1	The role of meiotic drive in chromosome number disparity between heterosporous and
2	homosporous plants
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11	Abstract
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13	In vascular plants, heterosporous lineages typically have fewer chromosomes than homosporous
14	lineages. The underlying mechanism causing this disparity has been debated for over half a
15	century. Although reproductive mode has been identified as critical to these patterns, the
16	symmetry of meiosis during sporogenesis has been overlooked as a potential cause of the
17	difference in chromosome numbers. In most heterosporous plants, meiosis during
18	megasporogenesis is asymmetric, meaning one of the four meiotic products survives to become
19	the egg. Comparatively, meiosis is symmetric in homosporous megasporogenesis and all meiotic
20	products survive. The symmetry of meiosis is important because asymmetric meiosis enables
21	meiotic drive and associated genomic changes, while symmetric meiosis cannot lead to meiotic
22	drive. Meiotic drive is a deviation from Mendelian inheritance where genetic elements are
23 24	preferentially inherited by the surviving egg cell, and can profoundly impact chromosome (and genome) size, structure, and number. Here we review how meiotic drive impacts chromosome
24 25	number evolution in heterosporous plants, how the lack of meiotic drive in homosporous plants
23 26	impacts their genomes, and explore future approaches to understand the role of meiotic drive on
27	chromosome number across land plants.
28	F F
29	Introduction
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31 Across vascular land plant lineages, there is great diversity and disparity of genome size and

chromosome number (Table 1). Specifically, flowering plants have smaller average nuclear
 genomes and lower chromosome numbers than most pteridophytes (ferns and lycophytes; Fig.

- genomes and lower chromosome numbers than most pteridophytes (ferns and lycophytes; Fig.
 1), a pattern that is correlated with a difference in reproductive mode (Wagner and Wagner 1979;
- 35 Klekowski and Baker 1966; Leitch and Leitch 2013; Nakazato et al. 2008). Seed plants
- 36 (including flowering plants) reproduce via separate megaspores and microspores that develop
- 37 into female and male gametophytes, respectively (Fig. 2A). In these heterosporous systems,
- 38 female meiosis is typically asymmetric, and male meiosis is symmetric (Fig. 3A, B). Although
- 39 some pteridophytes (Salviniales, Isoeteales, and Selaginellales) share this heterosporous system
- 40 of gamete production (Bell 1981; Pettitt 1977), most pteridophytes and all bryophytes have a
- 41 homosporous system (Fig. 2B) where symmetric meiosis leads to the production of only one type

- 42 of spore that germinates into a gametophyte that is capable of producing eggs, sperm, or both.
- 43 Homospory is the ancestral character state of land plants and heterospory has evolved
- 44 independently a minimum of 11 times throughout the history of tracheophytes (Bateman and
- 45 DiMichele 1994), including three extant lineages. These transitions from homospory to
- 46 heterospory are often accompanied by distinct changes in genome structure, including a decrease
- 47 in chromosome number and genome size (Clark et al. 2016; Carta, Bedini, and Peruzzi 2020).
- 48 One exception to this pattern is gymnosperms, which are heterosporous with small chromosome
- 49 numbers but unusually large genome sizes due to the accumulation of repetitive elements (e.g.,
- 50 (Nystedt et al. 2013) and few potential ancient WGD events (Li et al. 2015; One Thousand Plant
- 51 Transcriptomes Initiative 2019; Stull et al. 2021; Wan et al. 2018; Liu et al. 2022; Wan et al.
- 52 2022).
- 53

54 Several mechanisms have been proposed to explain the differences in genome organization

- 55 between homosporous and heterosporous plants (Barker and Wolf 2010; Kinosian, Rowe, and
- 56 Wolf 2022; Haufler and Soltis 1986; Haufler 1987; Wagner and Wagner 1979; Klekowski and
- 57 Baker 1966; Leitch and Leitch 2013). The most well-studied have been differences in whole
- 58 genome duplications (WGDs) and diploidization (Haufler 1987; Wagner and Wagner 1979;
- 59 Barker 2013; Klekowski and Baker 1966). Recent work has shown homosporous and
- 60 heterosporous lineages have roughly the same number of WGDs, but likely different
- 61 mechanisms or at least different rates of diploidization and fractionation (One Thousand Plant
- 62 Transcriptomes Initiative 2019; McKibben, Finch, and Barker 2024; Li et al. 2021, 2024). In
- 63 flowering plants, diploidization and fractionation can proceed rapidly via a reduction in
- 64 chromosome number (descending dysploidy) and extensive gene deletions by ectopic
- 65 recombination (Li et al. 2021; Wendel 2015). Diploidization in homosporous pteridophytes
- 66 appears to involve limited chromosome loss accompanied by gene silencing rather than deletion,
- 67 leading to their high chromosome numbers (Haufler 1987; Barker 2013; Barker and Wolf 2010;
- Li et al. 2024; Gastony 1991) and striking intra- and intergenomic collinearity (Li et al. 2024;
- 69 Huang et al. 2022). Heterosporous pteridophytes, however, have chromosome numbers akin to
- 70 heterosporous angiosperms (Klekowski and Baker 1966). The transition to heterospory might
- 71 lead to smaller chromosome numbers and genomes through the dynamics of post-WGD
- 72 diploidization, perhaps through fundamental changes in the symmetry of female meiosis
- 73 associated with heterospory.
- 74

Female meiosis in heterosporous plants is asymmetric in terms of cell fate: of the four meioticproducts, only one survives to become the egg cell and the three polar bodies do not enter the

- 77 germline (Figs. 1A, 3A; Schmerler and Wessel 2011; Burt and Trivers 2009; Haig 2020). This
- asymmetry allows for a deviation in Medelian inheritance called meiotic drive (also referred to
- 79 here as *female meiotic drive* or *true meiotic drive*), where genetic elements are preferentially
- 80 transmitted to the germline specifically during asymmetric female meiosis (Sandler and Novitski
- 81 1957; Pardo-Manuel de Villena and Sapienza 2001b). Meiotic drive is a type of transmission
- 82 ratio distortion (TRD) or non-random inheritance caused by different processes through an

