## 1 The X chromosome is a potential polarising signal for asymmetric cell

## 2 divisions in meiotic cells of a nematode

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# 12 ABSTRACT

The unequal partition of molecules and organelles during cell division results in daughter cells with different fates. Asymmetric cell divisions have been best characterised in systems in which extrinsic signals polarise the mother cell during

16 cell division. However, the mechanisms of asymmetric cell division mediated by

17 intrinsic signals, and the nature of these signals, are mostly unknown. Here we

18 report an asymmetric cell division in the nematode Auanema rhodensis that may be

19 cued by the X chromosome. In the wildtype XO male, the spermatocyte divides

20 asymmetrically to generate X-bearing spermatids that inherit components necessary

21 for sperm viability, and nullo-spermatids that inherits components to be discarded.

22 We found that in XX mutant pseudomales, sperm components co-segregate with the

23 X chromosome, supporting the hypothesis that the X chromosome is employed as a

24 polarising signal for partitioning essential cytoplasmic components for sperm

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### 35 INTRODUCTION

36 An asymmetric cell division (ACD) is a division that results in sister cells with 37 different sizes, morphology, chemical composition, or gene expression. ACDs are 38 present in both unicellular and multicellular organisms, are crucial for embryonic 39 development, cellular differentiation, and production of cell diversity (Gönczy, 2008; 40 Venkei and Yamashita, 2018), whereas their dysregulation may result in 41 tumorigenesis (Knoblich, 2010; Neumuller and Knoblich, 2009). 42 43 The polarity of a dividing cell is established by extrinsic or intrinsic factors. The best-44 known examples are the cases in which extrinsic signals provided by neighbouring 45 cells polarise the dividing cell (Fuller and Spradling, 2007). Relatively little is known 46 about intrinsic mechanisms controlling the polarity of an asymmetric dividing cell 47 (Freisinger et al., 2013). We have previously described a cell division that undergoes 48 asymmetric segregation that is likely to be guided by internal mechanisms. This 49 division occurs during the final division of the male spermatogenesis in the nematode 50 Auanema rhodensis (aka, Rhabditis sp. SB347) (Shakes et al., 2011; Winter et al., 51 2017). 52

53 Spermatogenesis is very well characterized in the nematode *Caenorhabditis* 

*elegans*. During the maturation of the spermatids, non-essential cytoplasmic

55 components are extruded into vesicles named residual bodies (Fig. 1A). Thus, the

56 final product of an XO spermatocyte division and differentiation is the formation of

57 four spermatids, two spermatids with no X chromosomes (nullo-X spermatids) and

58 two spermatids with one X chromosome each (X-bearing spermatids) (Fig. 1A).

59 Unexpectedly, spermatogenesis in *A. rhodensis* results in only two viable gametes,

60 each containing either one X chromosome (XO male, Fig. 1B) or two X

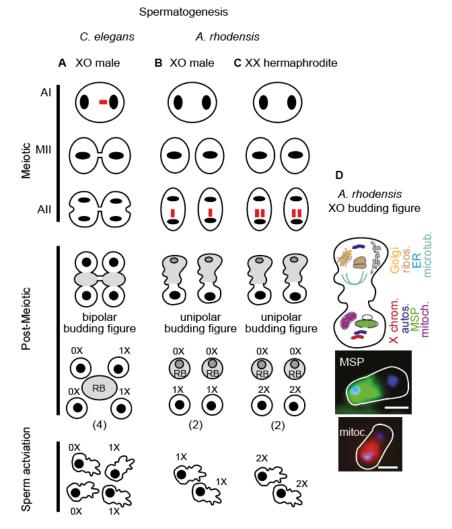
61 chromosomes (Shakes et al., 2011; Tandonnet et al., 2018) (XX hermaphrodite, Fig.

62 1C). In A. rhodensis the non-essential sperm components are discarded into the

63 nullo-X spermatids because of asymmetric partitioning of the cytoplasm during

anaphase II (Fig. 1D) (Shakes et al., 2011; Winter et al., 2017). Thus, the nullo-X

65 spermatids become cellular residual bodies.





