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The chromosomal basis of meiotic acentrosomal spindle assembly and function in oocytes

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Abstract

Several aspects of meiosis are impacted by the absence of centrosomes in oocytes. Here we review four aspects of meiosis I that are significantly affected by the absence of centrosomes in oocyte spindles. One, microtubules tend to assemble around the chromosomes. Two, the organization of these microtubules into a bipolar spindle is directed by the chromosomes. Three, chromosome biorientation and attachment to microtubules from the correct pole require modification of the mechanisms used in mitotic cells. Four, chromosome movement to the poles at anaphase cannot rely on polar anchoring of spindle microtubules by centrosomes. Overall, the chromosomes are more active participants during acentrosomal spindle assembly in oocytes, compared to mitotic and male meiotic divisions where centrosomes are present. The chromosomes are endowed with information that can direct the meiotic divisions and dictate their own behavior in oocytes. Processes beyond those known from mitosis appear to be required for their bi-orientation at meiosis I. As mitosis occurs without centrosomes in many systems other than oocytes, including all plants, the concepts discussed here may not be not limited to oocytes. The study of meiosis in oocytes has revealed mechanisms that are operating in mitosis and will probably continue to do so.

Keywords

meiosis; microtubule; spindle assembly; chromosome segregation; oocyte; acentrosomal

Introduction

A germ cell with two homologous alleles of each gene, one inherited from each parent, progresses through meiosis to produce four haploid cells with only one allele each. The haploid state is reached by replicating the genome once followed by two divisions. It is the first meiotic, or reductional, division that separates the two parental alleles into two daughter cells and is critical for Mendel's laws. Both meiotic divisions should be flawless because mistakes are usually lethal to the zygote. The success of meiosis usually depends on

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crossover formation between homologs to maintain a physical link between pairs of homologous chromosomes (bivalents) until anaphase I onset. That is why there is an inverse correlation between crossover frequency and nondisjunction of homologs (Baker and Hall 1976), and mutations that cause decreases in crossing over also increase nondisjunction. In addition, there is an increase in frequency of non-exchange chromosomes associated with human aneuploidy (Hassold and Hunt 2001; Nagaoka, et al. 2012).

The physical linkage of homologs that results from a crossover is called a chiasma, however, this linkage only directs segregation in combination with a bipolar spindle that orients homologs to opposite poles. Oocytes in many organisms have an added complication: spindle assembly occurs in the absence of centrosomes (referred to here as acentrosomal), the centriole-containing microtubule-organizing centers (MTOCs). The reason why centrioles are often eliminated prior to meiosis I in oocytes is not known but has been discussed elsewhere (Pimenta-Marques, et al. 2016; Severson, et al. 2016). The consequences of this centrosome elimination on spindle assembly, however, are dramatic. Instead of the chromosomes acting as passengers or cargo for the spindle, the chromosomes have the ability to organize meiotic spindles in a variety of systems (McKim and Hawley 1995).

The premise of this review is that chromosomes are active participants in the spindle assembly process. That the chromosomes carry information that can direct the meiotic divisions is long standing. For example, meiosis I chromosomes physically removed from a cell and placed into a meiosis II spindle still divide reductionally (Nicklas 1977; Paliulis and Nicklas 2000), showing the chromosomes dictate their own behavior. This review focuses on four aspects of meiosis I that are significantly affected by the absence of centrosomes in oocvte spindles: 1) assembly of microtubules around the chromosomes, 2) organization of these microtubules into a bipolar spindle, 3) chromosome bi-orientation and attachment to microtubules from the correct pole, and 4) chromosome movement to the poles at anaphase. Mitosis can and does occur without centrosomes in many systems other than oocytes, including all plants. Indeed, several reviews of cell division by acentrosomal mitosis have been published (Gatti, et al. 2012; Meunier and Vernos 2016; Moutinho-Pereira, et al. 2013; Wadsworth and Khodjakov 2004). Therefore, the knowledge gained from the study of oocytes is also relevant for many other cell types in a large range of organisms. In addition, for additional perspective on meiotic acentrosomal spindle assembly, a couple reviews have recently been published (Dumont and Desai 2012; Severson, et al. 2016).

Oocyte spindle assembly is dominated by chromosome-based mechanisms

Given the large size of oocytes (e.g. 120 µm diameter in humans) compared to most mitotically dividing cells (e.g. 40 µm diameter HeLa cell), nucleating and organizing microtubules must be combined with a mechanism to restrict spindle assembly to the vicinity of the chromosomes. Chromosome-directed spindle assembly has been observed in several oocyte systems. For example, early work in *Drosophila* and *Xenopus* oocytes and recent work in human oocytes show that spindle assembly begins with organization of microtubules around the chromosomes (Gard 1992; Holubcová, et al. 2015; Theurkauf and Hawley 1992). In some *Drosophila* mutants, oocytes chromosomes are ejected from the

main spindle and form their own spindles (Cullen, et al. 2005; Theurkauf and Hawley 1992). These results show that oocyte chromosomes can organize a bipolar spindle.