83 organism's life cycle, including meiosis, gametogenesis, fertilization, and development (Fishman

- and McIntosh 2019).
- 85

Female meiotic drive can have profound impacts on genome structure. In heterozygotes, a 86 87 driving allele can rapidly rise to fixation within a handful of generations by taking advantage of a 88 functionally asymmetric meiotic spindle pole and be preferentially transmitted into the surviving egg cell (Pardo-Manuel de Villena and Sapienza 2001b; Fishman and Willis 2005; Finseth 2023; 89 Akera et al. 2017). The mechanisms of drive often act on entire chromosomes: meiotic drive can 90 91 favor the inheritance of certain chromosomal rearrangements and morphologies, altering a 92 species' karyotype and chromosome number (Burt and Trivers 2009; Fishman et al. 2014; 93 Boman et al. 2024; Blackmon et al. 2019; Baack et al. 2015; Lindholm et al. 2016; Pardo-Manuel de Villena and Sapienza 2001a). In addition, female meiotic drive can lead to changes in 94 centromere and chromosome size (Zedek and Bureš 2016; Finseth, Nelson, and Fishman 2021; 95 96 Malik and Bayes 2006; Henikoff, Ahmad, and Malik 2001; Plačková et al. 2022). Drive has been 97 proposed as a mechanism of karyotype evolution in mammals (Pardo-Manuel de Villena and Sapienza 2001a; Blackmon et al. 2019), as well as examined in the context of symmetric and 98 99 asymmetric meiosis (Zedek and Bureš 2016; Plačková et al. 2024) and heterosporous and 100 homosporous reproductive modes (Kinosian, Rowe, and Wolf 2022) in plants. 101

102 We propose that female meiotic drive may be an important force behind differences in

chromosome number evolution across land plants. In heterosporous angiosperms, chromosome 103 number decreases by descending dysploidy following WGD (Li et al. 2024, 2021). Descending 104 105 dysploidy occurs at a slower rate in homosporous pteridophytes, about half that of angiosperms (Li et al. 2024). As discussed by (Kinosian, Rowe, and Wolf 2022) and (Plačková et al. 2024), 106 107 this disparity could be explained by the presence of meiotic drive in heterosporous lineages and its absence in homosporous lineages. Meiotic drive is a known mechanism for chromosomal 108 rearrangements to overcome a heterozygote disadvantage, and rearrangements involving the 109 110 centromere would be preferentially inherited (Searle and de Villena 2022; Searle 1993; White 111 1968). Specifically, rearrangements like Robertsonian translocations can be favored by drive and 112 reduce overall chromosome number and could be part of how dysploidy occurs during the 113 diploidization process (Searle and de Villena 2022; Escudero et al. 2014). Homosporous plants may go through slower post-WGD dysploid changes because they have strictly symmetric 114 115 meiosis, eliminating the possibility of genome restructuring by true meiotic drive. Other types of TRD can occur in lineages with symmetric meiosis, although these processes typically occur 116 117 before or after meiosis and often act on specific genotypes (Lindholm et al. 2016). Here we 118 review how female meiotic drive impacts genome evolution and chromosome number across 119 vascular land plants with asymmetric meiosis (focusing on angiosperms), what genome characteristics and types of TRD we would expect to see in a lineage with symmetric meiosis 120 121 (focusing on pteridophytes), how homosporous genomes may evolve in the absence of meiotic 122 drive, and conclude with future directions for incorporating meiotic drive into the growing body 123 of work on post-WGD genome evolution in plants.

125 The influence of meiotic drive on heterosporous genome structure

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127 Changes in genome structure as a consequence of female meiotic drive have been described across eukaryotes (e.g., Blackmon et al. 2019; Chmátal et al. 2014; Pardo-Manuel de Villena and 128 Sapienza 2001a; Fishman et al. 2014). Female meiotic drive can happen via different 129 mechanisms during meiosis I or II, depending on which aspects of cell division are available to 130 create preferential inheritance (Clark and Akera 2021). The most well-studied drive mechanism 131 is centromere drive, which has direct effects on chromosome number, shape, and size (Mayrose 132 133 and Lysak 2021; Dudka and Lampson 2022; Plačková et al. 2024; Blackmon et al. 2019). In centromere drive, cell signaling during meiosis I differentiates the egg (surviving, enters the 134 germ line) and cortical (polar body, does not enter the germline) sides of the spindle (Figs. 2A, 135 136 3A; Akera et al. 2017; Silva and Akera 2023). Larger centromeres have a stronger kinetochore-137 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. 4B; 138 Dudka and Lampson 2022; Akera et al. 2017; Peris et al. 2009). In drive systems that favor 139 140 larger centromeres, chromosome fusions can be preferentially inherited, reducing overall chromosome number (Pardo-Manuel de Villena and Sapienza 2001a; Lindholm et al. 2016). For 141 142 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 143 144 create stronger bond to the spindle fiber (Chmátal et al. 2014). Therefore, Robertsonian translocations are preferentially transmitted in human female meiosis despite deleterious effects 145 146 (Talbert and Henikoff 2022; Poot and Hochstenbach 2021; Daniel 2002; de Lima et al. 2024); similar fused chromosomes are also preferentially transmitted in *Drosophila* (Stewart et al. 147 148 2019). Chromosome number evolution mediated by chromosomal rearrangements has been documented across heterosporous plants, including Brassicaceae (Mandáková and Lysak 2008; 149 Lysak et al. 2006), Orchidaceae (Moscone et al. 2007), and Podocarpaceae (Hair and 150 151 Beuzenberg 1958; Jones 1979), with fusions being noted as a potential mechanism of genomic 152 downsizing and chromosome number reduction following polyploidy (Jones 1998; Mayrose and Lysak 2021; Mandáková and Lysak 2018). Chromosome fusions favored by meiotic drive could 153 154 be an important part of how heterosporous plants cytologically downsize (i.e., descending dysploidy) following WGD; the absence of meiotic drive in homosporous plants would eliminate 155 156 this process as a mechanism for genome downsizing. 157 158 In addition to chromosome number, meiotic drive can cause different morphologies (meta-, 159 acro-, or telocentric chromosomes) to be favorably inherited depending on the system, although 160 it is unclear exactly in what stage of meiosis this occurs (Burt and Trivers 2009; Dudka and

