail)	ZW	N.S.	N.S.	N.S.	COs evenly distributed in both sexes.	Calderon and Pigozzi (2006)
<i>Ficedula albicollis</i> (collared flycatcher)	ZW	M	M	N.S.	Elevated near telomeres in males. Very low at extreme terminal ends in females.	Smeds et al. (2016)
<i>Parus major</i> (great tit)	ZW	N.S.	N.S.	M*	Varies by chromosome. Most commonly male-biased in chromosome centers.	Van Oers et al. (2014)
<i>Taeniopygia guttata</i> (zebra finch)	ZW	N.S.	N.S.	N.S.	Elevated near telomeres in males and females.	Backström et al. (2010)
Reptiles						
<i>Crocodylus porosus</i> (saltwater crocodile)	ESD	F	M?	F	Elevated near telomeres in males. Relatively uniform in females. Did not provide direct comparison of male and female recombination rate at telomeres.	Miles et al. (2009)
Amphibians						
<i>Hyla arborea</i> (European tree frog)	XY	F	M	F*	Limited to telomeres in males. Relatively uniform in females.	Brelsford et al. (2016a)
<i>Rana temporaria</i> (common frog)	XY	F	M	F*	Limited to telomeres in males. Relatively uniform in females.	Brelsford et al. (2016b)

<i>Xenopus borealis</i> (Marsabit clawed frog)	ZW	F	M	F*	Elevated at ends in males. Elevated in chromosome centers in females.	Furman and Evans (2018)
<i>Xenopus laevis</i> (African clawed frog)	ZW	F	M	F*	Elevated at ends in males. Elevated in chromosome centers in females.	Furman and Evans (2018)
<i>Triturus cristatus</i> (northern crested newt)	XY / ESD	M	M	F*	Chiasmata relatively uniform in males. Localized near center of chromosomes in females.	Watson and Callan (1963)
<i>Triturus helveticus</i> (palmate newt)	XY / ESD	?	M	F*	Chiasmata localized near telomeres in males. Relatively uniform in females.	Watson and Callan (1963)
Teleost Fish						
<i>Danio rerio</i> (zebrafish)	Polygenic	F	M	F		Singer et al. (2002)
<i>Gasterosteus</i> spp. (stickleback)	XY	F	M	F	Decreases with distance from telomere and increases with distance from centromeres in males. More uniform in females.	Sardell et al. (2018)
<i>Notothenia coriiceps</i> (Bullhead notothen)	?	N.S.	M	F	Elevated near telomeres in males. Elevated near centromeres in females.	Amores et al. (2017)
<i>Hippoglossus hippoglossus</i> (Atlantic halibut)	XY	F	M	F*	Male-biased in telomeric half, female-biased in centromeric half.	Reid et al. (2007)
<i>Oncorhynchus mykiss</i> (rainbow trout)	XY	F	M	F*	Male-biased near telomeres, female-biased in middle of chromosome.	Sakamoto et al. (2000)
<i>Oreochromis</i> spp. (tilapia)	Polygenic / ESD	N.S.	M	F*	Male-biased near telomeres, female-biased in middle of chromosome.	Lee et al. (2005)
<i>Oreochromis niloticus</i> (Nile tilapia)	Polygenic / ESD	F	N.S.	N.S.	Suppressed near telomeres and elevated in middles in both sexes.	Conte et al. (2018)
<i>Paralichthys olivaceus</i> (Japanese flounder)	XY / ESD	M	M	F	Elevated near telomeres in males. Elevated near centromeres in females.	Castaño-Sánchez et al. (2010)