Enucleated mouse oocytes are unable to form spindles in a timely manner (<12 hrs) (Schuh and Ellenberg 2007). While other studies have reported spindle assembly in bisected (Brunet, et al. 1998) or enucleated oocytes (Yang, et al. 2007), these observations were made after a substantial period of time (18 hrs) and most spindles formed were grossly abnormal. In a result strikingly similar to what has been observed in *Drosophila*, dispersed clusters of mouse oocyte metaphase chromosomes, induced by nocodazole treatment, each organize a meiotic spindle (Maro, et al. 1986). Thus, it appears in mouse, and also pig oocytes (Sun, et al. 2001), that chromosomes organize microtubules, but other nuclear factors may also be critical for spindle assembly (Polanski, et al. 2005). In *C. elegans*, meiosis I oocyte spindle assembly appears to initiate with a microtubule "cage" that forms on the inside of the nuclear envelope (Wolff, et al. 2016) (although at meiosis II, the microtubules assemble closely around the chromosomes like the other systems). Observations like these raise the possibility that spindle assembly is coupled to the release of nuclear factors caused by disassembly of the nuclear envelope. This is a mechanism that oocytes could use to restrict spindle assembly to the vicinity of the chromosomes.

The first insights into identifying the molecules required for chromatin-based spindle assembly mechanisms came from work in Xenopus egg extracts. DNA-coated beads (Heald, et al. 1996) or sperm nuclei (Nachury, et al. 2001; Wilde and Zheng 1999) promote de novo spindle assembly in a process that depends on two mechanisms: a gradient of RanGTP centered on the chromosomes and the chromosomal passenger complex (CPC) (Carazo-Salas, et al. 1999; Sampath, et al. 2004). RanGTP, whose conversion from RanGDP is stimulated by chromosome-localized RCC1, causes the release of spindle assembly factors from the inhibitory effect of importins (Clarke and Zhang 2008; Meunier and Vernos 2016). Surprisingly, expression of a dominant-negative form of Ran (RanT24N) in human, mouse, and Drosophila oocytes demonstrated that the RanGTP pathway is not essential for assembly of the first meiotic spindle in vivo (Cesario and McKim 2011; Dumont, et al. 2007; Holubcová, et al. 2015). However, RanT24N caused delayed and disorganized assembly of the first meiotic spindle (Cesario and McKim 2011; Dumont, et al. 2007; Holubcová, et al. 2015). These data suggest that RanGTP contributes to the speed and efficiency of meiosis I spindle assembly in oocytes, but that other essential mechanisms are also present. Instead, the Xenopus, mouse and Drosophila studies suggest Ran becomes critical after meiosis I, during meiosis II and subsequent embryonic mitoses. For example, RanGTP appears to be essential for establishing the mouse metaphase II spindle (Dumont et al 2007).

Chromosome-based spindle assembly and the CPC

In the absence of the RanGTP gradient, the CPC promotes spindle assembly around sperm nuclei (Maresca, et al. 2009). The CPC is composed of four proteins: INCENP, Survivin, Borealin, and Aurora B (or C) kinase (Carmena, et al. 2012). The chromatin-focused enrichment of the CPC may facilitate local kinase activation, satisfying the spatial component to spindle assembly to be around the chromosomes (Kelly, et al. 2007). Two studies in *Drosophila* have supported a role for the CPC in acentrosomal spindle assembly in

In *Xenopus* egg extracts, both the INCENP centromere-targeting and microtubule-targeting domains are required to support chromatin-mediated spindle assembly. Therefore, it appears that the initiation of spindle assembly depends on simultaneous interactions between the CPC, the chromosomes and the microtubules (Tseng, et al. 2010). A partner for the CPC in this context could be motor proteins that bundle microtubules such as the *Drosophila* kinesin-6 Subito, which colocalizes with the CPC on the metaphase spindle in oocytes (Jang, et al. 2005). The bundling activity of the Subito appears to be activated only in the presence of the chromosomes after nuclear envelope breakdown (NEB) (Jang, et al. 2007). Thus, enforcement of spindle assembly around the chromosomes may also depend on the localized activation of motor proteins and their bundling activity.

The features of the chromatin that interact with the CPC to promote spindle assembly are not known. Sites that recruit the CPC include the centromeres, which may result in assembly of the kinetochores (Emanuele, et al. 2008; Kim and Yu 2015; Radford, et al. 2015; Rago, et al. 2015). Microtubules nucleated at or near kinetochores contribute to spindle assembly in somatic cells (Maiato, et al. 2004; Torosantucci, et al. 2008), but oocyte chromosomes lacking kinetochores still initiate spindle assembly (Brunet, et al. 1999; Deng, et al. 2009; Dumont, et al. 2010; Radford, et al. 2015). Therefore, there must be other chromosomal sites that make a significant contribution to oocyte spindle assembly mechanisms, although it is not yet clear what structures or molecules are important. In *Drosophila* mitotic cells, a chromosome-based spindle assembly pathway depends on the conserved Misato protein and the tubulin chaperone complex (Palumbo, et al. 2015). The role of these proteins in oocytes is not known.