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

- 162 in mammals and fishes, where a change in the polarity of meiotic drive (i.e., which meiotic
- 163 product survives, Fig. 4A) favors a specific chromosome morphology, causing rapid
- 164 rearrangement of chromosome structure and number (Blackmon et al. 2019; Molina et al. 2014).

165 For example, in mammals, some species have a karyotype comprising mostly metacentric

- 166 chromosomes, some have mostly acro- or telocentric chromosomes, and others have a mix of
- both chromosome morphologies (Burt and Trivers 2009; Pardo-Manuel de Villena and Sapienza2001b). Rates of chromosomal evolution are slower in species with one chromosome
- 169 morphology and higher in species with a mix of morphologies (Blackmon et al. 2019). Meiotic
- 170 drive is thought to stabilize chromosome morphology within a species, and drive can cause rapid
- 171 karvotype change if the polarity of drive changes (Blackmon et al. 2019; Pardo-Manuel de
- 172 Villena and Sapienza 2001a). Interestingly, most angiosperms species have only metacentric
- 173 chromosomes (Weiss-Schneeweiss and Schneeweiss 2013; Stebbins and Others 1971), perhaps
- because the polarity of meiosis is weaker compared to animals (Huang and Russell 1992).
- 175 However, within some lineages like monocots, for example, karyotypes can be much more
- 176 variable (Hamouche et al. 2010; Choi et al. 2008; Stedje 1989; McKain et al. 2012; Plačková et
- al. 2022). Research in the Brassicaceae suggests that a karyotype comprising multiple
- 178 chromosome morphologies is perhaps a transition state (Mandáková and Lysak 2008; Lysak et
- al. 2009; Weiss-Schneeweiss and Schneeweiss 2013) that could be created by drive if the
- 180 polarity of meiosis recently changed, perhaps following one of the many WGD in evolutionary
- history of the family (e.g., McKibben, Finch, and Barker 2024; Lysak et al. 2009; Mabry et al.
- 182 2020; Mandáková et al. 2017; Mabry et al. 2024).
- 183

184 Other types of meiotic drive can occur via mechanisms in meiosis II, although the exact

- 185 mechanisms by which this happens are unclear because the egg and polar sides of the spindle are
- 186 not established through the same signalling mechanisms as is meiosis I (Clark and Akera 2021).
- 187 A neocentromere on chromosome Ab10 in maize acts during meiosis II, where the driving
- 188 neocentromeres are transmitted to the outer cells of the tetrad, one of which survives to become
- the egg (Dawe et al. 2018). In addition, cytokinesis in female meiosis is asymmetrical, with a
- 190 majority of the cytoplasm remaining in the egg cell. Certain chromosomes can lag during
- 191 cytokinesis of meiosis II and remain in the egg cytoplasm, similar to biased transmission of B
- 192 chromosomes (Wu et al. 2019; Chen, Birchler, and Houben 2022).
- 193

As mentioned previously, the mechanisms and effects of female meiotic drive are incredibly 194 195 variable. One aspect of this variability is the polarity of meiotic drive, where either large or small centromeres are favored depending on which meiotic product survives to become the egg 196 (Blackmon et al. 2019; Plačková et al. 2024). It has been hypothesized that centromere size may 197 mediate chromosome size, with larger centromeres associated with larger chromosomes, and 198 199 vice versa (Plačková et al. 2022, 2024). Indeed, centromere size has been observed to scale with 200 chromosome size in Agavoideae (Plačková et al. 2022) and grasses (Zhang and Dawe 2012; 201 Bennett et al. 1981). This could extend to meiotic drive favoring different chromosome traits, depending on the polarity of drive. For example, there is evidence that centromere drive can 202 203 favor chromosome fusions in one species of butterfly, but conserve chromosome structure by 204 selecting against chromosome fusions in another (Boman et al. 2024). In addition, in Mimulus it 205 is thought that chromosome fissions supported by meiotic drive are responsible for chromosome

number doubling, not polyploidy (Fishman et al., 2014). Considering that drive is variable across
systems (e.g., Lindholm et al. 2016), additional work is needed to understand the mechanisms at
play.

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210 It is important to note that there are many ways in which angiosperm megasporogenesis happens (Huang and Russell 1992; Kaur, Kathpalia, and Koul 2024), with different types of cells 211 212 developing into the megaspore resulting in variable mechanisms of meiotic drive (Fig. 3D, 4A). 213 Indeed, a correlation between the type of sporogenesis and nuclear genome size has been made 214 in angiosperms species that undergo disporic or tetrasporic megasporogenesis have larger 215 genomes than those with monosporic sporogenesis (Bharathan 1996). In monosporic 216 megasporogenesis, following meiosis one of the four resulting one-nucleate cells survived to become the egg cell; in disporic megasporogenesis, one of the two resulting two-nucleate cells 217 218 survives; and in tetrasporic megasporogenesis, a single four-nucleate cell survives (e.g., Drews 219 and Koltunow 2011; Haig 1990, 2020). Meiotic drive would be the strongest in a monosporic system, and present but less strong in a disporic system. In tetrasporic megasporogenesis, the egg 220 221 cell develops much later, and within an embryo derived from one genetic background (Yadegari 222 and Drews 2004). As a result, drive would act differently and potentially be weaker than in the 223 other two types of megasporogenesis.