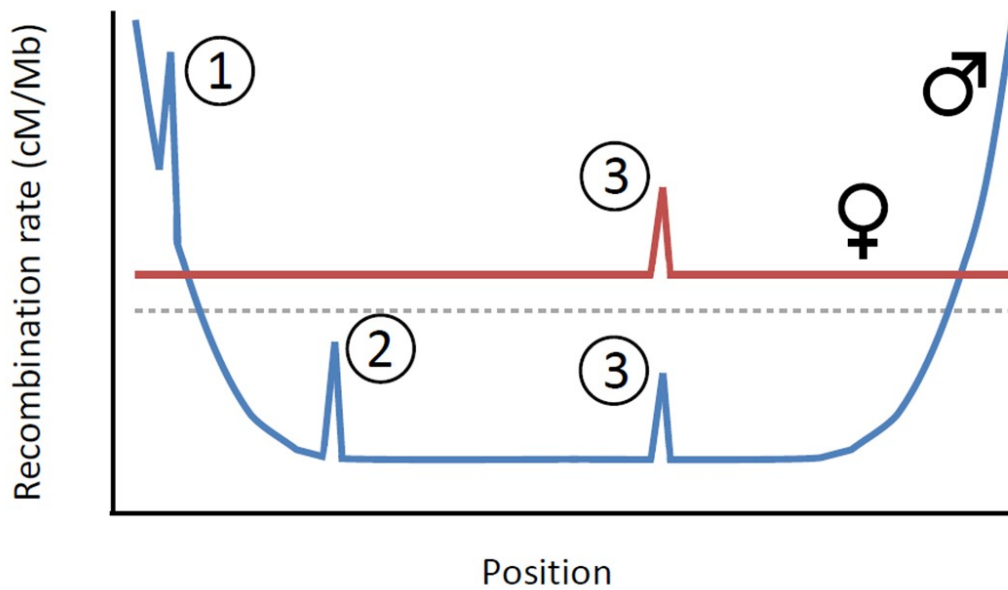
<i>Poecilia reticulata</i> (Trinidadian guppy)	XY	F	M	F	Limited to telomeres in males. Relatively uniform in females.	Bergero et al. (2019)
<i>Salmo salar</i> (Atlantic salmon)	XY	F	M	F	Elevated near telomeres and reduced near centromeres in males. More uniform in females.	Moen et al. (2008), Lien et al. (2011)
<i>Salmo trutta</i> (brown trout)	XY	F	M	F*	Elevated near telomeres in males but not females.	Gharbi et al. (2006)
<i>Salvelinus alpinus</i> (Arctic char)	XY	F	M	F*	Elevated near telomeres and reduced near putative centromeres in males.	Woram et al. (2004)
<i>Salvelinus fontinalis</i> (Brook char)	XY	F	M	F*	Elevated near telomeres in males. Elevated toward center of chromosomes in females.	Sutherland et al. (2017)
<i>Sparus aurata</i> (gilthead sea bream)	Hermaphrodite	F	Variable	Variable	Variable: some chromosomes male-biased near telomeres, others female-biased in telomeres.	Franch et al. (2006)
Invertebrates						
<i>Crassostrea</i> spp. (oysters)	Hermaphrodite	F	M	F	Elevated near telomeres in males. Elevated near centromeres in females.	Wang et al. (2016a)
<i>Pinctada maxima</i> (silver-lipped pearl oyster)	Hermaphrodite	F	M	F		Jones et al. (2013)
<i>Chorthippus jucundus</i> (large green grasshopper)	XO	N.S.	M*	M	Higher in males across whole chromosome, but biased more towards telomeres and away from centromeres in females relative to males.	Cano and Santos (1990)
<i>Chorthippus parallelus</i> (meadow grasshopper)	XO	M	M*	M	Higher in males across whole chromosome, but biased more towards telomeres and away from centromeres in females relative to males.	Cano and Santos (1990)

<i>Chorthippus vagans</i> (steppe grasshopper)	XO	M	M*	M	Higher in males across whole chromosome, but biased more towards telomeres in females relative to males.	Cano and Santos (1990)
<i>Euchorthippus chopardi</i> (Iberian straw grasshopper)	XO	M	M*	M	Higher in males across whole chromosome, but biased much more towards telomeres and away from centromeres in females.	Cano and Santos (1990)
<i>Euchorthippus pulvinatus</i> (eastern straw grasshopper)	XO	M	M*	M	Higher in males across whole chromosome, but biased more towards telomeres in females.	Cano and Santos (1990)
<i>Omocestus panteli</i> (Pantel's grasshopper)	XO	M	M*	M	Higher in males across whole chromosome, but biased more towards telomeres in females.	Cano and Santos (1990)
<i>Litopenaeus vannamei</i> (Pacific white shrimp)	ZW	N.S.	N.S.	N.S.	No significant difference between sexes.	Jones et al. (2017)
Plants						
<i>Arabidopsis thaliana</i> (thale cress)	Monoecious	M	M	N.S.	Elevated near telomeres in males. No difference in middles of chromosomes.	Giraut et al. (2011)
<i>Brassica nigra</i> (black mustard)	Monoecious	M	M	F	Elevated near telomeres in males. Elevated near centromeres in females.	Lagercrantz and Lydiate (1995)
<i>Hordeum bulbosum</i> (barley)	Monoecious	M	M	N.S.	Elevated near telomeres in males.	Devaux et al. (1995), Phillips et al. (2015)
<i>Solanum</i> spp. (tomato)	Monoecious	F	F	F	Reduced chromosome-wide in males.	de Vicente and Tanksley (1991)
<i>Zea mays</i> (maize)	Monoecious	N.S.	N.S.	N.S.	Fine-scale, but no broad-scale sex differences in recombination.	Kianian et al. (2018)