Evidence of a single CPC-based pathway promoting initiation of spindle assembly in Drosophila oocytes has not been replicated in all organisms. For example, knocking down CPC components such as INCENP does not cause spindle assembly failure in mouse (Sharif, et al. 2010). The CPC may not be required for the initiation of spindle assembly in *C. elegans*, although it is required during meiosis because oocytes lacking the CPC have disorganized spindles (Schumacher, et al. 1998; Wignall and Villeneuve 2009). It may be that the existence of multiple Aurora homologs provides redundancy. For example, in mouse, like other mammals, there are three Auroras - AURKA, AURKB, and AURKC that may compensate for each other (Balboula and Schindler 2014; Fernández-Miranda, et al. 2011; Schindler, et al. 2012). Many eukaryotes express at least two Aurora kinases, one that is pole-associated (Aurora A) and one that transits from chromosomes to the central spindle (Aurora B/C). However, the designation of an Aurora kinase as either A or B/C does not always reflect their evolutionary relationships. For example, while mammal AURKB and AURKC are more similar to one another than they are to AURKA, AURKA is phylogenetically more closely related to AURKB and AURKC than to Aurora A in Drosophila (Brown, et al. 2004) (Figure 1).

In mouse oocytes, AURKC is the predominant catalytic subunit in the CPC (Balboula and Schindler 2014; Sharif, et al. 2010; Yang, et al. 2010). AURKC localizes to centromeres and chromosome arms during metaphase I and is restricted to centromeres at metaphase II, whereas AURKB localizes to microtubules (Balboula and Schindler 2014; Sharif, et al. 2010; Shuda, et al. 2009). A recent finding is that AURKC is recruited to spindle poles by Haspin kinase and is required for MTOC clustering (Balboula, et al. 2016). However, in oocytes from *Aurkc^{-/-}* mice, AURKB compensates, adopting AURKC localization and activity (Balboula and Schindler 2014; Schindler, et al. 2012). Conversely, embryos lacking *Aurkb* develop to the blastocyst stage because of AURKC (Fernández-Miranda, et al. 2011). Because of this compensation between AURKB and AURKC, establishing a requirement for initiating spindle assembly as a conserved function of Aurora kinase in oocytes has been challenging. Despite the evidence obtained from *Xenopus* and *Drosophila*, the role of the Aurora kinases in meiotic spindle assembly, and, more generally, the chromosome-based signal(s) that regulates spindle assembly continue to be an important topic of study.

Microtubule nucleation, bundling, and stabilization in oocyte spindle

assembly

Microtubules may be nucleated at many cytoplasmic locations such as membranes in *Drosophila* (Cha, et al. 2001; Theurkauf, et al. 1992), or at dispersed, small-sized cytoplasmic foci of pericentriolar material (PCM) also known as MTOCs in mouse oocytes (Luksza, et al. 2013). Mouse meiosis I oocytes contain a large number of acentriolar MTOCs (Szollosi, et al. 1972) that assemble microtubules at a rate similar to centriole-containing centrosomes in mitotic cells (Schuh and Ellenberg 2007). These MTOCs are also observed in preimplantation mouse embryos, which are also acentrosomal (Courtois, et al. 2012). Prior to NEB, an oocyte has 1–3 MTOCs, and this number increases to ~30 upon NEB, possibly due to MTOC fragmentation rather than *de novo* synthesis (Clift and Schuh 2015). Their biogenesis may require AURKA, which is localized at the poles, because oocytes from transgenic mice overexpressing AURKA showed an increase in MTOC numbers (Solc, et al. 2012).

Depletion of Pericentrin, an important component of acentriolar MTOCs, in mouse oocytes causes misaligned chromosomes and perturbs, but does not eliminate, bipolar spindle formation and maturation to metaphase II (Ma and Viveiros 2014). Despite a lack of γ -tubulin at these MTOCs, microtubules still nucleate, albeit the total amount of α -tubulin incorporated into the spindle is reduced. This reduction is similar to that of oocytes expressing dominant-negative RanT24N (Dumont, et al. 2007; Schuh and Ellenberg 2007). These results suggest that the Ran and MTOC pathways are not essential for spindle assembly in mouse oocytes. Similarly, in *Drosophila sas-4* mutant brain cells, centrosomes are absent and acentriolar MTOCs form, but they do not increase the efficiency of acentrosomal spindle assembly (Baumbach, et al. 2015).

Microtubules may also be nucleated within an existing spindle, which is usually associated with the Augmin pathway (Meunier and Vernos 2016; Sanchez-Huertas and Luders 2015). Augmin is not required to initiate mitotic spindle assembly, but instead enhances spindles by

building on pre-existing microtubules. In *Drosophila* oocytes, the Augmin pathway contributes to acentrosomal spindle function but is not required for assembly (Meireles, et al. 2009). Proteins like Dgp71WD/Nedd1, which recruit microtubule nucleating factors such as γ -TuRC, have also been implicated in microtubule nucleation within spindles (Reschen, et al. 2012). *Drosophila* oocytes lacking Dgp71WD have reduced density of tubulin and spindle width. It is possible these pathways are downstream targets of the CPC that contribute to specific populations of microtubules within the context of acentrosomal spindle assembly.