224

225 Megasporogenesis in heterosporous ferns has similar variability in the degree of asymmetry as

angiosperm megasporogenesis. In the heterosporous fern *Marsila*, 1 out of 32 megaspores are

227 ultimately viable. Megasporogenesis in *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). The final surviving megaspore typically comes from the tetrad closest to the attachment

230 point of the sporangium to its stalk (Sheffield and Bell 1987). Comparatively, in the

231 heterosporous lycophyte *Selaginella* one or two meiotic products survive megasporogenesis

232 (John M. Pettitt 1971), and micosporogenesis can also be asymmetric (Pettitt 1977). Finally, all

233 meiotic products in the heterosporous lycophyte *Isoetes englemanii* survive (Pettitt 1976),

eliminating the possibility of meiotic drive in this lineage. More research on the dynamics of

meiosis in heterosporous ferns and lycophytes is needed to better understand the asymmetriesand types of drive in these lineages.

237

It is important to note that microsporogenesis can be asymmetric in the case of angiosperms with pseudomonad pollen, which presents an opportunity for meiotic drive (Furness and Rudall 2011).

At maturity, pollen often comprises solitary grains (monads), but can be in groups of four

241 (tetrads). It is hypothesized that pseudomonad pollen evolved from tetrad pollen where three of

the four cells produced by meiosis do not mature (Walker and Doyle 1975). Pseudomonad pollen

243 has evolved independently multiple times in angiosperms, including in the monocot family

244 Cyperaceae (sedges) and the eudicot subfamily Styphelioideae (Ericaeae, heathers), with the

245 developmental mechanisms being slightly different in each lineage (Brown and Lemmon 2000;

246 Furness 2009). Several authors have speculated that meiotic drive is present in lineages with

- 7
- 247 pseudomonad pollen, perhaps causing the large variation in chromosome number observed in
- these groups (Furness and Rudall 2011; Hipp 2007). Cyperaceae and Styphelioideae have a high
- frequency of aneuploidy, with the most extreme example being the large genus *Carex*
- 250 (Cyperaceae) varying from *n*=6-68 (Davies 1956). Interestingly, Cyperaceae frequently have
- 251 holocentric chromosomes (diffuse centromeres) while Styphelioideae do not, presenting a unique
- 252 comparative system to look at how holocentric chromosomes and meiotic drive interact (e.g.,
- (Krátká et al. 2021). These examples of asymmetric male meiosis in heterosporous plants offer a
 different way to look at the role of meiotic drive in chromosome number and genome structure
- 255 evolution, as the asymmetry of microsporogeneis could have similar consequences as
- 256 asymmetric megasporogenesis, despite different mechanisms. In addition, it will be important to
- consider the possibility of asymmetrical meiosis in homosporous plants and the different ways
- 258 drive may impact those systems.
- 259

260 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 261 262 rearrangements involving and strengthening centromeres. The potential for meiotic drive may be 263 part of the reason why heterosporous plants, in particular angiosperms, have a large diversity and 264 relatively rapid evolution of genome structure and organization. Competition among centromeres 265 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. 266 267 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 268

- 269 stable and high chromosome numbers over time.
- 270

271 Meiotic drive and symmetric meiosis

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273 In contrast to the genomes of the heterosporous angiosperms, the genomes of plants with 274 symmetric meiosis—namely homosporous pteridophytes and bryophytes—cannot be influenced 275 by female meiotic drive. Homosporous pteridophytes have a unique kind of symmetric meiosis, different from that of male meiosis in heterosporous plants. In leptosporangiate ferns, spores are 276 277 produced via four rounds of mitosis and one round of meiosis (Fig. 3C). In eusporangiate ferns, hundreds of spores are produced in tetrads via one round of meiosis, akin to male meiosis in 278 279 heterosporous plants (Brown and Lemmon 2001). Sporogenesis in bryophytes is also symmetric; however, unlike pteridophytes, some bryophytes have sex chromosomes, which could lead to 280 other mechanisms of TRD (e.g., Presgraves, Severance, and Wilkinson 1997; Fedvk, Bajkowska, 281 and Chetnicki 2005; Úbeda, Patten, and Wild 2015). In these plants without true meiotic drive, 282 283 we expect to see slower chromosomal change, limited selection pressure on chromosome size and morphology, and larger genomes. Through empirical work on homosporous pteridophytes, 284 285 we observe high chromosome numbers (Klekowski and Baker 1966) and large genomes (Pellicer 286 and Leitch 2020; Kinosian, Rowe, and Wolf 2022), slow rates of dysploidy (Li et al. 2024) and 287 genome size change (Clark et al. 2016), as well as limited gene order (collinearity) change across

- hundreds of millions of years (Huang et al. 2022; Li et al. 2024). These observations are
- consistent with the hypothesis that diploidization in plants with symmetric meiosis is
- 290 fundamentally different than in plants with asymmetric meiosis, potentially due to the absence of
- 291 meiotic drive.
- 292

293 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 294 evolutionary analyses of histones, in particular CenH3, which is often rapidly evolving with 295 signals of positive selection in lineages with asymmetric meiosis, but evolutionarily constrained 296 297 and under purifying selection in organisms with symmetric meiosis (Zedek and Bureš 2016; F. 298 R. Finseth, Nelson, and Fishman 2021; Plačková et al. 2024). Centromere histones are thought to 299 be rapidly evolving to counteract the changes caused by centromere drive. Thus, rapid histone 300 evolution is not expected or observed in lineages with symmetric meiosis because centromeres 301 are conserved in the absence of meiotic drive (Zedek and Bureš 2016; Plačková et al. 2024). In 302 previous work, all pteridophytes have been categorized as having symmetric meiosis, so 303 investigating the evolution of CenH3 in heterosporous pteridophytes would reveal if this pattern 304 held across all lineages with asymmetric meiosis.