68 Figure 1. Comparison of spermatogenesis between C. elegans and A. rhodensis. In C. 69 elegans males (A), the unpaired X chromosome (in red) lags during anaphase I, leading to 70 the formation of two secondary spermatocytes that divide symmetrically to generate either 71 two gametes with 1X or two gametes containing only autosomes (0X). A central residual 72 body (RB) is formed in the postmeiotic phase. In A. rhodensis males (B), the sister 73 chromatids of the X chromosome separate prematurely in meiosis I and the X chromosome 74 lags during anaphase II. High rates of non-disjunction in A. rhodensis hermaphrodite 75 spermatogenesis leads to the formation of 2X sperm (C). Only two functional gametes are 76 formed during spermatogenesis of A. rhodensis (**B**, **C**), with the nullo-X sperm acting as a 77 residual body. Cytoplasmic components partition during the second meiotic division (D), 78 whereby specific organelles and proteins partition in opposite directions. Mitochondria and 79 the sperm cytoskeleton protein Major Sperm Protein (MSP) partition with the X chromosome 80 to one of the poles of the cell, whereas ribosomes, Golgi complex, and microtubules migrate 81 to the opposite side of the dividing cell, which has only autosomes (Winter et al., 2017).

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- 83 We hypothesized that the *A. rhodensis* X chromosome acts as a polarizing signal,
- 84 guiding the partitioning of sperm-specific cytoplasmic contents to one of the poles of
- 85 the dividing secondary spermatocyte. As a result of this ACD during
- 86 spermatogenesis, A. rhodensis males and other species of the same genus produce
- mostly X-bearing sperm (Chaudhuri et al., 2015; Shakes et al., 2011; Winter et al.,
- 88 2017). Thus, crosses between males and females result mostly in XX progeny,
- 89 which can be either female or hermaphrodite (Chaudhuri et al., 2011a).
- 90 Hermaphrodites, which produce nullo-X oocytes and XX-sperm, produce mostly XX
- 91 self-progeny. To determine if the X chromosome acts as a polarizing signal, we
- 92 isolated a sex determination mutant that is phenotypic male but that harbours two X
- 93 chromosomes. We show that sperm components co-segregate with the X
- 94 chromosome, further supporting that this chromosome acts as a polarizing signal.
- 95

# 96 MATERIALS AND METHODS

# 97 Nematodes strains and cultures

- 98 The Auanema rhodensis inbred strains APS4 and APS6 (Tandonnet et al., 2018)
- 99 were maintained according to standard conditions for *C. elegans*, at 20 °C
- 100 (Stiernagle, 2006). Nematodes were cultured on NGM plates seeded with the
- 101 Escherichia coli strain OP50 or with the E. coli streptomycin-resistant strain OP50-1.
- 102

# 103 Mutagenesis

- 104 A. rhodensis APS4 was mutagenized with the chemical mutagen ethyl
- 105 methanesulfonate (EMS), as previously described (Chaudhuri et al., 2011b; Pires-
- 106 daSilva and Sommer, 2004). To screen for a masculinizing phenotype (XX
- 107 pseudomales), we isolated 521 F1 hermaphrodites derived from mutagenized P0s
- and transferred them to single plates to let them self-fertilize. To simplify screening
- 109 for mutants that generate high rates of (pseudo)males, we screened for F2s derived
- 110 from 3-day old hermaphrodites. We adopted this procedure because hermaphrodites
- 111 of this age tend to generate less XO self-offspring (~3%) than younger
- 112 hermaphrodites (~8%) (Chaudhuri et al., 2015). From plates in which potential
- 113 pseudomale mutants were found (containing ~25% of phenotypic males), 10-15
- 114 sister hermaphrodites were isolated to single plates to maintain the mutations as a
- 115 heterozygous strain. Heterozygous hermaphrodites were selected based on the

- 116 production of excess male progeny, which is consistent with the anticipated
- 117 production of 25% XX male progeny
- 118