In the previous section, we emphasized the role of the chromosomes in stabilizing and bundling microtubules. In this section, we emphasized microtubule nucleation, which may have several mechanisms. To integrate these concepts, we propose a model with three distinct phases required in all acentrosomal systems: microtubule nucleation, bundling of microtubules, and stabilization around the chromosomes (Meunier and Vernos 2016) (Figure 2). In all systems, before NEB, the cytoplasm contains diffuse microtubules. At some point in meiotic progression, which may differ between organisms, there is a profound reorganization of cytoplasmic microtubules that may reflect enhanced bundling activity. This reorganization seems most dramatic in mouse oocytes with the appearance of MTOCs prior to NEB. However, in other cases this phase may be linked to NEB, which would be consistent with observations in human, *Drosophila* and *C. elegans* oocytes. Microtubule reorganization results in the bundling of the previously dispersed microtubules and depends on microtubule motors, such as the mouse kinesin-5 Kif11 (Clift and Schuh 2015). Such bundling may help restrict the microtubules to the vicinity of the chromosomes and, as described in the next section, may contribute to the assembly of a bipolar spindle.

Once NEB occurs, bundles of microtubules appear to be attracted to or stabilized by the chromosomes. Future research will be important to uncover how chromosome-based spindle assembly mechanisms are coordinated with non-chromosome-based mechanisms to form a functional oocyte spindle. While cytoplasmic elements may nucleate microtubules, and bundling may even occur prior to NEB, paramount is a mechanism that forces spindles to form around chromosomes. In all these cases, spindles reach a similar state after NEB, with a disorganized accumulation of microtubules near the chromosomes.

Bipolarity is established through a central spindle

Centrosomes function not only as a source of microtubule nucleation, but they also establish spindle bipolarity. An important question, therefore, is how oocytes accomplish bipolarity in the absence of centrosomes. Although oocytes lack centrosomes, the poles contain PCM components. The composition and assembly of these pole structures has been recently reviewed (Severson, et al. 2016). Some of these studies have shown that pole-localized proteins, and by inference pole focusing, are important for chromosome segregation. For example, proteins known to associate with acentrosomal poles include TACC, MSPS (Cullen and Ohkura 2001) and ASP in *Drosophila* (Riparbelli, et al. 2002) and *C. elegans* (van der Voet, et al. 2009; Wignall and Villeneuve 2009). Defective spindle poles and a pole-focusing defect were found in mouse oocytes lacking NuMA (Kolano, et al. 2012). The spindles were longer than normal and initially lacked focused poles. Despite kinetochore attachments

occurring normally, the spindle pole defects resulted in aneuploidy. These results suggest that pole focusing is important for chromosome segregation, but is not required for spindle bipolarity.

Results in *Drosophila* show that pole focusing is not sufficient to ensure only two poles. In *Drosophila* oocytes, a central spindle, composed of the antiparallel overlap of nonkinetochore microtubules, forms at metaphase I. The kinesin-6 motor protein Subito localizes to the central spindle (Jang, et al. 2005). In *subito* mutants, the central spindle is missing and oocyte spindles are frequently monopolar or tripolar, suggesting the central spindle is required to maintain a bipolar spindle (Giunta, et al. 2002) (Figure 3). Several other proteins that localize to the central spindle at anaphase in somatic cells also localize to the central spindle at metaphase in *Drosophila* oocytes, including the CPC and the centralspindlin complex (Das, et al. 2016; Jang, et al. 2005). Interestingly, loss of centralspindlin does not lead to the same spindle polarity defects seen in *subito* mutants (Das, et al. 2016), suggesting that Subito itself may provide the antiparallel microtubule crosslinking function required for oocyte spindle bipolarity.

These results are consistent with computational models that simulate key features of acentrosomal spindle assembly through sliding and bundling of microtubules by a plus-end-directed motor and clustering of minus ends by a minus-end-directed motor (Burbank, et al. 2007) or through microtubule crosslinking and microtubule depolymerization at minus ends (Janson, et al. 2007; Loughlin, et al. 2010). These models rely heavily on the antiparallel arrangement of microtubules that is typical of non-kinetochore microtubules. Non-kinetochore microtubules are a prominent feature of oocyte spindles, with estimates of their prevalence as high as 95% (compared to 5% kinetochore microtubules) in *Xenopus* egg extract spindles (Ohi, et al. 2007). The known properties of the kinesin-6 family are consistent with these models. Kinesin-6 proteins, which include the Subito mammalian homolog MKLP2, bundle antiparallel microtubules, an activity best known for its role in assembling the midzone at anaphase (Cesario, et al. 2006; Gruneberg, et al. 2004; Mishima, et al. 2002; Neef, et al. 2003; Nislow, et al. 1992; Tao, et al. 2016).