305

306 The absence of meiotic drive in homosporous plants means no possibility for the preferential

- 307 inheritance of certain karyotypes, chromosome morphologies, or rearrangements (Burt and
- 308 Trivers 2009), and could be part of the reason why homosporous pteridophytes have relatively
- 309 stable chromosome structures (Clark et al. 2016; Wagner and Wagner 1979; Bomfleur,
- 310 McLoughlin, and Vajda 2014), uniform chromosome sizes (Clark et al. 2016; Wagner and
- 311 Wagner 1979; Manton 1950; Nakazato et al. 2008), and relatively slow rates of chromosome
- number evolution (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
- 315 material in each chromosome (Liu et al. 2019), but also may come from the lack of meiotic drive 316 influencing chromosome morphological change (Plačková et al. 2024). Despite the extensive
- influencing chromosome morphological change (Plačková et al. 2024). Despite the extensive
 research on pteridophyte cytology (e.g., Manton 1950; Klekowski and Baker 1966), limited
- 318 information has been published on their chromosome morphology. We surveyed the literature for
- 319 pteridophytes chromosome morphology data, finding only ten studies describing eleven taxa and
- 320 a broad range of chromosome morphologies (Table 2). Further studies on homosporous and
- 321 heterosporous pteridophyte chromosome morphology and karyotype structure, ideally in a
- 322 phylogenetic context, are needed to understand what processes are influencing these traits.
- 323

324 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 (Fishman and Willis
- 2005; Sandler and Novitski 1957; Pardo-Manuel de Villena and Sapienza 2001b). One example
- 327 of post-meiotic drive is spore drive, which results from competition between spores containing
- 328 different haplotypes, often called killer and alternative haplotypes (Lindholm et al. 2016;

Grognet et al. 2014; Harvey et al. 2014; Raju 1994). 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 (Rose and Dassler 2017; Conant 1978), 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

- 341 (Lindholm et al. 2016).
- 342

343 Another example of post-meiotic drive is male drive, which occurs when a drive locus kills sperm with a target locus (e.g., Presgraves, Severance, and Wilkinson 1997; Taylor 1994). 344 Similar to spore drive, one genotype will kill an alternate genotype regardless of the fitness 345 346 consequences (Lindholm et al. 2016; Rice 2013). There is limited research on fern sperm 347 competition, but observational studies have described competition in the archegonium before 348 sperm meet the egg (Lopez-Smith and Renzaglia 2008). Further work is needed to understand competition and potential male drive among sperm of homosporous plants, as well as in other 349 350 flagellate plants.

351

352 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 353 homosporous plants by removing certain genotypes but likely does not shape the genome in the 354 same ways as true meiotic drive. Several homosporous pteridophyte genome assemblies are now 355 356 published (Marchant et al. 2022; Li et al. 2024; Yu et al. 2023; Fang et al. 2021; Huang et al. 357 2022), 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 358 synteny over millions of years (Huang et al. 2022; Li et al. 2024); the same is true for hornworts 359 (Schafran et al. 2025). An exception is *Ceratopteris richardii*, which has a rate of fractionation 360 rivaling many angiosperm species (Marchant et al. 2022; Nakazato et al. 2006), but the 361 mechanisms causing this rapid diploidization are unclear. Broad patterns in homosporous 362 pteridophytes suggest that, while genic diploidization and fractionation occur, in the apparent 363 364 absence of meiotic drive there are limited mechanisms or forces causing cytological restructuring 365 of the genome following WGD. 366

- 367 Conclusions and future work
- 368

- 370 genome architecture among land plants, specifically influencing chromosome number and
- 371 structure of heterosporous plants. Other types of TRD or drive can occur in homosporous
- 372 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; Kinosian, Rowe, and Wolf 2022; Finseth 2023; Lindholm et al. 2016).
- **376** Pteridophytes are an ideal system to study the effects of drive on genome structure, as there are
- 377 sister lineages with asymmetric and symmetric meiosis in both ferns and lycophytes. Here we
- 378 discuss a few ways to investigate how drive or other types of transmission ratio distortion might
- affect homosporous vascular land plant genomes.
- 380

381 Finding natural examples of meiotic drive can be challenging, as a driving locus is often rapidly 382 fixed in a population, and therefore undetectable. However, these can re-emerge following a hvbridization event, or even be part of hybrid sterility (Lindholm et al. 2016). An example of this 383 is in yellow monkeyflowers, where hybrids between *Mimulus guttatus* and *M. nasutus* exhibit 384 385 strong female meiotic drive and the driving locus is nearly fixed in only a handful of generations (Fishman and Willis 2005; Fishman et al. 2001; Finseth, Nelson, and Fishman 2021). Testing for 386 387 such cryptic drive systems could be a potential avenue of research in pteridophytes, specifically in those with known hybrid sterility. Pteridology has a rich history of work on hybrid species, 388 389 including studies on the model system Ceratopteris (e.g., Hickok and Klekowski 1974; Hickok 1973). Genomic resources available for *Ceratopteris* (Marchant et al. 2022; Nakazato et al. 390 391 2006) would make testing for drive in a hybrid mapping population feasible (e.g., Fishman and 392 Willis 2005). Even without genomic resources, detecting TRD in pteridophytes is possible by 393 genotyping gametophyte progeny arrays to determine meiotic product ratios (described in Kinosian et al. 2022). An important step for understanding meiotic drive and TRD in 394 395 pteridophytes will be to sequence and assemble centromere diversity, ideally among closely related species. Cytology suggests that ferns have distinct monocentric chromosomes, rather than 396 397 dispersed holocentric chromosomes (e.g., Manton, 1950); however, no published fern genomes have assembled centromeres. Interestingly, bryophyte taxa typically have monocentric 398 399 chromosomes, but have small units of euchromatic and heterochromatin dispersed along a chromosome (Schafran et al. 2025; Hisanaga et al. 2023), rather than the large regions of 400 401 heterochromatin which characterize centromeres in monocentric angiosperms (Guerra 2000). Considering that fern genes are more or less evenly distributed along chromosomes (Marchant et 402 403 al. 2022; Rabinowicz et al. 2005; Wolf et al. 2015), their chromatin landscapes may look more 404 similar to bryophytes than to angiosperms. 405