### 119 A. rhodensis masculiniser (Arh-mas-1) crosses

- 120 The A. rhodensis masculiniser was named Arh-mas-1(brz-3), following the
- 121 nomenclature described in Wormbase (<u>www.wormbase.org</u>). Arh-mas-1(brz-3) was
- 122 backcrossed with the wildtype APS4 for three generations to remove background
- 123 mutations generated during the mutagenesis.
- 124
- 125 To ascertain that the pseudo male has two X chromosomes, we crossed it to the
- 126 polymorphic strain APS6. Arh-mas-1 crosses with APS6 females were performed for
- 127 ~24 h at 20 °C. To isolate females for crosses, we left a 1-day old hermaphrodite to
- 128 lay eggs, which were picked to individual wells of a 24-well plate. After 72 hours,
- 129 those eggs develop into males, females or hermaphrodites. Females were
- 130 distinguished from males by their tail morphology, and from hermaphrodites by their
- 131 inability to self-fertilize (Kanzaki et al., 2017).
- 132

#### 133 Single nematode genotyping

- 134 The X chromosome genetic markers (markers 9686 and 12469) generated for A.
- 135 *rhodensis* (Tandonnet et al., 2019), together with the primer sequences, restriction
- 136 enzymes and fragment sizes are detailed in
- 137 https://data.mendeley.com/datasets/63d7rrrx28/3#file-16ff094d-6c74-478a-a3f5-
- 138 <u>8878e89fd72f</u>.
- 139

#### 140 Antibody staining

- 141 To detect the ER, we used an antibody against cytochrome P450 (CYPP33-E1). For
- 142 mitochondria, we used an antibody against the beta-subunit of ATP synthase.
- 143

## 144 **RESULTS**

- 145 Arh-mas-1 has a male phenotype and XX karyotype
- 146 We performed chemical mutagenesis in *A. rhodensis* to screen for a masculinizing
- 147 sex determination mutant. This mutant would allow us to determine whether the X
- 148 chromosome acts as a signal to partition the cytoplasm in dividing spermatocytes. If
- 149 this hypothesis is correct, the expectation was that the two homologous X

150 chromosomes segregate to opposite poles in the first and second meiotic divisions, 151 following a Mendelian pattern (Fig. 2A, Model 1), to generate viable 1X spermatids. 152 Although A. rhodensis XX females follow the canonical meiosis, XX hermaphrodites 153 do not: the X chromosomes undergo premature chromatid separation in the first 154 meiotic division, forming viable 2X spermatids and non-viable nullo-X spermatids 155 (Tandonnet et al., 2018). Thus, similar to hermaphrodites, pseudomales may follow 156 the pattern found in hermaphrodite spermatogenesis (Fig. 2A, Model 2). 157 Alternatively, those X chromatids may segregate equally to each pole to make viable 158 1X sperm. A third possibility is that the two X chromosomes segregate to one pole in 159 the first meiotic division, generating spermatocytes that either make only nullo-X 160 spermatids or 0-4X spermatids. (Fig. 2A, Model 3). The fourth possibility is a mixture 161 of Models 2 and 3. The hypothesis of the X chromosome acting as a polarising 162 signal for cytoplasmic partitioning would be falsified in case the cellular components 163 important for post-meiotic sperm do not co-segregate with the X chromosomes in 164 anaphase II.

165

166 We found an XX A. rhodensis sex determination mutant (Arh-mas-1) with a male

167 morphological phenotype that is almost indistinguishable from XO wildtype males

168 (Fig. 2B-E). Adult Arh-mas-1 XX pseudomales that are three days or older display a

169 gut pigmentation pattern distinct from XO males (Fig. 2B), which we used as a