The role of Subito in other organisms in organizing the meiotic central spindle is not known. Other microtubule-bundling kinesins, for example kinesin-5, may have a similar function in mouse oocytes. Antiparallel microtubule crosslinking and sliding can be performed by kinesin-5 (known as Eg5, KIF11, BimC, or KLP61F). Inhibition of kinesin-5 in mouse oocytes with the small molecule inhibitor monastrol induces monopolar spindles (Mailhes, et al. 2004) (Figure 3). This phenotype is, however, distinct from that observed in *subito* mutants. In the absence of Subito, spindle poles undergo continuous cycles where they collapse into existing poles and then re-form, resulting in continual transitions between tripolar, bipolar, and monopolar spindles (Colombié, et al. 2008). In contrast, inhibition of kinesin-5 simply results in collapse of the spindle poles into a monopole (Mailhes, et al. 2004). These differences suggest that kinesin-5 and kinesin-6 have distinct roles in generating the bipolarity of oocyte spindles. In *C. elegans*, kinesin-5 is not essential for viability. A different microtubule-bundling kinesin, the kinesin-12, KLP-18, is required for spindle bipolarity (Segbert, et al. 2003). Much like *Drosophila*, at least conceptually, there is a ring containing the CPC and a motor protein (KLP-19, a kinesin-4) that forms around the

Although the role of the central spindle in promoting spindle bipolarity has not been confirmed in organisms other than *Drosophila*, some studies suggest it is important. As described below, bi-orientation in mouse may depend on a "prometaphase belt", which may be similar to the *Drosophila* meiotic central spindle. HURP, a conserved microtubule-associated protein, localizes to mouse oocyte central spindles in a kinesin-5-dependent manner and promotes microtubule stability in this region. Importantly, HURP is required for establishment and maintenance of a bipolar meiotic spindle through MTOC sorting and clustering (Breuer, et al. 2010). Like Subito, in the absence of HURP, there is a reduction in microtubule density specifically in the region of antiparallel overlap. These results are consistent with the idea that acentrosomal spindles require a robust central spindle to establish spindle bipolarity.

Bi-orientation without "search and capture" – the importance of converting lateral to end-on attachments

Formation of a stable, bipolar spindle is a critical step in cell division, but the ultimate goal is for this structure to direct the accurate segregation of the chromosomes. For accurate chromosome segregation to occur, chromosomes must become bi-oriented. One model for how bi-orientation occurs is "search and capture" – where microtubules growing from the poles randomly make end-on attachments to kinetochores (Kirschner and Mitchison 1986; Nicklas 1997). This process, however, may not be efficient enough to find and orient all the chromosomes in a typical cell. Cells containing centrosomes probably require additional mechanisms to supplement search and capture (Heald and Khodjakov 2015; Wollman, et al. 2005). This problem is only magnified in acentrosomal spindles with the absence of defined microtubule growth from the poles. One of these additional mechanisms supporting chromosome alignment is congression, where prometaphase chromosomes are collected to the center of the spindle (Cai, et al. 2009; Kapoor, et al. 2006). An important contributor to this process is lateral attachments between the chromosomes and the microtubules.

Several studies show that lateral attachments are important for the process of homolog biorientation on acentrosomal spindles (Figure 4). During prometaphase in mouse oocytes, chromosomes move towards the outside edges of the developing spindle and then congress via lateral attachments with microtubules to a ring around the central part of the spindle, or the "prometaphase belt" (Kitajima, et al. 2011; Magidson, et al. 2011). There are several important features of the "prometaphase belt" model. One, it highlights an important role for the central spindle, which is a prominent feature of oocytes in several systems. Not only an organizer of bipolarity, the central spindle and its structure of antiparallel microtubule overlaps is a rich source of microtubule plus ends. In this context, the "prometaphase belt" facilitates and enhances the rate of bi-orientation by bringing kinetochores into the vicinity of a high density of microtubule plus ends, which leads to stable kinetochore-microtubule attachments. Another important feature of the "prometaphase belt" model is the prominent

role of lateral chromosome-microtubule attachments. Oocytes experience a prolonged period during which stable end-on kinetochore-microtubule attachments are not observed (Brunet, et al. 1999). However, during a stage (2–4 hrs after NEB) when end-on attachments are not observed, the bivalents appear to stretch, indicating that there is enough stability in lateral attachments to provide force on the chromosomes and promote bi-orientation (Yoshida, et al. 2015).

The predominance of lateral attachments and absence of end-on kinetochore-microtubule attachments has also been observed in C. elegans oocytes (Wignall and Villeneuve 2009). C. *elegans* is an interesting case because the chromosomes are holocentric and form cup-shaped kinetochores. Lateral attachments sandwich the chromosomes between parallel bundles of microtubules, and this constrains and bi-orients the chromosomes (Wignall and Villeneuve 2009). The kinetochore protein KNL-1 is required for chromosome alignment in C. elegans oocytes (Dumont, et al. 2010), suggesting that these lateral attachments between chromosomes and microtubules are mediated by the kinetochore. Work in Drosophila oocytes has demonstrated that kinetochores are indeed essential for prometaphase chromosome movements and alignment (Radford, et al. 2015). Importantly, lateral kinetochore-microtubule attachments are sufficient to drive prometaphase chromosome movements in Drosophila oocytes (Radford, et al. 2015), consistent with the "prometaphase belt" model. Thus, in nematode, Drosophila and mouse, lateral attachments are important for bi-orientation and sufficient for substantial chromosome movement and interkinetochore stretch. End-on attachments may only be required to stabilize bi-oriented homologs (Radford, et al. 2015).

