

Across vascular land plant lineages, there is great diversity and disparity of genome size and 31

chromosome number (Table 1). Specifically, flowering plants have smaller average nuclear 32

genomes and lower chromosome numbers than most pteridophytes (ferns and lycophytes), a 33

pattern that is correlated with a difference in reproductive mode (Klekowski and Baker, 1966; 34

Wagner and Wagner, 1979; Nakazato *et al.*, 2008; Leitch and Leitch, 2013). Seed plants 35

- (including flowering plants) reproduce via separate megaspores and microspores that develop 36
- into female and male gametophytes, respectively (Fig. 1A). In these heterosporous systems, 37
- female meiosis is typically asymmetric, and male meiosis is symmetric (Fig. 2A, B). While some 38
- pteridophytes share this heterosporous system of gamete production (Pettitt, 1977; Bell, 1981), 39
- most pteridophytes and all bryophytes have a homosporous system (Fig. 1B) where symmetric 40
- meiosis leads to the production of only one type of spore, which germinates into a gametophyte 41
- that is capable of producing eggs, sperm, or both. Homospory is the ancestral state of land plants 42
- and heterospory has evolved a minimum of 11 times throughout the history of tracheophytes 43
- (Bateman and DiMichele, 1994). These transitions from homospory to heterospory are often 44
- accompanied by distinct changes in genome structure, including a decrease in chromosome 45
- number and genome size (Clark *et al.*, 2016; Carta, Bedini and Peruzzi, 2020). One exception to 46
- this pattern are gymnosperms, which are heterosporous with small chromosome numbers but 47
- unusually large genome sizes due to the accumulation of repetitive elements (e.g., Nystedt *et al.*, 48
- 2013) and few potential ancient WGD events (Li *et al.*, 2015; One Thousand Plant 49
- Transcriptomes Initiative, 2019; Stull *et al.*, 2021). 50
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Several mechanisms have been proposed to explain the differences in genome organization 52

- between homosporous and heterosporous plants (Klekowski and Baker, 1966; Wagner and 53
- Wagner, 1979; Haufler and Soltis, 1986; Haufler, 1987; Barker and Wolf, 2010; Leitch and 54
- Leitch, 2013; Kinosian, Rowe and Wolf, 2022). The most well-studied have been differences in 55
- whole genome duplications (WGDs) and diploidization (Klekowski and Baker, 1966; Wagner 56
- and Wagner, 1979; Haufler, 1987; Barker, 2013). Recent work has shown homosporous and 57
- heterosporous lineages have roughly the same number of WGDs, but likely different 58
- mechanisms or at least different rates of diploidization and fractionation (One Thousand Plant 59
- Transcriptomes Initiative, 2019; Li *et al.*, 2021; McKibben, Finch and Barker, 2024; Z. Li *et al.*, 60
- 2024). In flowering plants, diploidization and fractionation can proceed rapidly via a reduction in 61
- chromosome number (descending dysploidy) and extensive gene deletions by ectopic 62
- recombination (Wendel, 2015; Li *et al.*, 2021). Diploidization in homosporous pteridophytes 63
- appears to involve limited chromosome loss accompanied by gene silencing rather than deletion, 64
- leading to their high chromosome numbers (Haufler, 1987; Barker and Wolf, 2010; Barker, 65
- 2013; Z. Li *et al.*, 2024) and striking intra- and intergenomic collinearity (Huang *et al.*, 2022; C. 66
- Li *et al.*, 2024). Heterosporous pteridophytes, however, have chromosome numbers akin to 67
- heterosporous angiosperms (Klekowski and Baker, 1966). The transition to heterospory might 68
- lead to smaller chromosome numbers and genomes through the dynamics of post-WGD 69
- diploidization, perhaps through fundamental changes in the symmetry of female meiosis 70
- associated with heterospory. 71
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Female meiosis in heterosporous plants is asymmetric in terms of cell fate: of the four meiotic 73

- products, only one survives to become the egg cell and the three polar bodies do not enter the 74
- germline (Figs. 1A, 3A; (Burt and Trivers, 2009; Schmerler and Wessel, 2011; Haig, 2020). This 75
- asymmetry allows for a deviation in Medelian inheritance called meiotic drive (also referred to 76
- here as *female meiotic drive* or *true meiotic drive*), where genetic elements are preferentially 77
- transmitted to the germline specifically during asymmetric female meiosis (Sandler and Novitski, 78
- 1957; Pardo-Manuel de Villena and Sapienza, 2001b). Meiotic drive is a type of transmission 79
- ratio distortion (TRD) or non-random inheritance caused by different processes through an 80
- organism's life cycle, including meiosis, gametogenesis, fertilization, and development (Fishman 81
- and McIntosh, 2019). 82

