S1 Appendix

Recursion equations

In each generation we census the genotype frequencies in male and female gametes/gametophytes (hereafter, gametes) between meiosis (and any meiotic drive) and gametic competition. At this stage we denote the frequencies of X- and Y-bearing gametes from males and females x_i° and y_i° . The superscript $\circ \in \{\mathcal{S}, \mathcal{Q}\}$ specifies the sex of the diploid that the gamete came from. The subscript $i \in \{1, 2, 3, 4\}$ specifies the genotype at the selected locus **A** and at the novel sex-determining locus **M**, where 1 = AM, 2 = aM, 3 = Am, and 4 = am. The gamete frequencies from each sex sum to one, $\sum_i x_i^{\circ} + y_i^{\circ} = 1$.

Competition then occurs among gametes of the same sex (e.g., among eggs and among sperm separately) according to the genotype at the A locus $(w_1^\circ = w_3^\circ = w_A^\circ, w_2^\circ = w_4^\circ = w_a^\circ$, see Table 1). The genotype frequencies after gametic competition are $x_i^{\circ,s} = w_i^\circ x_i^\circ / \bar{w}_H^\circ$ and $y_i^{\circ,s} = w_i^\circ y_i^\circ / \bar{w}_H^\circ$, where $\bar{w}_H^\circ = \sum_i w_i^\circ x_i^\circ + w_i^\circ y_i^\circ$ is the mean fitness of male $(\circ = \varnothing)$ or female $(\circ = \varnothing)$ gametes.

Random mating then occurs between gametes to produce diploid zygotes. The frequencies of XX zygotes are then denoted as xx_{ij} , XY zygotes as xy_{ij} , and YY zygotes as yy_{ij} , where **A** and **M** locus genotypes are given by $i, j \in \{1, 2, 3, 4\}$, as above. In XY zygotes, the haplotype inherited from an X-bearing gamete is given by i and the haplotype from a Y-bearing gamete is given by j. In XX and YY zygotes, individuals with diploid genotype ij are equivalent to those with diploid genotype ji; for simplicity, we use xx_{ij} and yy_{ij} with $i \neq j$ to denote the average of these frequencies, $xx_{ij} = (x_i^{Q,s} x_j^{d,s} + x_j^{Q,s} x_i^{d,s})/2$ and $yy_{ij} = (y_i^{Q,s} y_j^{d,s} + y_j^{Q,s} y_i^{d,s})/2$. Denoting the **M** locus genotype by $b \in \{MM, Mm, mm\}$ and the **X** locus genotype by

Denoting the **M** locus genotype by $b \in \{MM, Mm, mm\}$ and the **X** locus genotype by $c \in \{XX, XY, YY\}$, zygotes develop as females with probability k_{bc} . Therefore, the frequencies of XX females are given by $xx_{ij}^{Q} = k_{bc}xx_{ij}$, XY females are given by $xy_{ij}^{Q} = k_{bc}xy_{ij}$, and YY females are given by $yy_{ij}^{Q} = k_{bc}yy_{ij}$. Similarly, XX male frequencies are $xx_{ij}^{d} = (1 - k_{bc})xx_{ij}$, XY male frequencies are $xy_{ij}^{d} = (1 - k_{bc})xy_{ij}$, and YY males frequencies are $yy_{ij}^{d} = (1 - k_{bc})yy_{ij}$. This notation allows both the ancestral and novel sex-determining regions to determine zygotic sex according to an XY system, a ZW system, or an environmental sex-determining system. In addition, we can consider any epistatic dominance relationship between the two sex-determining loci. Here, we assume that the ancestral sex-determining system (**X** locus) is XY ($k_{MMXX} = 1$ and $k_{MMXY} = k_{MMYY} = 0$) or ZW ($k_{MMZZ} = 0$ and $k_{MMZW} = k_{MMWW} = 1$) and epistatically recessive to a dominant novel sex-determining locus, **M** ($k_{Mmc} = k_{mmc} = k$).

Selection among diploids then occurs according to the diploid genotype at the **A** locus, $l \in \{AA, Aa, aa\}$, for an individual of type ij (see Table 1). The diploid frequencies after selection in sex \circ are given by $xx_{ij}^{\circ,s} = w_l^{\circ}xx_{ij}/\bar{w}_D^{\circ}$, $xy_{ij}^{\circ,s} = w_l^{\circ}xy_{ij}/\bar{w}_D^{\circ}$, and $yy_{ij}^{\circ,s} = w_l^{\circ}yy_{ij}/\bar{w}_D^{\circ}$, where $\bar{w}_D^{\circ} = \sum_{i=1}^4 \sum_{j=1}^4 w_l^{\circ}xx_{ij} + w_l^{\circ}xy_{ij} + w_l^{\circ}yy_{ij}$ is the mean fitness of diploids of sex \circ .

Finally, these diploids undergo meiosis to produce the next generation of gametes. Recombination and sex-specific meiotic drive occur during meiosis. Here, we allow any relative locations for the \mathbf{X} , \mathbf{A} , and \mathbf{M} loci by using three parameters to describe the recombination rates between them. \mathbf{R} is the recombination rate between the \mathbf{A} and \mathbf{M} loci, ρ is the recombination rate between the \mathbf{M} and \mathbf{X} loci, and \mathbf{r} is the recombination rate between the \mathbf{A} and \mathbf{X} loci (Fig 1). S1 Table shows replacements that can be made for each possible ordering of the loci assuming that there is no cross-over interference. During meiosis in sex \circ , meiotic drive occurs such that,

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in Aa heterozygotes, a fraction α° of gametes produced carry the A allele and $(1 - \alpha^{\circ})$ carry the a allele.