Bi-orientation, error correction and establishment of end-on attachments

Lateral attachments between kinetochores and microtubules, likely involving antiparallel bundles of microtubules in the central spindle, appear to be a conserved feature of acentrosomal meiosis. How these attachments lead to bi-orientation is not known. Several observations suggest bi-orientation in oocytes involves a mechanism of error correction. Homologous chromosomes undergo multiple rounds of stretching and relaxation in mouse oocytes (Kitajima, et al. 2011), reminiscent of "error correction". Similarly, real-time imaging of achiasmate *Drosophila* chromosomes suggest they also undergo multiple rounds of attachment to each spindle pole (Hughes, et al. 2009). Genetic studies have shown that mutants in checkpoint genes, such as in the *Drosophila* homolog of Mps1, display chromosome segregation errors in oocytes (Gilliland, et al. 2007; Gilliland, et al. 2005; O'Tousa 1982). Mps1 is also required for chromosome segregation in mouse oocytes (Hached, et al. 2011), although further studies are required to determine if Mps1 is required for error correction.

Homologous chromosome bi-orientation is often defined as the establishment of stable endon kinetochore-microtubule attachments, with sister chromatids (mitosis or meiosis II) or homologous chromosomes (meiosis I) making connections to opposite spindle poles. This places the chromosomes under tension, which can be observed as the stretching apart of sister or homologous centromeres. Errors are identified by a lack of end-on kinetochoremicrotubule attachment or a lack of the tension provided by these attachments. Surprisingly,

during meiosis I in mouse oocytes, homologous centromeres become stretched prior to the establishment of stable end-on kinetochore-microtubule attachments (Yoshida, et al. 2015). This is corroborated by data from *Drosophila* oocytes that shows homologous centromeres stretch apart in the absence of end-on kinetochore-microtubule attachments (Radford, et al. 2015). Similarly, *C. elegans* pairs of homologous chromosomes stretch towards opposite poles in the absence of end-on attachments (McNally, et al. 2016). These results suggest that end-on attachment and tension are separated in oocytes. Further research is needed to determine how incorrect attachments are identified, what defines bi-orientation and how it is monitored in meiosis I oocytes.

Does Aurora B/C have a role in error correction?

As in mitotic cells, Aurora B/C kinase could have a pivotal role in error correction. The spatial separation model for error correction suggests that incorrectly attached kinetochores are in close proximity to Aurora B kinase, which destabilizes the microtubule attachments, while correctly bi-oriented chromosomes are pulled away from Aurora B (Kalantzaki, et al. 2015; Lampson, et al. 2004). There are two problems with this model in meiosis. First, unlike mitosis and meiosis II, the sister centromeres of meiosis I orient towards the same pole; therefore, tension would not be predicted to displace kinetochores from Aurora B at inner centromeres (Watanabe 2012). Furthermore, this model probably does not apply to lateral kinetochore-microtubule attachments, which are resistant to Aurora B-mediated destabilization (Kalantzaki, et al. 2015). Because lateral attachments may have a role in oocyte bi-orientation, how error correction operates on lateral kinetochore-microtubule attachment of correctly bi-oriented chromosomes that are primed for the stabilization of end-on attachments is an important question for further study.

Additionally, evidence from the mouse suggests that Aurora B/C may not be regulated by attachment status. The conversion from lateral to stable end-on attachments in mouse oocytes depends on the timing of PP2A and CDK1 activity, rather than bi-orientation or tension/stretching status (Yoshida, et al. 2015). Even stretched apart and correctly bi-oriented chromosomes are treated like errors and "corrected" in an Aurora B-dependent manner in mouse oocytes (Yoshida, et al. 2015). During prometaphase, active Aurora B/C is located at the centromeres of stretched homologous chromosomes, which may explain the lack of stable end-on kinetochore-microtubule attachments (Yoshida, et al. 2015). Late in prometaphase, CDK1 activity increases (Davydenko, et al. 2013) and the phosphatase PP2A-B56 is recruited to centromeres (Yoshida, et al. 2015), which may promote the formation of stable end-on kinetochore-microtubule attachments, regardless of bi-orientation status. These observations help explain the failure to observe end-on attachments during most of mouse meiosis I (Brunet, et al. 1999).

These results appear to challenge the view that Aurora B/C is required to prevent incorrect attachments in oocytes. Numerous studies, however, have found that Aurora B is required for bi-orientation (Davydenko, et al. 2013; Kitajima, et al. 2011; Lane, et al. 2010; Rattani, et al. 2013; Shuda, et al. 2009; Yang, et al. 2010; Yoshida, et al. 2015). One possibility is that Aurora B/C destabilizes premature end-on attachments that occur during early prometaphase that may be at a high risk of forming stable improper attachments. Later,

Aurora B/C activity decreases, allowing the stabilization of end-on attachments. Interestingly, in *Drosophila sentin* mutant oocytes (Sentin is an EB1 interacting protein), end-on attachments are stabilized prematurely and the result is bi-orientation errors (Głuszek, et al. 2015). Thus, oocytes may use a bi-orientation mechanism that depends on destabilizing all attachments until there is a high probability that all are correct (Figure 2). In addition, pole-localized Aurora A activity can promote error correction of chromosomes that have wandered close to the poles (Chmátal, et al. 2015; Ye, et al. 2015), but whether this represents the primary mechanism for error correction in oocytes is not known. More research is needed to understand how such a system works.