- Female meiotic drive can have profound impacts on genome structure. In heterozygotes, a 84
- driving allele can rapidly rise to fixation within a handful of generations by taking advantage of a 85
- functionally asymmetric meiotic spindle pole and be preferentially transmitted into the surviving 86
- egg cell (Pardo-Manuel de Villena and Sapienza, 2001b; Fishman and Willis, 2005; Akera *et al.*, 87
- 2017; Finseth, 2023). The mechanisms of drive often act on entire chromosomes: meiotic drive 88
- can favor the inheritance of certain chromosomal rearrangements and morphologies, altering a 89
- species' karyotype and chromosome number (Pardo-Manuel de Villena and Sapienza, 2001a; 90
- Burt and Trivers, 2009; Fishman *et al.*, 2014; Baack *et al.*, 2015; Lindholm *et al.*, 2016; 91
- Blackmon *et al.*, 2019; Boman *et al.*, 2024). In addition, female meiotic drive can lead to 92
- increased centromere and chromosome size (Henikoff, Ahmad and Malik, 2001; Malik and Bayes, 2006; Zedek and Bureš, 2016; Finseth, Nelson and Fishman, 2021; Plačková *et al.*, 2022). 93 94
- Drive has been proposed as a mechanism of karyotype evolution in mammals (Pardo-Manuel de 95
- Villena and Sapienza, 2001a; Blackmon *et al.*, 2019), as well as examined in the context of 96
- symmetric and asymmetric meiosis in plants (Zedek and Bureš, 2016; Plačková *et al.*, 2024); 97
- however, it has not previously been explored as a potential mechanism for the remarkable 98
- difference in genome structure and chromosome number between heterosporous and 99
- homosporous plants. 100
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We propose that female meiotic drive may be an important force behind differences in 102

chromosome number evolution across land plants. In heterosporous angiosperms, chromosome 103

- number decreases by descending dysploidy following WGD (Li *et al.*, 2021; Z. Li *et al.*, 2024). 104
- Descending dysploidy occurs at a slower rate in homosporous pteridophytes, about half that of 105
- angiosperms (Z. Li *et al.*, 2024). This disparity could be explained by the presence of meiotic 106
- drive in heterosporous lineages and its absence in homosporous lineages. Meiotic drive is a 107
- known mechanism for chromosomal rearrangements to overcome a heterozygote disadvantage, and rearrangements involving the centromere would be preferentially inherited (White, 1968; 108 109
- Searle, 1993; Searle and de Villena, 2022). Specifically, rearrangements like Robertsonian 110
- translocations can be favored by drive and reduce overall chromosome number and could be part 111
- of how dysploidy occurs during the diploidization process (Escudero *et al.*, 2014; Searle and de 112
- Villena, 2022). Homosporous plants may go through slower post-WGD dysploid changes 113
- because they have strictly symmetric meiosis, eliminating the possibility of genome restructuring 114
- by true meiotic drive. Other types of TRD can occur in lineages with symmetric meiosis, 115
- although these processes typically occur before or after meiosis and often act on specific 116
- genotypes (Lindholm *et al.*, 2016). Here we review how female meiotic drive impacts genome 117
- evolution and chromosome number across vascular land plants (focusing on angiosperms), what 118
- genome characteristics we would expect to see in a lineage without meiotic drive (focusing on 119
- pteridophytes), how post-meiotic drive may be influencing homosporous genomes, and conclude 120
- with future directions for incorporating meiotic drive into the growing body of work on post-121
- WGD genome evolution in plants. 122
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The influence of meiotic drive on heterosporous genome structure 124