Among gametes from sex o, the frequencies of haplotypes (before gametic competition) in the next generation are given by

$$x_{1}^{o'} = xx_{11}^{o,s} + xx_{13}^{o,s}/2 + (xx_{12}^{o,s} + xx_{14}^{o,s})a^{\circ} - R(xx_{13}^{o,s} - xx_{23}^{o,s})a^{\circ} + (xy_{11}^{o,s} + xy_{13}^{o,s})/2 + (xy_{12}^{o,s} + xy_{13}^{o,s})/2 + (xy_{12}^{o,s} + xy_{13}^{o,s})/2 + (xy_{12}^{o,s} - xy_{21}^{o,s})a^{\circ} - \rho(xy_{13}^{o,s} - xy_{31}^{o,s})/2 + [-(R+r+\rho)xy_{14}^{o,s} + (R+\rho-r)xy_{41}^{o,s} + (R+r-\rho)xy_{23}^{o,s} + (R+\rho-r)xy_{32}^{o,s}]a^{\circ}/2 + (R+r-\rho)xy_{23}^{o,s} + (R+\rho-r)xy_{32}^{o,s})a^{\circ} - R(xx_{23}^{o,s} - xx_{14}^{o,s})a^{\circ} - R(xx_{23}^{o,s} - xx_{14}^{o,s})a^{\circ} - \rho(xy_{24}^{o,s} + xy_{23}^{o,s})(1-a^{\circ}) - r(xy_{21}^{o,s} - xy_{12}^{o,s})/2 + (xy_{21}^{o,s} + xy_{23}^{o,s})(1-a^{\circ}) - r(xy_{21}^{o,s} - xy_{12}^{o,s})/2 + (R+\rho-r)xy_{23}^{o,s} + (R+\rho-\rho)xy_{23}^{o,s} + xy_{33}^{o,s})/2 + (R+\rho-\rho)xy_{23}^{o,s} + (R+\rho-\rho)xy$$

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$$y_{2}^{o'} = yy_{22}^{\circ,s} + yy_{24}^{\circ,s}/2 + (yy_{12}^{\circ,s} + yy_{23}^{\circ,s})\alpha^{\circ} - R(yy_{23}^{\circ,s} - yy_{14}^{\circ,s})/2 + (xy_{12}^{\circ,s} + xy_{32}^{\circ,s})(1 - \alpha^{\circ}) - r(xy_{12}^{\circ,s} - xy_{21}^{\circ,s})(1 - \alpha^{\circ}) - \rho(xy_{42}^{\circ,s} - xy_{24}^{\circ,s})/2 + \left[-(R + r + \rho)xy_{32}^{\circ,s} + (R + \rho - r)xy_{23}^{\circ,s} + (R + r - \rho)xy_{41}^{\circ,s} + (R + \rho - r)xy_{14}^{\circ,s} \right] (1 - \alpha^{\circ})/2$$

$$y_{3}^{o'} = yy_{33}^{\circ,s} + yy_{13}^{\circ,s}/2 + (yy_{23}^{\circ,s} + yy_{34}^{\circ,s})\alpha^{\circ} - R(yy_{23}^{\circ,s} - yy_{14}^{\circ,s})\alpha^{\circ} - R(yy_{23}^{\circ,s} - yy_{14}^{\circ,s})\alpha^{\circ} - \rho(xy_{13}^{\circ,s} - xy_{34}^{\circ,s})/2 + (xy_{23}^{\circ,s} + xy_{33}^{\circ,s})/2 + (r + r - \rho)xy_{23}^{\circ,s} + (R + \rho - r)xy_{32}^{\circ,s} + (R + r - \rho)xy_{14}^{\circ,s} + (R + \rho - r)xy_{32}^{\circ,s} + (R + r - \rho)xy_{14}^{\circ,s} + (R + \rho - r)xy_{31}^{\circ,s} + (R + r - \rho)xy_{14}^{\circ,s} + (R + \rho - r)xy_{23}^{\circ,s} + (R + r - \rho)xy_{14}^{\circ,s} + (R + r - \rho)xy_{23}^{\circ,s} + (R + r - \rho)xy_{23}^{\circ,s})/2 + (r + r - \rho)xy_{14}^{\circ,s} + (r + r - \rho)xy_{14}^{\circ,s} + (r + r - \rho)xy_{14}^{\circ,s} + (r + r - \rho)xy_{24}^{\circ,s} + xy_{24}^{\circ,s})/2 + (r + r - \rho)xy_{34}^{\circ,s} + (r + \rho - r)xy_{34}^{\circ,s} +$$

The full system is therefore described by 16 recurrence equations (three diallelic loci in two sexes, $2^3 \times 2 = 16$). However, not all diploid types are produced under certain sex-determining systems. For example, with the M allele fixed and an ancestral XY sex-determining system, there are XX females and XY males $(x_3^\circ = x_4^\circ = y_3^\delta = y_4^\delta = y_i^\varrho = 0, \forall i)$. In this case, the system only involves six recursion equations, which we assume below to calculate the equilibria.

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