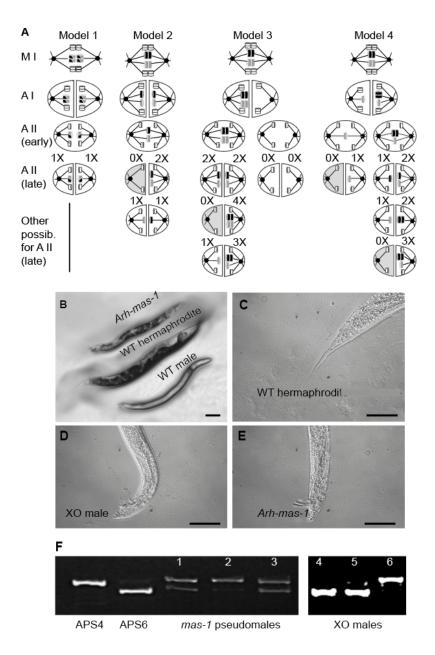
170 marker to distinguish between the two karyotypes. Arh-mas-1 pseudomales do not

171 show signs of partial feminization, as encountered in similar mutants in other

172 nematode species (Hill et al., 2006; Hodgkin, 1979; Pires-daSilva and Sommer,

173 2004). Arh-mas-1 pseudomales display mating behaviours and produce viable

174 offspring when crossing with females (see below).





177

178 Figure 2. Arh-mas-1 masculinizes XX animals. (A) Models for possible meiotic events in 179 XX pseudomales. Homologous X chromosomes are in grey and black, and autosomes are in 180 white. In the models, nullo-X sperm are discarded as residual bodies (grey cytoplasm) when 181 the sister cell has an X chromosome(s). (B) The Arh-mas-1 pseudomale has a gut dark 182 pigmentation pattern distinct from the wildtype male. Bar, 10 µm. The tail of the 183 hermaphrodite (C) is long and slender, whereas the male (D) and Arh-mas-1 (E) are blunt 184 and with spicules. Bar, 50 µm. (F) Genotyping of the X chromosome (chromosome marker 185 9686) in animals derived from crosses between pseudomales (APS4 background) and 186 females (APS6).

#### 187

188 To confirm the XX karyotype of pseudomales, we crossed *Arh-mas-1* pseudomales

189 (in APS4 background) with wildtype females (APS6 background). Using markers for

190 the X chromosome, we genotyped pseudomales derived from self-progeny of F1

- 191 hybrid hermaphrodites. We found that *Arh-mas-1* pseudomales were heterozygous
- 192 for the X chromosome markers (Fig. 2F, samples 1-3), confirming that this is a strain
- 193 with an XX karyotype.
- 194

## 195 Arh-mas-1 XX pseudomales produce 0-4X sperm

- 196 In crosses between XX pseudomales and females, most of the F1 offspring is
- 197 hermaphrodite or female (Fig. 3A), indicating that most of the Arh-mas-1
- 198 spermatogenesis events generate viable 1X sperm. Cytological studies indicate that
- there is high variation in the way the X chromosomes segregate (Fig. 3B). In contrast
- to *C. elegans* (Fig. 1B), we never observed a lagging chromosome in anaphase I of
- 201 A. rhodensis wildtype XO males. Instead, the unpaired X chromosome lags in
- anaphase II (Fig. 1B, Fig. 3B)(Shakes et al., 2011). Although the majority (66%,
- 203 N=224) of the primary spermatocytes in XX pseudomales (Arh-mas-1) segregate
- 204 DNA symmetrically, we also observed primary spermatocytes that segregate
- unequal amounts of DNA to the daughter cells (33%, N= 224). These occasional
- 206 asymmetric cell divisions are supported by the observation of lagging X
- 207 chromosomes during anaphase I (Fig. 3B). The different sizes of lagging
- 208 chromosome masses could be due to (1) segregation of homologous X
- 209 chromosomes to the same side, and (2) the premature separation of X chromatids in
- 210 anaphase I (see models in Fig. 2A).
- 211
- 212 During anaphase II in spermatocytes of XX pseudomales, lagging chromosomes are
- also segregated symmetrically (59%, N= 54) (Fig. 3B) or asymmetrically (35%, N=
- 54)(Fig. 3B). Chromosome-lagging has been attributed to the attachment of
- 215 microtubules to both sides of an unpaired chromosome, delaying its segregation to
- one of the poles (Fabig et al., 2020).
- 217
- About 1% of the F1s from XX pseudomales and females were dumpy animals (Fig.
- 3A), possibly as the result of the fertilization of XX-sperm with an X-bearing oocyte.
- 220 These dumpy animals are unlikely to be the result of background mutations because