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Changes in genome structure as a consequence of female meiotic drive has been described across eukaryotes (e.g., Pardo-Manuel de Villena and Sapienza, 2001a; Chmátal *et al.*, 2014; Fishman *et al.*, 2014; Blackmon *et al.*, 2019). Female meiotic drive can happen via different mechanisms during meiosis I or II, depending on which aspects of cell division are available to create preferential inheritance (Clark and Akera, 2021). The most well-studied drive mechanism is centromere drive, which has direct effects on chromosome number, shape, and size (Blackmon *et al.*, 2019; Mayrose and Lysak, 2021; Dudka and Lampson, 2022; Plačková *et al.*, 2024). In centromere drive, cell signaling during meiosis I differentiates the egg (surviving, enters the germ line) and cortical (polar body, does not enter the germline) sides of the spindle (Figs. 2A, 3A; (Akera *et al.*, 2017; Silva and Akera, 2023). Larger centromeres have a stronger kinetochore-spindle connection, resulting in the larger homologous centromere (and attached homolog) being preferentially attached to the more stable microtubules on the egg side of the spindle (Fig. 3B; (Peris *et al.*, 2009; Akera *et al.*, 2017; Dudka and Lampson, 2022). In drive systems that favor larger centromeres, chromosome fusions can be preferentially inherited, reducing overall chromosome number (Pardo-Manuel de Villena and Sapienza, 2001a; Lindholm *et al.*, 2016). For example, Robertsonian translocations are a type of chromosome fusion where two acro- or telocentric chromosomes fuse to form a new metacentric chromosome; these fused centromeres create stronger bond to the spindle fiber (Chmátal *et al.*, 2014). Therefore, Robertsonian translocations are preferentially transmitted in human female meiosis despite deleterious effects (Daniel, 2002; Poot and Hochstenbach, 2021; Talbert and Henikoff, 2022; de Lima *et al.*, 2024); similar fused chromosomes are also preferentially transmitted in *Drosophila* (Stewart *et al.*, 2019). Chromosome number evolution mediated by chromosomal rearrangements has been documented across heterosporous plants, including Brassicaceae (Lysak *et al.*, 2006; Mandáková and Lysak, 2008), Orchidaceae (Moscone *et al.*, 2007), and Podocarpaceae (Hair and Beuzenberg, 1958; Jones, 1979), with fusions being noted as a potential mechanism of genomic downsizing and chromosome number reduction following polyploidy (Jones, 1998; Mandáková and Lysak, 2018; Mayrose and Lysak, 2021). Chromosome fusions favored by meiotic drive could be part of how heterosporous plants cytologically downsize (i.e., descending dysploidy) following WGD; the absence of meiotic drive in homosporous plants would eliminate this process as a mechanism for genome downsizing. 126 127 128 129 130 131 132 133 134 135 136 137 138 139 140 141 142 143 144 145 146 147 148 149 150 151 152 153 154 155 156

In addition to chromosome number, meiotic drive can cause different morphologies (meta-, 157

acro-, or telocentric chromosomes) to be favorably inherited depending on the system, although 158

it is unclear exactly in what stage of meiosis this occurs (Burt and Trivers, 2009; Dudka and 159

Lampson, 2022). Chromosome morphology changes caused by meiotic drive are known to occur 160

in mammals and fishes, where a change in the polarity of meiotic drive (i.e., which meiotic 161

product survives, Fig. 3A) favors a specific chromosome morphology, causing rapid 162

rearrangement of chromosome structure and number (Molina *et al.*, 2014; Blackmon *et al.*, 163

2019). For example, in mammals, some species have a karyotype comprising mostly metacentric 164

- chromosomes, some have mostly acro- or telocentric chromosomes, and others have a mix of 165
- both chromosome morphologies (Pardo-Manuel de Villena and Sapienza, 2001b; Burt and 166
- Trivers, 2009). Rates of chromosome evolution are slower in species with one chromosome 167
- morphology and higher in species with a mix of morphologies (Blackmon *et al.*, 2019). Meiotic 168
- drive is thought to stabilize chromosome morphology within a species, and drive can cause rapid karyotype change if the polarity of drive changes (Pardo-Manuel de Villena and Sapienza, 169 170
- 2001a; Blackmon *et al.*, 2019). Interestingly, most angiosperms species have only metacentric 171
- chromosomes (Stebbins and Others, 1971; Weiss-Schneeweiss and Schneeweiss, 2013), perhaps 172
- because the polarity of meiosis is weaker compared to animals (Huang and Russell, 1992). 173
- However, within some lineages like monocots, for example, karyotypes can be much more 174
- variable (Stedje, 1989; Choi *et al.*, 2008; Hamouche *et al.*, 2010; McKain *et al.*, 2012; Plačková 175
- *et al.*, 2022). Research in the Brassicaceae suggests that a karyotype comprising multiple 176
- chromosome morphologies is perhaps a transition state (Mandáková and Lysak, 2008; Lysak *et* 177
- *al.*, 2009; Weiss-Schneeweiss and Schneeweiss, 2013), which could be created by drive if the 178
- polarity of meiosis recently changed, perhaps following one of the many WGD in evolutionary 179
- history of the family (e.g., Lysak *et al.*, 2009; McKibben, Finch and Barker, 2024). 180
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- Other types of meiotic drive can occur via mechanisms in meiosis II, although the exact 182
- mechanisms by which this happens are unclear because the egg and polar sides of the spindle are 183
- not established through the same signalling mechanisms as is meiosis I (Clark and Akera, 2021). 184
- A neocentromere on chromosome Ab10 in maize acts during meiosis II, where the driving 185
- neocentromeres are transmitted to the outer cells of the tetrad, one of which survives to become 186
- the egg (Dawe *et al.*, 2018). In addition, cytokinesis in female meiosis is asymmetrical, with a 187
- majority of the cytoplasm remaining in the egg cell. Certain chromosomes can lag during 188
- cytokinesis of meiosis II and remain in the egg cytoplasm, similar to biased transmission of B 189
- chromosomes (Wu *et al.*, 2019; Chen, Birchler and Houben, 2022). 190
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As mentioned previously, the mechanisms and effects of female meiotic drive are incredibly variable. One aspect of this variability is the polarity of meiotic drive, where either large or small 192