Consistent with the idea that the emphasis for oocyte bi-orientation is making correct attachments rather than having to correct errors, the presence of one or even several univalents does not cause progression delays (Gui and Homer 2012; Kolano, et al. 2012; Lane, et al. 2012; LeMaire-Adkins, et al. 1997; Nagaoka, et al. 2012; Nagaoka, et al. 2011; Sebestova, et al. 2012). In some cases, univalents bi-orient (Kouznetsova, et al. 2007), which could satisfy the checkpoint. One interpretation is that oocytes lack a robust mechanism to correct erroneous attachments. Another possibility is that the checkpoint doesn't monitor chromosomes, but instead controls progression (i.e. the checkpoint as a timer hypothesis). This second possibility is consistent with the results described above from mouse oocytes in which stabilization of end-on attachments results from an increase throughout prometaphase in PP2A activity at centromeres, which silences the destabilizing activity of the CPC and depends on CDK1 and BubR1 (Davydenko, et al. 2013; Yoshida, et al. 2015). Therefore, in oocytes, the checkpoint and error correction machinery may function within an inflexible timescale, which could be tied to the developmental restrictions associated with oocyte maturation.

Segregation at anaphase

The bulk of studies into meiosis in oocytes (and also the bulk of this review) have been dedicated to understanding spindle assembly and homolog bi-orientation. It may be assumed that once these events have occurred, anaphase merely executes the pulling of end-on kinetochore-microtubule attachments to drive chromosome poleward movement. One problem with this simplistic view is that, in the absence of centrosomes, how are the poles anchored? Indeed, surprising results were found in *C. elegans*, where the poles are observed to move towards the chromosomes. At metaphase, the *C. elegans* meiotic spindle shortens until the kinetochores become embedded in the spindle poles (McNally, et al. 2016). The chromosomes then separate with sliding of the microtubules between them. This resembles Anaphase B (spindle elongation) and does not depend on kinetochore proteins (Dumont, et al. 2010). *C. elegans* meiosis also shows evidence of Anaphase A (chromosome movement), based on the observation that the chromosome move through channels in the microtubule array (Muscat, et al. 2015).

Pushing the chromosomes apart with microtubule sliding may also operate in mouse, where in meiosis II, anaphase B has been proposed to occur before anaphase A (FitzHarris 2012). This mechanism depends on spindle elongation driven by kinesin-5. The existence of these forces may explain observations in mouse that chromatin-coated beads move poleward at

anaphase (Deng, et al. 2009). Similar mechanisms may be operating in *Drosophila* based on elongation of the spindle observed in precocious anaphase, whereas chromosome to pole distance may not decrease (McKim, et al. 1993). These results suggest that anaphase includes a combination of A- and B-type chromosome movements, although relative to mitosis, possibly in reverse order or at the same time.

A final issue when considering anaphase is the concept and consequences of the asymmetric cell division in oocytes. Although meiosis can generate four haploid products, only one of these becomes the large oocyte, while the rest are retained in much smaller polar bodies. In a variety of organisms, the oocyte spindle is positioned asymmetrically prior to the division. Not only is the spindle located close to the cortex, but it may also rotate from an initial parallel orientation to perpendicular at anaphase I or II (McNally 2013). The mechanism of spindle positioning involves interactions with the cortex ("cortical pulling") mediated by actin and/or tubulin (Almonacid, et al. 2014; Brunet and Verlhac 2011). While the asymmetric division may only reflect elements in the cytoplasm, it is also possible that, as a consequence or cause, the meiotic spindle is asymmetric, with consequences on segregation. Indeed, the asymmetry in *C. elegans* meiosis causes a bias in the inheritance of univalents (Cortes, et al. 2015).

Because only one of the four meiotic products is incorporated into the zygote, an asymmetric spindle could have significant consequences on Mendelian segregation. If a pair of homologs have kinetochores that interact unequally with the spindle, then there is the potential for biased inheritance (Ross and Malik 2014). While this mechanism is plausible and attractive, it has proven difficult to observe. One study in mouse oocytes has revealed a link between asymmetric segregation and a property of the chromosomes (Chmátal, et al. 2014). This study demonstrated that some chromosomes (i.e. fusion chromosomes in natural populations) have stronger centromeres than other chromosomes (i.e. telocentric chromosomes in the same populations). These stronger centromeres appear to load more kinetochore proteins, and therefore interact with more microtubules, than weaker centromeres. It needs to be reiterated that such a difference in centromere strength only results in biased inheritance if there is an asymmetric spindle. Given the scarcity of such evidence (Hewitt 1976), more study is needed on the basis and consequences of the asymmetric spindle.

Summary

Chromosomes are the focal point of acentrosomal spindle assembly in oocytes. Amazingly, despite intensive knowledge about various spindle assembly pathways and proteins that mediate microtubule dynamics, there is still much to learn about what recruits the microtubules and what are the chromosome-based signals. For example, although the CPC is required in *Drosophila* and *Xenopus*, the generality of these findings is not clear. Most targets for acentrosomal spindle assembly come from mitotic candidates, which puts a limitation on finding factors involved in meiotic spindle assembly. New factors may be identified in screens, as has been reported in mouse oocytes, for spindle assembly factors (Pfender, et al. 2015). In addition to the central role of the chromosomes, the central spindle has a function at metaphase that is critical for spindle organization and bi-orientation. It

replaces the centrosomes with an array of sliding antiparallel microtubules that can establish bipolarity. The central spindle may also provide a kinetochore-independent source of microtubules.