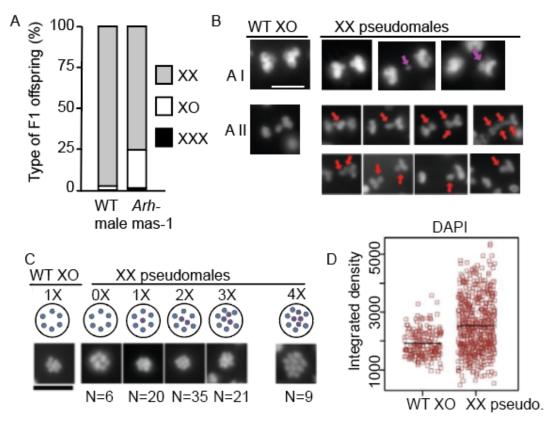
the strain was backcrossed three times and because we have not observed dumpy
 phenotype offspring from hermaphrodites harbouring the *Arh-mas-1* mutation. In *C. elegans*, XXX animals have a dumpy phenotype because of dosage compensation

defects (Hodgkin, 1979; Vargas et al., 2017).

225

226 About a quarter of the offspring of a cross between a XX Arh-mas-1 pseudomale and 227 a XX female is XO male (Fig. 3A). Those F1 XO males are likely the result of the 228 fertilization between an X-bearing sperm and a nullo-oocyte. The formation of a 229 nullo-oocyte in females is rare, but it occurs in ~3% of the meiosis (Shakes et al., 230 2011). Confirming this hypothesis, we found that 23% (N= 57) of the tested F1 males 231 harbour the paternal X chromosome (Fig. 2F, samples 4, 5). An alternative scenario 232 for the formation of XO offspring is the fertilization of an X-bearing oocyte by a nullo-233 X sperm. This seems to have been the case for the remaining 77% (N=57) of the 234 cases (Fig. 2F, sample 6), in which F1 males inherited the maternal X chromosome. 235 236 To determine more directly the number of chromosomes in sperm, we examined

237 spermatocytes stained with the DNA-staining dyes DAPI or Hoechst 33342. In 238 metaphase plates, in which the number of chromosomes can be quantified, we 239 confirmed the previous observation that A. rhodensis XO males have 6 autosomes 240 and 1X chromosome (Fig. 3C) (Shakes et al., 2011). In XX pseudomales we observed cells only with autosomes (0X), or with X chromosomes that varied in 241 242 number between 1-4 (Fig. 4C). To identify mature, functional sperm, we used the 243 cytoskeleton protein Major Sperm Protein (MSP) as a marker. In those cells, the 244 range of DNA intensity and size of those sperm was broader than in wild-type XO 245 males (Fig. 3D), confirming the variability in the number of chromosomes in XX 246 pseudomales. These variations reflect models 1-4 (Fig. 2A) and are useful to test if 247 the X chromosome guides the partitioning of cytoplasmic components (see below).



249

250 Figure 3. Types of F1 offspring after crossing *Arh-mas-1* pseudomales with females.

(A) Bar graph representing the proportion of XX animals (females or hermaphrodites), XO
 (males) and XXX (dumpy). 19 crosses were performed between APS4 wildtype males and

253 females (N= 5629 offspring) and 23 crosses were performed between Arh-mas-1

254 pseudomales and APS4 females (N= 3135 offspring). (B) Meiosis in wildtype A. rhodensis

255 XO males and XX pseudomales. The pink arrow is a lagging chromosome (stained with

- 256 Hoechst) in anaphase I (A I) and red arrows indicate lagging chromosomes (stained with
- 257 DAPI) in anaphase II (A II). Scale bar= 5 µm. (C) A variable number of chromosomes in

258 metaphase plates, stained with DAPI. N= number of spermatocytes observed for each

category, out of a total of 91. (**D**) Quantification of DAPI in MSP-stained sperm.