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- centromere are favored depending on which meiotic product survives to become the egg 194
- (Blackmon *et al.*, 2019; Plačková *et al.*, 2024). It has been hypothesized that centromere size 195
- may mediate chromosome size, with larger centromeres associated with larger chromosomes, 196
- and vice versa (Plačková *et al.*, 2022, 2024). Indeed, centromere size has been observed to scale 197
- with chromosome size in Agavoideae (Plačková *et al.*, 2022) and grasses (Bennett *et al.*, 1981; 198
- Zhang and Dawe, 2012). This could extend to meiotic drive favoring different chromosome 199
- traits, depending on the polarity of drive. For example, there is evidence that centromere drive 200
- can favor chromosome fusions in one species of butterfly, but conserve chromosome structure by 201
- selecting against chromosome fusions in another (Boman *et al.*, 2024). In addition, in *Mimulus* it 202
- is thought that chromosome fissions supported by meiotic drive are responsible for chromosome 203
- number doubling, not polyploidy (Fishman et al., 2014). Considering that drive is variable across 204

systems (e.g., (Lindholm *et al.*, 2016), additional work is needed to understand the mechanisms at play. 205 206

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It is important to note that there are many ways in which angiosperm megasporogenesis happens (Huang and Russell, 1992; Kaur, Kathpalia and Koul, 2024), all having different types of cells develop into the megaspore resulting in variable mechanisms of meiotic drive. Indeed, a correlation between the type of sporogenesis and nuclear genome size has been made in angiosperms, where species that undergo disporic or tetrasporic megasporogenesis have larger genomes than those with monosporic sporogenesis (Bharathan, 1996). In monosporous megasporogenesis, following meiosis one of the four resulting one-nucleate cells survived to become the egg cell (Fig. 3A); in disporic megasporogenesis, one of the two resulting twonucelate cells survives; and in tetrasporic megasporogenesis, a single four-nucleate cell survives (e.g., Haig, 1990, 2020; Drews and Koltunow, 2011). Meiotic drive would be the strongest in a monosporic system, and present but less strong in a disporic system. In tetrasporic megasporogenesis, the egg cell develops much later, and within an embryo derived from one genetic background (Yadegari and Drews, 2004); drive would act much differently and potentially be weaker than in the other two types of megasporogenesis. Interestingly, megasporogenesis in heterosporous ferns could have stronger meiotic drive than in seed plants. While 1 out of 4 potential megaspores are ultimately viable in angiosperm monosporic sporogenesis, 1 out of 32 megaspores are ultimately viable in the germ *Marsilea*. Megasporogenesis in the fern *Marsilea* produces eight tetrads of megaspores (and in each tetrad just one megaspore survives), but only one of the eight survives to maturity (Bell, 1981). Interestingly, spores are produced in a tetrad with no polarity as to which megapore survives (Bell 1985). More research of the dynamics of meiosis in heterosporous ferns and lycophytes is needed to better understand the asymmetries in these lineages. In summary, meiotic drive can influence genome structure by the non-Mendelian inheritance of chromosomes with larger centromeres, and chromosome number by inheritance of chromosomal rearrangements involving and strengthening centromeres. The potential for meiotic drive may be part of the reason why heterosporous plants, in particular angiosperms, have a large diversity and relatively rapid evolution of genome structure and organization. Competition among centromeres and chromosomes in plant lineages with meiotic drive may lead to lower chromosome numbers especially if fusions or larger centromeres and chromosomes are favored in their drive systems. In contrast, the absence of meiotic drive in homosporous plants may explain the comparative lack of diversity and slower evolution in their genome structure and organization such as largely stable and high chromosome numbers over time. 208 209 210 211 212 213 214 215 216 217 218 219 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235 236 237 238 239 240

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Meiotic drive and symmetric meiosis 242