406 Consistent with the absence of meiotic drive, past studies have shown that homosporous ferns

- 407 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.
- 409 2015), perhaps because of a lack of meiotic drive selecting for genome downsizing or,

alternatively, pseudogenization caused by transposons in ferns (Lisch 2013). Long terminal
repeat retrotransposon (LTR-RT) have a high birth but low death rate in homosporous ferns, with

- 412 the inverse occurring in heterosporous lycophytes (Yu et al. 2023; Wang et al. 2021). Older
- 413 LTR-RTs are associated with larger haploid nuclear genome size in pteridophytes (Baniaga and
- 414 Barker 2019). The exact role of transposons in unclear in pteridophytes, but more research is
- 415 warranted, especially because transposons are known to be associated with meiotic drive
- 416 (Eickbush, Young, and Zanders 2019; Vogan et al. 2021). In addition, the expansion of different
- 417 gene families as well as the rate of meiosis gene evolution is greater in heterosporous
- 418 angiosperms than in homosporous pteridophytes (Li et al. 2024; Dhakal, Harkess, and Wolf
- 419 2025), suggesting that different selection pressures, perhaps one of those being meiotic drive, are
- 420 acting on meiosis in heterosporous lineages. It will also be important to look at TRD and sex
- 421 chromosome drive in bryophytes. In particular, mosses have dynamic genomes (Fujiwara et al.
- 422 2025; Patel, Budke, and Bainard 2025) that may be affected by sex chromosome drive (e.g.,
- 423 (Úbeda, Patten, and Wild 2015). Segregation distortion of loci and biased sex ratios have been
- 424 detected in the moss *Ceratodon* (McDaniel, Willis, and Shaw 2007). While our understanding of
- 425 bryophyte genome evolution is improving (Schafran et al. 2025; Fujiwara et al. 2025; Patel,
- 426 Budke, and Bainard 2025), TRD has not been explored in hornworts and liverworts.
- 427
- 428 A post-meiotic TRD mechanism that could occur in homosporous plants is spore drive. In fungi,
- 429 spores are affected by drive when a certain genotype kills an alternative genotype. Considering
- 430 the large number of spores produced within a sporangium by both eusporangiate and
- 431 leptosporangiate lineages, the ingredients for spore drive are present. DNA expression studies
- 432 have identified spore drive systems, revealing that the driving element can lethally target specific
- 433 genes (Urquhart and Gardiner 2023). A consistent reduction in spore number per sporangium in
- 434 leptosporangiate ferns has been observed (Barker and James Hickey 2006; A. L. Grusz, Pers.
- 435 Ob.), suggesting something may be happening during sporogenesis to reduce spore number.
- 436 Additional work counting spores could better characterize this pattern. If more concrete patterns
- 437 of spore reduction are observed and connected to a spore drive system, the model fern
- 438 *Ceratopteris* could help understand the genetic mechanisms for homosporous spore drive.
- 439

440 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 441 drive in Carnivora: smaller populations had greater variation in chromosome number than large 442 populations (Jonika et al. 2024). Ferns appear to have generally high gene flow among 443 populations (Pelosi and Sessa 2021), which could partially explain their consistently high stable 444 chromosome numbers. Another hypothesis for the evolution of chromosome number is the 445 minimum interaction theory, which postulates smaller chromosomes are selected to reduce 446 reciprocal translocation; consequently, chromosome number will increase as more chromosomes 447 are needed to contain the genome (Imai et al. 1986). Ferns are the only lineage with a positive 448 449 correlation between genome size and chromosome number (Nakazato et al. 2008; Bainard et al. 2011; Clark et al. 2016; Fujiwara et al. 2021; Kinosian, Rowe, and Wolf 2022). It has been 450

- 451 suggested that fern chromosome size is limited by the amount of genetic material per
- 452 chromosome (Liu et al. 2019), however, the minimum interaction theory could explain the
- 453 mechanism behind this limitation.
- 454

455 Exploring ecology and natural history in homosporous plants could be another avenue to

- 456 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 lifecycles, compared to those with longer life cycles (Cavalier-Smith 2005; Smith and Donoghue
- 458 cycles, compared to those with longer life cycles (Cavalier-Smith 2005; Smith and Donoghue459 2008). There is also a correlation between vascular structure (i.e., the presence or absence of
- 460 tracheids) and genome size (Cavalier-Smith 2005). Heterosporous ferns (Salviniales) and
- 461 *Selaginella* have tracheids and small genome sizes, but gymnosperms and the heterosporous
- 462 lycophyte *Isoëtes* lack tracheids and have large genomes (Leitch et al. 2005; Cavalier-Smith
- 463 1978). Comparing life history and genome traits in a phylogenetic context could help tease apart
- 464 complex interactions between these variables (Soltis and Soltis 1990; Pelosi and Sessa 2021).465
- 466 The proliferation of a selfish element is key in female meiotic drive and TRD systems, yet the
- 467 mechanisms by which this occurs are incredibly varied (Saupe and Johannesson 2022; Lindholm
- 468 et al. 2016; Burt and Trivers 2009). Female meiotic drive is potentially a mechanism behind the
- 469 rapid diploidization and downsizing following WGD in heterosporous plants. TRD in
- 470 pteridophytes and bryophytes is probably distinct from true meiotic drive in angiosperms, fungi,
- 471 or animals. The absence of meiotic drive would help explain large and stable genomes, uniform
- 472 chromosome sizes, relatively static chromosome numbers following WGDs, as well as the
- 473 current challenges with assembling centromeres in homosporous fern and lycophyte genomes.
- 474 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
- 476 lineage.