260

## 261 **Post-meiotic sperm components co-segregate with the X chromosomes**

To examine the pattern of segregation of cytoplasmic components relative to the X chromosome segregation, we performed immunostainings on fixed sperm spreads. We chose cytoplasmic components that are essential for post-meiotic sperm (MSP and mitochondria) and others that are discarded into residual bodies (endoplasmic reticulum, alpha-tubulin and Small Ubiquitin-like Modifier (SUMO)). MSP is a protein contained within organelles called fibrous bodies in spermatocytes and is required in

268 mature sperm for sperm motility (Smith, 2014). SUMO conjugation has been 269 implicated in a variety of cellular processes (Drabikowski et al., 2018), including the 270 regulation of meiotic proteins in *C. elegans* (Davis-Roca et al., 2018; Pelisch et al., 271 2017). Except for mitochondria and SUMO, we previously described the distribution 272 of MSP, tubulin and endoplasmic reticulum (ER) in XO A. rhodensis spermatocytes 273 (Shakes et al., 2011; Winter et al., 2017). 274 275 During the first meiotic division in wildtype XO males, there is symmetric segregation 276 of X chromatids, autosomal homologous chromosomes and all five cytoplasmic 277 components (Fig. S1A-D). In XX pseudomales, we observe the same pattern of 278 bipolar distribution for all markers (Fig. S1A-D). Equal partitioning of cytoplasmic 279 components occurs even when there is asymmetric segregation of chromosomes 280 (63/194 primary spermatocytes), with one pole receiving more DNA than the other. 281 During early anaphase II, when the lagging X is not yet in contact with autosomes, 282 we also observe the equal distribution of cytoplasmic components in XO males and

- 283 XX pseudomales for all markers (N= 407 secondary spermatocytes) (Fig. S1A-D).
- 284

In late anaphase II, when the X chromosome is biased towards one of the poles,

asymmetries start to appear for most of the tested cytoplasmic components (Fig. 4).

287 In wildtype XO animals, microtubules show a dense and elongated shape on the

pole with the X chromosome (Fig. 4A). Similarly, mitochondria and MSP start to

show a polarization towards the side with the X chromosome. The ER, however,

tends to stay in a more central position of the dividing cell (Fig. 4C). SUMO antibody

staining includes labelling of the lagging X chromosome, which is distinctly

asymmetric (Fig. 4D).

293

294 The post-meiotic partitioning stage has the most pronounced asymmetries of

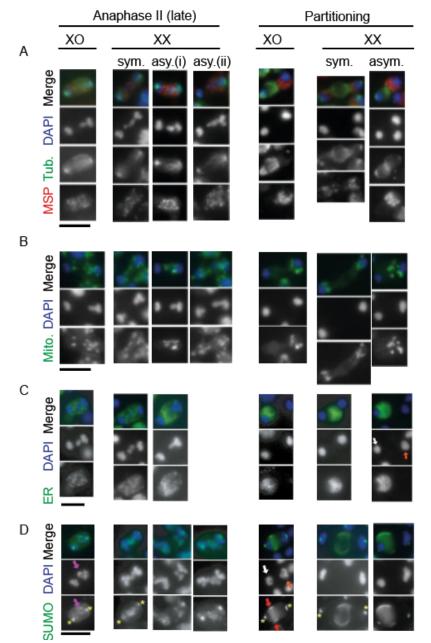
295 cytoplasmic components. By then, all components are distributed in a unipolar

296 pattern, with MSP and mitochondria at the pole with the highest DNA content,

whereas tubulin, ER and SUMO with the opposite pattern (Fig. 4A-D). The pattern of

distribution of SUMO in the cortex of the cell to become the residual body is like the

299 previously described distribution of actin (Winter et al., 2017).