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- In contrast to the genomes of the heterosporous angiosperms, the genomes of plants with 244
- symmetric meiosis—namely homosporous pteridophytes and bryophytes—cannot be influenced 245
- by female meiotic drive. Homosporous pteridophytes have a unique kind of symmetric meiosis, 246
- different from that of male meiosis in heterosporous plants. In leptosporangiate ferns, spores are 247
- produced via four rounds of mitosis and one round of meiosis (Fig. 2C). In eusporangiate ferns, 248
- hundreds of spores are produced in tetrads via one round of meiosis, akin to male meiosis in heterosporous plants (Brown and Lemmon, 2001). Sporogenesis in bryophytes is also 249 250
- symmetric; however, unlike pteridophytes, some bryophytes have sex chromosomes, which 251
- could lead to alternate mechanisms of TRD (e.g., Presgraves, Severance and Wilkinson, 1997; 252
- Fedyk, Bajkowska and Chętnicki, 2005). In these plants without true meiotic drive, we expect to 253
- see slower chromosomal change, limited selection pressure on chromosome size and 254
- morphology, and larger genomes. In addition, post-WGD diploidization is fundamentally 255
- different in lineage with symmetric meiosis compared to those with asymmetric meiosis; 256
- diploidization in plants with symmetric meiosis likely involves gene silencing and fractionation, 257
- without major structural changes to the genome (Haufler, 1987; Barker and Wolf, 2010; 258
- Marchant *et al.*, 2022; C. Li *et al.*, 2024; Z. Li *et al.*, 2024). 259
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Previous work has shown that centromere drive is absent in lineages with symmetric meiosis (Zedek and Bureš, 2016; Plačková *et al.*, 2024). Evidence for this comes from molecular evolutionary analyses of histones, in particular CenH3, which is often rapidly evolving with 261 262 263

- signals of positive selection in lineages with asymmetric meiosis, but evolutionarily constrained 264
- and under purifying selection in organisms with symmetric meiosis (Zedek and Bureš, 2016; 265
- Finseth, Nelson and Fishman, 2021; Plačková *et al.*, 2024). Centromere histones are thought to 266
- be rapidly evolving to counteract the changes caused by centromere drive; thus, rapid histone
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- evolution is not expected or observed in lineages with symmetric meiosis because centromeres 268
- are conserved in the absence of meiotic drive (Zedek and Bureš, 2016; Plačková *et al.*, 2024). 269
- Comparing the evolution of CenH3 between heterosporous and homosporous pteridophytes 270
- would reveal if this pattern held across all lineages with asymmetric meiosis. 271
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The absence of meiotic drive in homosporous plants means no possibility for the preferential inheritance of certain karyotypes, chromosome morphologies, or rearrangements (Burt and Trivers, 2009), and could be part of the reason why homosporous pteridophytes have relatively stable chromosome structures (Wagner and Wagner, 1979; Bomfleur, McLoughlin and Vajda, 2014; Clark *et al.*, 2016), uniform chromosome sizes (Manton, 1950; Wagner and Wagner, 1979; Nakazato *et al.*, 2008; Clark *et al.*, 2016), and relatively slow rates of chromosome number evolution (Z. Li *et al.*, 2024). The size range variation in homosporous fern chromosomes is only 31-fold, compared to 1300-fold in heterosporous angiosperms (Clark *et al.*, 2016). It has been hypothesized that this stability and uniformity is due to a limit on the amount of genetic material in each chromosome (Liu *et al.*, 2019), but also may come from the lack of meiotic drive influencing chromosome morphological change (Plačková *et al.*, 2024). Despite the extensive research on pteridophyte cytology (e.g., Manton, 1950; Klekowski and Baker, 1966), limited information has been published on their chromosome morphology. We surveyed the literature for pteridophytes chromosome morphology data, finding only ten studies describing eleven taxa and 273 274 275 276 277 278 279 280 281 282 283 284 285 286

- a broad range of chromosome morphologies (Table 2). Further studies on homosporous and heterosporous pteridophyte chromosome morphology and karyotype structure, ideally in a 287 288
- phylogenetic context, are needed to understand what processes are influencing these traits. 289
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Transmission ratio distortion can operate in organisms with symmetric meiosis, but these types of TRD occur post-meiotically, and so are different from true meiotic drive (Sandler and Novitski, 1957; Pardo-Manuel de Villena and Sapienza, 2001b; Fishman and Willis, 2005). One example of post-meiotic drive is spore drive, which results from competition between spores containing different haplotypes, often called killer and alternative haplotypes (Raju, 1994; Grognet *et al.*, 2014; Harvey *et al.*, 2014; Lindholm *et al.*, 2016). Spore drive is best characterized in ascomycete fungi, where the spores are packaged together in an ascus. The proximity of spores in an ascus allows for a "killer" haplotype to kill or disable neighboring spores with an alternate or "target" haplotype. Similar spore production occurs in pteridophytes and bryophytes with spores packaged in a sporangium. Spore drive has not been tested in ferns, but reductions in spore numbers within a sporangium has been observed across populations of homosporous ferns (Barker and James Hickey, 2006; A. L. Grusz, Pers. Ob.) which suggests it could occur. In addition, dispersal can also influence spore drive, with drive being more prevalent if spores do not travel far from one another (Lindholm *et al.*, 2016). The majority of fern spores disperse only a few meters from the parent plant (Conant, 1978; Rose and Dassler, 2017), but because ferns produce vast numbers of spores, some do travel great distances (e.g., Tryon, 1970). The greater distance that spores disperse from one another, the smaller the potential effect from spore drive (Lindholm *et al.*, 2016). 291 292 293 294 295 296 297 298 299 300 301 302 303 304 305 306 307 308