477

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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	(Pellicer and Leitch 2020; Klekowski and Baker 1966; McKibben, Finch, and Barker 2024; One Thousand Plant Transcriptomes Initiative 2019; Z. Li et al. 2024)
Gymnosperm	15.51	11.69	1.63	Yes	Centromere, chromosome, female meiotic drive	(One Thousand Plant Transcriptomes Initiative 2019; Pellicer and Leitch 2020; A. Rice et al. 2015; Z. Li et al. 2024)
Heterosporous pteridophyte (Salviniales, Isoetales, Selaginellales)	1.01	13.62	2.4	Yes	Centromere, chromosome, female meiotic drive	(Pellicer and Leitch 2020; Z. Li et al. 2024; Klekowski and Baker 1966)
Homosporous pteridophyte	15.13	57.5	2.82	No	Spore drive, other non- meiotic TRD	(Pellicer and Leitch 2020; Z. Li et al. 2024; Klekowski and Baker 1966)
Bryophytes	0.72	10.23	2.28	No	Spore drive, other non- meiotic TRD, sex chromosome drive	(Pellicer and Leitch 2020; Z. Li et al. 2024; Klekowski and Baker 1966) N. Patel, et al. 2025)

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

- **Table 2.** Published descriptions of chromosome morphology for homosporous and heterosporous
 pteridophytes.

Taxon	Reproductive mode	Chromosome morphology	Citation
Claytosmunda claytoniana	homosporous	acrocentric, very few metacentric	(Seizi Tatuno and Yoshida 1967)
Osmundastrum cinnamomeum	homosporous	acrocentric, very few metacentric	(Seizi Tatuno and Yoshida 1967)
Plenasium banksiaefolium	homosporous	acrocentric, very few metacentric	(Seizi Tatuno and Yoshida 1967)
Hymenophyllum barbatum	homosporous	acrocentric, some metacentric	(Seizi Tatuno and Takei 1969)
Asplenium incisum	homosporous	acrocentric, very few metacentric	(S. Tatuno and Kawakami 1969)
Pteris	homosporous	acrocentric	(Kawakami 1971)
Regnellidium diphyllum (Salviniales)	heterosporous	Submetacentric and subtelocentric	(Kuriachan 1994)
Selaginella (Selaginellales)	heterosporous	Metacentric, telocentric	(Adriana Buarque Marcon, Barros, and Guerra 2005; Takamiya 1993)
Danaea	homosporous	submetacentric	(Benko-Iseppon, Rodrigues, and da Fonsêca 2000)
Acrostichum	homosporous	subtelocentric	(Adriana B. Marcon, Barros, and Guerra 2003)
Doryopteris triphylla	homosporous	telocentric	(Neira et al. 2017)

Figures

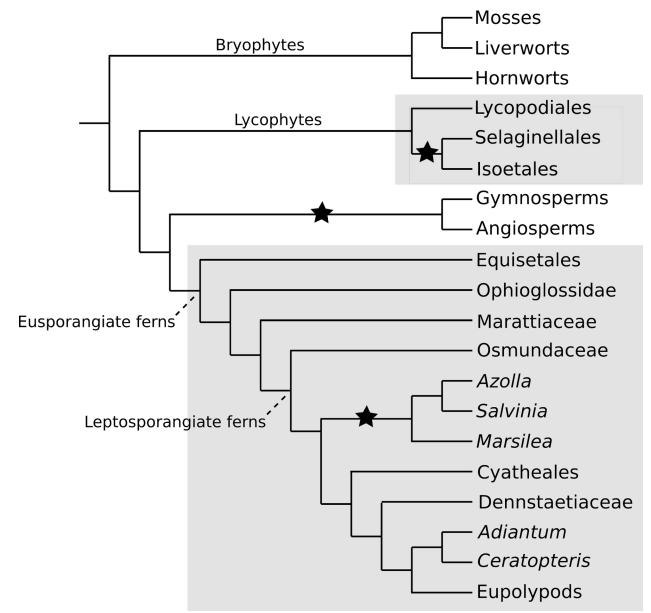


Figure 1. Cartoon phylogeny of land plants, including notable groups discussed in this paper but
not all major lineages. Grey boxes distinguish pteridophytes, a paraphyletic group comprising
lycophytes (above) and ferns (below). Black stars distinguish the three extant heterosporous
plant clades. Modified from (PPG I 2016) and (Patel, Budke, and Bainard 2025).

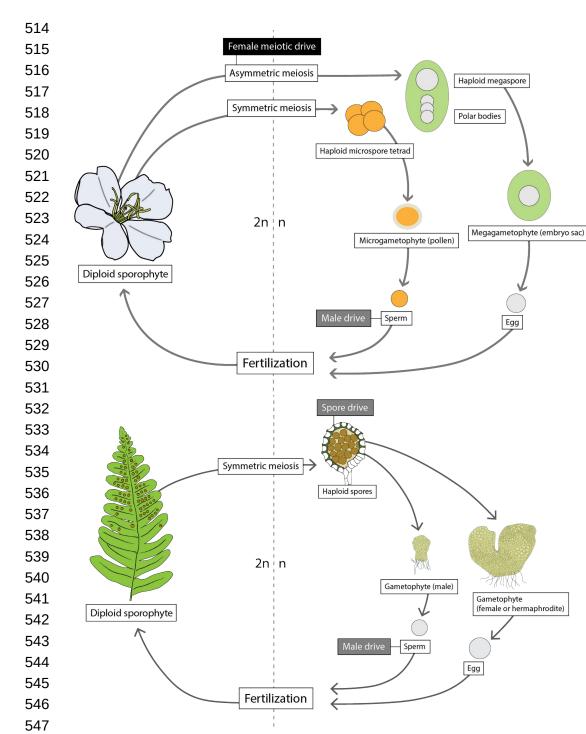


Figure 2. Life cycles of heterosporous (top) and homosporous (bottom) land plants. Locations of
female meiotic drive and other types of transmission ratio distortion are denoted in black and
gray boxes, respectively.