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787 Table Notes - “M” = males, “F” = females, “N.S.” = no significant sex difference, “ESD” = environmental sex determination

788 **Supp. Figure 1** – The standard definition for a recombination hotspot fails to identify some sex-
789 specific recombination peaks, and confounds effects that are sex specific with those that are not.
790 Feature (1) is identified as male-specific hotspot because its peak is above the chromosome
791 average (dashed gray line). Feature (2) is not because its peak lies below the average. Feature
792 (3), which has identical effects in males and females, is only recognized as a hotspot in females.



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795 **Supp. Material:** A model of sexually antagonistic *cis*-acting epistasis

796 We developed a simple model for the evolution of sex differences in recombination resulting from
797 sexually antagonistic *cis* epistasis (SACE). As explained in the text, the model is motivated by Equation
798 (32) of Lenormand (2003).

799 The model is of three loci. Loci *R* and *C* experience frequency-dependent selection which
800 maintains them polymorphic at a specified allele frequency. In the results shown here, we simply assumed
801 both loci have allele frequencies fixed at 0.5. In males, the loci interact epistatically such that haplotypes
802 R_0-C_0 and R_1-C_1 have fitness $(1 + \epsilon)$ relative to the other two haplotypes. We assume fitnesses are
803 multiplicative across chromosomes, so the fitness of a diploid is the product of the fitnesses of its
804 haplotypes. Epistatic selection is absent from females.

805 The third locus, *M*, is a neutral recombination modifier. Initially the ancestral allele M_0 is fixed,
806 and recombination rates are equal in males and females. The linkage map is *R-C-M*, and the
807 recombination rates between adjacent loci are r_{RC} and r_{CM} . We study the fate of a rare mutation M_1 that
808 modifies the recombination rates. In males, the modifier multiplies both recombination rates by a factor K
809 (where $K < 1$ corresponds to decreased recombination). In females, recombination is increased by the
810 same amount, and so the sex-averaged recombination rate is unchanged.

811 We introduce the mutation at a very low frequency (10^{-6}) and simulate the dynamic equations to
812 determine how the mutation's frequency changes. After a few generations of initial transients, the
813 frequency increases or decreases by a constant proportion that represents the system's dominant
814 eigenvalue. This eigenvalue can be interpreted as the modifier's "effective selection coefficient", s_{eff} ,
815 which is the selection coefficient acting on a nonneutral locus that would produce the same rate of
816 change.

817 The model described here does not strictly speaking involve sexually antagonistic epistasis, but
818 rather sex-limited epistasis. We show that case here for the sake of simplicity. Because we are assuming

819 no epistasis in females, the effect of the modifier on recombination in females has no impact on the
820 outcome. Qualitatively similar outcomes do result with sexually antagonistic epistasis (that is, when
821 epistasis in females opposes that in males) so long as epistatic selection is stronger in males than females.
822 In that case, a modifier that decreases recombination in males and increases it in females will spread, as
823 predicted by Lenormand (2003).

824 Results from several representative simulations are shown below. The simulation code is
825 available from the authors on request.

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828	ϵ	r_{RC}	r_{CM}	K	s_{eff}
829	0	0.005	0.1	1.00	0
830	0.01	0.005	0.1	0.5	2.6×10^{-7}
831	0.01	0.005	0.1	0.25	4.1×10^{-7}
832	0.05	0.005	0.1	0.25	9.6×10^{-6}
833	0.05	0.005	0.2	0.25	7.6×10^{-6}

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