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Another example of post-meiotic drive is male drive, which occurs when a drive locus kills 310

sperm with a target locus (e.g., Taylor, 1994; Presgraves, Severance and Wilkinson, 1997). 311

Similar to spore drive, one genotype will kill the an alternate genotype regardless of the fitness 312

consequences (Rice, 2013; Lindholm *et al.*, 2016). There is limited research on fern sperm competition, but observational studies have described competition in the archegonium before 313 314

sperm meet the egg (Lopez-Smith and Renzaglia, 2008). Further work is needed to understand 315

competition and potential male drive among sperm of homosporous plants, as well as in other 316

- flagette plants. 317
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Because of fundamental differences in meiosis, meiotic drive cannot happen in homosporous pteridophytes in the same way as in heterosporous plants. TRD can still influence the genomes of homosporous plants by removing certain genotypes but likely does not shape the genome in the same ways as true meiotic drive. Several homosporous pteridophyte genome assemblies are now published (Fang *et al.*, 2021; Huang *et al.*, 2022; Marchant *et al.*, 2022; Yu *et al.*, 2023; C. Li *et al.*, 2024), giving us insight into plant genome structure in the absence of female meiotic drive. Perhaps most striking is that these species have highly conserved intra- and intergenomic synteny over millions of years (Huang *et al.*, 2022; C. Li *et al.*, 2024); the same is true for hornworts (Schafran *et al.*, 2025). An exception is *Ceratopteris richardii*, which has a rate of 319 320 321 322 323 324 325 326 327

fractionation rivaling many angiosperm species (Nakazato *et al.*, 2006; Marchant *et al.*, 2022), 328

but the mechanisms causing this rapid diploidization are unclear. Broad patterns in homosporous 329

pteridophytes suggest that, while genic diploidization and fractionation occur, there are limited 330

mechanisms for cytologically restructuring the genome following WGD. 331

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Conclusions and future work 333

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Here we describe how meiotic drive may be a potential mechanism behind the disparity in genome architecture among land plants, specifically influencing chromosome number and structure of heterosporous plants. Other types of TRD can occur in homosporous lineages, but not true meiotic drive, as sporogenesis is symmetric. Although the symmetry of meiosis has not been explicitly explored in the context of homosporous plants, other authors have noted that genome evolution is likely dependent on the mechanics of meiosis (e.g., Burt and Trivers, 2009; Lindholm *et al.*, 2016; Kinosian, Rowe and Wolf, 2022; Finseth, 2023). Pteridophytes are an ideal system to study the effects of drive on genome structure, as there are sister lineages with asymmetric and symmetric meiosis in both ferns and lycophytes. Here we discuss a few ways to investigate how drive or other types of transmission ratio distortion might affect homosporous 335 336 337 338 339 340 341 342 343 344

- vascular land plant genomes. 345
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Consistent with the absence of meiotic drive, past studies have shown that homosporous ferns have strong Mendelian inheritance (Andersson-Kottö, 1927; Dedera and Werth, 1987). Ferns have relatively low gene densities compared to flowering plants (Rabinowicz *et al.*, 2005; Wolf *et al.*, 2015), perhaps because of a lack of meiotic drive selecting for genome downsizing or, 347 348 349 350

- alternatively, pseudogenization caused by transposons in ferns (Lisch, 2013). Long terminal 351
- repeat retrotransposon (LTR-RT) have a high birth but low death rate in homosporous ferns, with the inverse occurring in heterosporous lycophytes (Wang *et al.*, 2021; Yu *et al.*, 2023). Older 352 353
- LTR-RTs are associated with larger haploid nuclear genome size in pteridophytes (Baniaga and 354
- Barker, 2019). The exact role of transposons in unclear in pteridophytes, but more research is 355
- warranted, especially because transposons are known to be associated with meiotic drive 356
- (Eickbush, Young and Zanders, 2019; Vogan *et al.*, 2021). In addition, the expansion of different 357
- gene families as well as the rate of meiosis gene evolution is greater in heterosporous 358
- angiosperms than in homosporous pteridophytes, suggesting that different selection pressures, 359
- perhaps one of those being meiotic drive, are acting on meiosis in heterosporous lineages 360
- (Dhakal, Wolf and Harkess, 2024; Z. Li *et al.*, 2024). Building on these observations, future 361
- work could look more explicitly for evidence of TRD in homosporous plants and investigate the mechanics of homosporous meiosis. 362 363
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Finding natural examples of meiotic drive can be challenging, as a driving locus is often rapidly 365