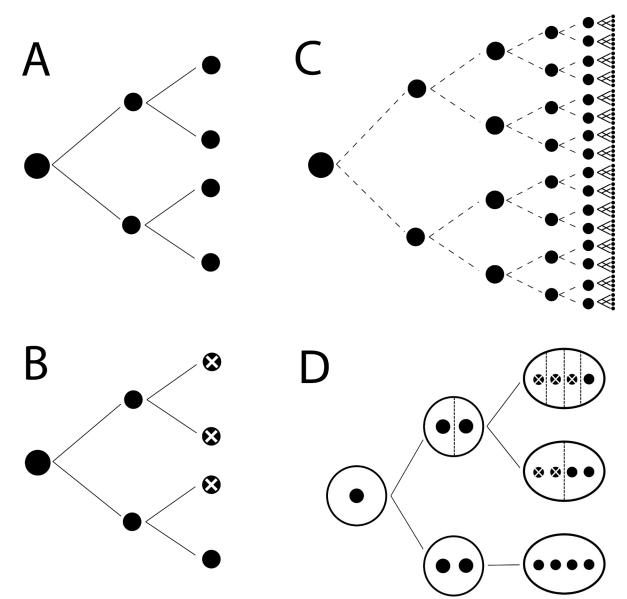


Figure 3. Examples of different types of meiosis. In each panel, the far left circle depicts the 554 555 sporocyte, and succeeding divisions proceeding to the right. Solid lines indicate meiosis and dotted lines indicate mitosis. A) Heterosporous female meiosis, where only one meiotic product 556 survives and the remaining three polar bodies die (can be either top or bottom, resulting from the 557 cell on the chalazal or micropylar end surviving, see Fig. 4A). B) Heterosporous male meiosis, 558 559 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. D) Types 560 561 of angiosperm megasporogenesis: monosporoic (top), bisporic (middle), and tetrasporic (bottom), modified from (Haig 1990, 2020); meiosis proceeds similarly to (A), but differing 562 563 numbers of nuclei survive due to variable cytokinesis in meiosis II. 564

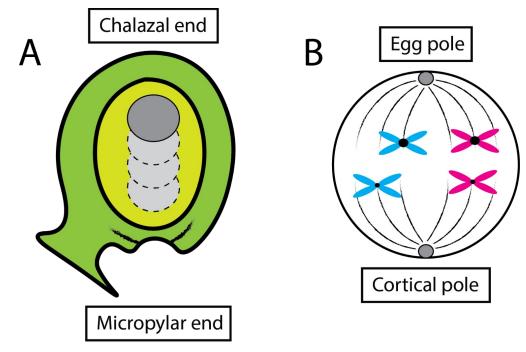




Figure 4. A) Organization of megasporogenesis in a heterosporous ovule, the cell on the chalazal
end often survives, but alternatively the cell on the micropylar end can become the egg. B)
Centromere drive, where larger centromeres are preferentially transmitted to the egg pole in
meiosis I (anaphase I is depicted); the egg pole is often associated with the chalazal end of the
ovule (see panel A), and the cortical side the micropylar end.

- 591
- 592 Transmission ratio distortion non-random inheritance of parental alleles in the offspring
 593 population, occuring before, during, or after meiosis.
- 594 **Meiotic drive** non-random inheritance of alleles, chromosomes, or other genome structures
- 595 specifically during asymmetrical meiosis.
- 596 Spore drive observed in ascomycete fungi where spores are packages together in an ascus, a
- 597 spore with the "killer" haplotype will kill spores with the alternative haplotype, particularly in 598 when spores are in close proximity.
- 599 Male drive biased transmission that occurs during or after male meiosis; typically involves an
- 600 allele that kills certain sperm cells.
- 601 Sporogenesis the generation of a spore cell by meiosis (in heterosporous plants) or mitosis (in602 homosporous plants)
- 603 **Megasporogenesis** creation of spores that will develop into a female gametophyte
- 604 (megagametophyte)
- 605 Microsporogenesis creation of spores that will develop into a male gametophyte
- 606 (megagametophyte)
- 607 Homosporous an organism that produces one type of spore. This spore develops into a
- 608 gametophyte which can produce male and/or female gametes.
- 609 Heterosporous an organism that produces two types of spores. These are the mega- and
- 610 microspores, which develop into gametophytes capable of producing eggs or sperm, respectively.
- 611 **Pteridophyte** a plant classified as a fern or lycophyte. This group is paraphyletic but is
- 612 functionally important because all taxa are seedless vascular plants.
- 613 **Leptosporangiate fern** ferns that produce sporangia with a long stalk that arise from a single
- epidermal cell. Their spore production proceeds from a single spore mother cell through four
- rounds of mitosis and one round of meiosis to form 64 spores.
- 616 **Eusporangiate fern** ferns that produce sporangia with a short or no stalk that arise from
- several epidermal cells. Hundreds to thousands of spores are produced in each sporangium viameiosis.
- 619 **Metacentric** a chromosome with the centromere positioned in the middle, with both arms
- 620 being the same length.
- 621 **Acrocentric** a chromosome with the centromere close to one end, with one arm being much
- 622 longer than the other.
- **Telocentric** a chromosome where the centromere is at one end, causing there to be effectivelyonly one arm rather than two.
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