301 Figure 4. Asymmetric distribution of cytoplasmic components. Sperm spreads of XO 302 wildtype males and XX pseudomales stained with antibodies against alpha-tubulin (A), MSP 303 (A), mitochondrial beta-subunit of ATP synthase (B), ER-located cyp-33E1 (cytochrome 304 P450 family) (C) and SUMO (D). In cells dividing asymmetrically (asym.), they may have one 305 (i) or more chromosomes (ii) migrating to opposite poles. The orange and white arrows 306 indicate larger and smaller DNA masses, respectively. The red arrows for the SUMO 307 antibody staining indicate the expanding cortical bands. The yellow asterisks are 308 centrosomes. Scale bar= 5 µm. 309

310 In XX pseudomales, X chromosomes segregate symmetrically in 30% of the cases 311 (N=829), with both poles containing equal amounts of DNA. The remaining cells 312 (70%, N=829), which segregate DNA asymmetrically, may have one or more 313 chromosomes segregating to one of the poles. With no exceptions, we observe that 314 MSP distribution is located around the centre of the symmetrically dividing cells in 315 the late anaphase II and partitioning phase, but it is unipolar with a bias towards the 316 side with more DNA in asymmetrically dividing cells (Fig. 4A). Similarly, the 317 mitochondria distribute equally in cells that divide the chromosomes symmetrically. 318 but with a bias towards the pole with more DNA (Fig. 4B). In instances in which there 319 are lagging chromosomes on both sides, but with unequal amounts of DNA (Fig. 4B), 320 mitochondria partition to both sides but shows a higher prevalence in the pole with 321 more DNA.

322

323 For the components that are discarded into polar bodies in wildtype XO males 324 (tubulin, ER, SUMO), in XX pseudomales they follow the predicted patterns of 325 distributing equally or centrally in cells that divide with equal amounts of DNA. In 326 asymmetric segregation of DNA, these components partition to the pole with less 327 DNA. In some of the cells with symmetric amounts of DNA segregation, tubulin is 328 located centrally in the partitioning phase (Fig. 4A). This is reminiscent of residual 329 body formation in C. elegans (Ward et al., 1983), which forms centrally in this 330 organism but never in wildtype A. rhodensis (Shakes et al., 2011). The ER 331 distribution in XX pseudomales remains at the centre of symmetrically dividing cells 332 in both anaphase II and partitioning phase (Fig. 4C). In cells with asymmetric 333 segregation of chromosomes, the ER distribution concentrates in the pole with less 334 DNA (Fig. 4C). The SUMO distribution in late anaphase II reflects in large part the 335 staining of lagging chromosomes, which can be equally partitioned in cells dividing 336 with equal amounts of DNA or biased towards one of the poles in asymmetrically 337 dividing cells (Fig. 4D). When partitioning, SUMO localization is in the central part of 338 the cortex of symmetrically dividing cells, or in the cortex of the cell with less DNA 339 (Fig. 4D).

340

#### 341 **DISCUSSION**

The initial observation of a few male offspring derived from *Auanema* male and female crosses is the result of a modification in spermatogenesis: males discard

male-generating spermatids (nullo-X sperm) and promote the production of Xbearing sperm (Shakes et al., 2011; Winter et al., 2017). This is a mechanism that
generates biased sex ratios, which provides adaptive advantages in certain
ecological circumstances (Van Goor et al., 2021). Here we provide support for the
hypothesis that the X chromosome acts as a signal for an ACD to generate viable Xbearing sperm cells. The X chromosome would be an example of an intrinsic signal
for polarising cell divisions.

351 Asymmetric segregation of proteins, RNAs and organelles during cell division 352 occurs in prokaryotes and eukaryotes (Horvitz and Herskowitz, 1992; Inaba and 353 Yamashita, 2012; Morrison and Kimble, 2006; Sunchu and Cabernard, 2020). In 354 mitotic cells, biased distribution of cytoplasmic components may result in cells with 355 different cell fate determinants, organelles and protein aggregates. Previously 356 reported examples for intrinsic cues include random events (Broadus and Doe, 357 1997) and co-segregation with organelles that are intrinsically asymmetric (Chen and 358 Yamashita, 2021).