- fixed in a population, and therefore undetectable. However, these can reemerge following a 366
- hybridization event, or even be part of hybrid sterility (Lindholm *et al.*, 2016). An example of 367
- this is in yellow monkeyflowers, where hybrids between *Mimulus guttatus* and *M. nasutus* 368

exhibit strong female meiotic drive and the driving locus is nearly fixed in only a handful of generations (Fishman *et al.*, 2001; Fishman and Willis, 2005; Finseth, Nelson and Fishman, 2021). Testing for such cryptic drive systems could be a potential avenue of research in 369 370 371

- pteridophytes, specifically in those with known hybrid sterility. Pteridology has a rich history of 372
- work on hybrid species, including studies on the model system *Ceratopteris* (e.g., Hickok, 1973; 373
- Hickok and Klekowski, 1974). Genomic resources available for *Ceratopteris* (Nakazato *et al.*, 374
- 2006; Marchant *et al.*, 2022) would make testing for drive in a hybrid mapping population 375
- feasible (e.g., Fishman and Willis 2005). It will also be important to look at TRD and sex chromosome drive in bryophytes. Segregation distortion of loci and biased sex ratios have been 376
- detected in the moss *Ceratodon* (McDaniel, Willis and Shaw, 2007), but less is known about 377 378
- hornworts and liverworts. 379
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A post-meiotic TRD mechanism that could be happening in homosporous plants is spore drive. 381

In fungi, spores are affected by drive when a certain genotype kills an alternative genotype. 382

Considering the large number of spores produced within a sporangium by both eusporangiate 383

and leptosporangiate lineages, the ingredients for spore drive are present. DNA expression 384

studies have identified spore drive systems, revealing that the driving element can lethally target 385

specific genes (Urquhart and Gardiner, 2023). A consistent reduction in spore number per sporangium in leptosporangiate ferns has been observed (Barker and James Hickey, 2006; A. L. 386 387

Grusz, Pers. Ob.), suggesting something may be happening during sporogenesis to reduce spore 388

number. Additional work counting spores could better characterize this pattern. If more concrete 389

patterns of spore reduction are observed and connected to a spore drive system, the model fern 390

Ceratopteris could help understand the genetic mechanisms for homosporous spore drive. 391 392

Other processes outside of meiotic and related drive systems could affect chromosome number and genome size. Population size can be a greater force on chromosome number than meiotic drive in Carnivora: smaller populations had greater variation in chromosome number than large populations (Jonika *et al.*, 2024). Another hypothesis for the evolution of chromosome number is the minimum interaction theory, which postulates smaller chromosomes are selected to reduce reciprocal translocation; consequently, chromosome number will increase as more chromosomes are needed to contain the genome (Imai *et al.*, 1986). Exploring ecology and natural history in homosporous plants could be another avenue to understand their distinct genome organization outside of meiotic drive. It is well-established that the rate of molecular evolution is higher and genome size is smaller in plants with shorter life cycles, compared to those with longer life cycles (Cavalier-Smith, 2005; Smith and Donoghue, 2008). There is also a correlation between vascular structure (i.e., the presence or absence of tracheids) and genome size (Cavalier-Smith, 2005). Heterosporous ferns (Salviniales) and *Selaginella* have tracheids and small genome sizes, but gymnosperms and the heterosporous lycophyte *Isoëtes* lack tracheids and have large genomes (Cavalier-Smith, 1978; Leitch *et al.*, 2005). Comparing life history and genome traits in a phylogenetic context could help tease apart complex interactions between these variables. 393 394 395 396 397 398 399 400 401 402 403 404 405 406 407 408 409

- mechanisms by which this occurs are incredibly varied (Burt and Trivers, 2009; Lindholm *et al.*,
- 2016; Saupe and Johannesson, 2022). Female meiotic drive is potentially a mechanism behind
- the rapid diploidization and downsizing following WGD in heterosporous plants. TRD in
- pteridophytes and bryophytes is probably distinct from true meitoic drive in angiosperms, fungi,
- or animals. The absence of meiotic drive would help explain large and stable genomes, uniform chromosome sizes, relatively static chromosome numbers following WGDs, as well as the
- current challenges with assembling centromeres in homosporous fern and lycophyte genomes.
- Although almost certainly only one piece of the puzzle, researching meiotic drive in
- homosporous plants is important to broaden our understanding of the unique genomes of this
- lineage.
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Group	Average Genome size $(1C)$	Average Haploid Chromosome Number (n)	Average Inferred rounds of WGD	Asymmetric meiosis	Potential types of drive	Citations
Angiosperm	4.4	15.99	3.5	Yes	Centromere, chromosome, female meiotic drive	(Klekowski and Baker, 1966; One Thousand Plant Transcriptomes Initiative, 2019; Pellicer and Leitch, 2020; McKibben, Finch and Barker, 2024; Z. Li et al., 2024)
Gymnosperm	15.51	11.69	1.63	Yes	Centromere, chromosome, female meiotic drive	(Rice et al., 2015; One Thousand Plant Transcriptomes Initiative, 2019; Pellicer and Leitch, 2020; Z. Li et al., 2024)
Heterosporous pteridophyte	1.01	13.62	2.4	Yes	Centromere, chromosome, female meiotic drive	(Klekowski and Baker, 1966; Pellicer and Leitch, 2020; Z. Li et al., 2024)
Homosporous pteridophyte	15.13	57.5	2.82	No	Spore drive, other non- meiotic TRD	(Klekowski and Baker, 1966; Pellicer and Leitch, 2020; Z. Li et al., 2024)
Bryophytes	0.72	10.23	2.28	$\rm No$	Spore drive, other non- meiotic TRD, sex chromosome drive	(Klekowski and Baker, 1966; Pellicer and Leitch, 2020; Z. Li et al., 2024) N. Patel, unpublished data)