359 This type of cell division found in *Auanema* spermatogenesis, in which 360 residual material is partitioned to a cell that contains chromosomes, is atypical in 361 metazoans (Gorelick et al., 2016). Residual bodies, which are often observed in 362 ACDs during spermiogenesis, are usually devoid of DNA (Gorelick et al., 2016; 363 Zakrzewski et al., 2021). There are a few exceptions, such as in the nematode 364 *Rhabdias*, in which one of the X chromosomes is discarded into a residual body 365 during the hermaphrodite spermatogenesis (Runey et al., 1978). During residual 366 body formation In C. elegans spermatogenesis, motor proteins (SPE-15, myosin VI; 367 NMY-2, non-muscle myosin II)(Hu et al., 2019; Kelleher et al., 2000), and a cargo 368 adaptor protein (GIPC, RGS-GAIP-interacting protein C) (Hu et al., 2019) have been 369 implicated in the correct segregation of organelles. However, the polarisation signal 370 for this asymmetric segregation is unknown.

Contrary to the canonical meiosis, the unpaired X chromosome in *Auanema* males separates its sister chromatids already in the first meiotic division (Shakes et al., 2011; Winter et al., 2017). The result is that each secondary spermatocyte has an X chromosome. Thus, during cell division of these cells, there is potential for the X chromosome to act as a polarising signal. Such a scenario does not occur in *C. elegans* males (Fig. 1), as this nematode follows the canonical meiosis (reviewed in (Chu and Shakes, 2013)): each secondary spermatocyte will either have the X

378 chromosome or not. When the secondary spermatocytes divide, each daughter cell 379 will contain the same chromosomal content (either with an X or without) as the 380 parental cell. Therefore, the polarisation mechanisms for the correct partition of 381 cytoplasmic content for the residual bodies is likely to be different in C. elegans. 382 The mechanism by which the X chromosome may act as a polarising cue in 383 Auanema is still unclear. It is possible, for instance, that a secondary nucleation 384 centre for cytoskeleton proteins, such as non-centrosomal microtubule-organising 385 centres, emanate from the X chromosome (Sanchez and Feldman, 2017). Further 386 genetics studies, as well as electron microscopy and live-cell imaging will help to 387 elucidate the possible mechanisms.

388 389

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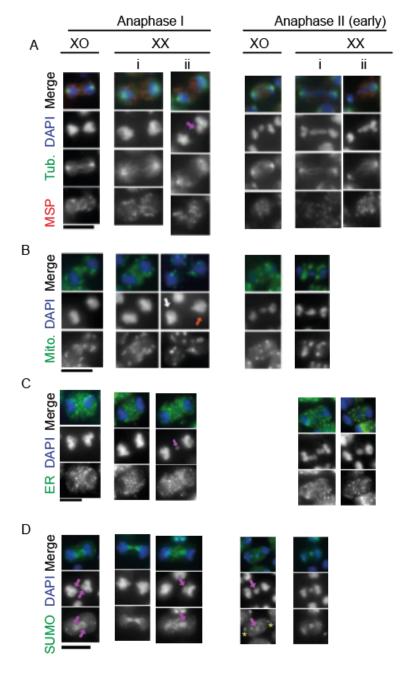
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# 507 Supplemental Figure 1. Cytoplasmic components segregate symmetrically in

508 anaphase I and early anaphase II. In wildtype XO males and XX pseudomales, the

- 509 cytoplasmic components MSP (A), alpha-tubulin (A), mitochondria (B), endoplasmic
- 510 reticulum (ER) (C) and SUMO (D) distribute equally to both poles of the dividing
- 511 spermatocytes. The pink arrow indicates lagging chromosomes, whereas the red and white
- 512 arrows indicate larger and smaller DNA masses, respectively. Scale bar= 5  $\mu$ m.
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