Table 1. Genome traits and potential types of meiotic drive for land plants. 451

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 Diploid sporophyte

Figure 1. Life cycles of heterosporous (top) and homosporous (bottom) land plants. Locations of female meiotic drive and other types of transmission ratio distortion are denoted in black and gray boxes, respectively.

Fertilization

Gametophyte (male)

 $-$ Sperm

Gametophyte
(female or hermaphrodite)

Egg

Figure 2. Examples of different types of meiosis. In each panel, the far left circle depicts the sporocyte, and succeeding divisions proceeding to the right. Solid lines indicate meiosis and dotted lines indicate mitosis. A) Heterosporous female meiosis, where only one meiotic product survives and the remaining three polar bodies die (can be either top or bottom, resulting from the cell on the chalazal or micropylar end surviving, see Fig. 3A) ; B) Heterosporous male meiosis, where all meiotic products survive; C) Leptosporagiate sporogenesis where 64 spores are generated through four rounds of mitosis and one round of meiosis; all spore survive. 504 505 506 507 508 509 510

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- **Figure 3**. A) Organization of megasporogenesis in a heterosporous ovule, the cell on the chalazal 513
- end often survives, but alternatively the cell on the micropylar end can become the egg. B) 514
- Centromere drive, where larger centromeres are preferentially transmitted to the egg pole in 515
- meiosis I (anaphase I is depicted); the egg pole is often associated with the chalazal end of the 516
- ovule (see panel A), and the cortical side the micropylar end. 517

Glossary 518

- 519
- **Transmission ratio distortion** non-random inheritance of parental alleles in the offspring 520

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- population, occuring before, during, or after meiosis. 521
- **Meiotic drive** non-random inheritance of alleles, chromosomes, or other genome structures 522
- specifically during asymmetrical meiosis. 523
- **Spore drive** observed in ascomycete fungi where spores are packages together in an ascus, a 524
- spore with the "killer" haplotype will kill spores with the alternative haplotype, particularly in when spores are in close proximity. 525 526
- **Male drive** biased transmission that occurs during or after male meiosis; typically involves an 527
- allele that kills certain sperm cells. 528
- **Sporogenesis** the generation of a spore cell by meiosis (in heterosporous plants) or mitosis (in homosporous plants) 529 530
- **Megasporogenesis** creation of spores that will develop into a female gametophyte 531
- (megagametophyte) 532
- **Microsporogenesis** creation of spores that will develop into a male gametophyte 533
- (megagametophyte) 534
- **Homosporous** an organism that produces one type of spore. This spore develops into a 535
- gametophyte which can produce male and/or female gametes. 536
- **Heterosporous** an organism that produces two types of spores. These are the mega- and 537
- microspores, which develop into gametophytes capable of producing eggs or sperm, respectively. 538
- **Pteridophyte** a plant classified as a fern or lycophyte. This group is paraphyletic but is 539
- functionally important because all taxa are seedless vascular plants. 540
- **Leptosporangiate fern** ferns that produce sporangia with a long stalk that arise from a single 541
- epidermal cell. Their spore production proceeds from a single spore mother cell through four 542
- rounds of mitosis and one round of meiosis to form 64 spores. 543
- **Eusporangiate fern** ferns that produce sporangia with a short or no stalk that arise from 544
- several epidermal cells. Hundreds to thousands of spores are produced in each sporangium via meiosis. 545 546
- **Metacentric** a chromosome with the centromere positioned in the middle, with both arms 547
- being the same length. 548
- **Acrocentric** a chromosome with the centromere close to one end, with one arm being much 549
- longer than the other. 550
- **Telocentric** a chromosome where the centromere is at one end, causing there to be effectively only one arm rather than two. 551 552
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