This review has emphasized the importance of the chromosomes because most studies have focused on the trans-acting spindle assembly factors. Almost nothing is known about the chromatin features that promote acentrosomal spindle assembly. The kinetochores are clearly neither sufficient nor necessary. There could be epigenetic factors such as histone modifications (Kelly, et al. 2010; Wang, et al. 2010; Yamagishi, et al. 2010) or proteins like HP1 (Abe, et al. 2016; Ainsztein, et al. 1998), which are known to recruit the CPC. An interesting recent development is that noncoding RNAs may have a role in chromatin-mediated spindle assembly (Blower 2016; Chen, et al. 2015; Grenfell, et al. 2016; Jambhekar, et al. 2014).

In general, there are more examples of conserved concepts (eg. chromatin-mediated spindle assembly and the central spindle) than conserved proteins that carry out these functions. This could represent differences in mechanisms, such as in the case of the CPC and spindle assembly. In other cases, it could be due to lack of studies. For example, while Subito is required for bipolarity of the *Drosophila* oocyte spindle, *C. elegans* lacks a true orthologue of Subito and there is a lack of studies on the mouse orthologue MKLP2. It is possible, however, that a similar role to Subito is taken by kinesin-12 (in *C. elegans*) or kinesin-5 (in mouse). Similarly, there are few studies on the motors that regulate dynamics, although depolymerizing kinesins (kinesin-13) are important for regulating spindle length and microtubule attachments in *Drosophila* (Radford, et al. 2012) and *C. elegans* (Connolly, et al. 2015). It is not known if a shift from dynamic to stable microtubules accompanies the transition from prometaphase to metaphase in meiosis as it does in mitosis (Godek, et al. 2015; Kabeche and Compton 2013).

Lateral microtubule attachments with the kinetochore, probably involving the central spindle, are a conserved element of the mechanism for bi-orientation. The stability provided by the central spindle is important in this context when, early in meiosis, there are few end-on attachments and ill-defined poles. Lateral attachments probably predominate over any search and capture type of mechanism. However, much remains to be learned about how bi-orientation is measured, especially if this occurs at kinetochores that are making lateral attachments with microtubules. Indeed, the observation that end-on attachments are stabilized late and in a temporal manner, and that some lateral attachments may be stable, suggests that bi-orientation is established using mechanisms that do not depend on end-on attachments. The models from mitosis are insufficient to explain bi-orientation at meiosis I, and more interestingly, lessons learned from meiosis may reveal mechanisms that are operating in mitosis.

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Figure 1. Evolutionary relationships between Aurora kinases

Unrooted phylogenetic tree, assembled using the Maximum Likelihood method based on the JTT matrix-based model (Felsenstein 1992; Jones, et al. 1992). The tree with the highest log likelihood is shown. Initial trees for the heuristic search were obtained automatically by applying Neighbor-Join and BioNJ algorithms to a matrix of pairwise distances estimated using a JTT model, and then selecting the topology with superior log likelihood value. The tree is drawn to scale, with branch lengths measured in the number of substitutions per site. The analysis involved 22 amino acid sequences. All positions containing gaps and missing

data were eliminated. There were a total of 261 positions in the final dataset. Evolutionary analyses were conducted in MEGA7 (Kumar, et al. 2016). All three mammalian Auroras cluster together, indicating independent evolution of Aurora A/B in mammals, *Drosophila* and *C. elegans*.

Radford et al.



lateral MT attachment

end-on MT attachment

Figure 2. Model for acentrosomal spindle assembly

The emphasis of this model is that the accumulation and bundling of microtubules are regulated to occur in close proximity to the chromosomes. A) Oocyte before nuclear envelope breakdown. In most organisms, microtubules are visible outside the nuclear envelope. In mouse, these cluster in MTOCs. B) After nuclear envelope breakdown, there may be bundles or clusters of microtubules in close proximity, but distinct from, the chromosomes. This is seen in mouse and nematodes but not in fly or human. C) In all systems, the initial interactions between the chromosomes and microtubules may appear quite disorganized or appear as a ball of microtubules. D) A bipolar spindle forms and then lateral attachments predominate. E) Prior to anaphase, end-on attachments are stabilized. This last step is not clear in nematodes which lack point centromeres. In B and C, the kinetochores (red) are drawn relative to the chromosomes (blue) to show they are not stretched by the microtubules (green). In D and E, the kinetochores are drawn relative to the chromosomes to indicate they are stretched towards the poles.

Radford et al.



Figure 3. Defects in spindle bipolarity in Drosophila and mouse oocytes

A) In *Drosophila* oocytes lacking the Subito protein, tripolar spindles form. Microtubules are in green, DNA in blue, and centromeres (CENP-C) in red. Scale bar is 10 μ M. B) Metaphase II eggs from a control with bipolar spindle, and monastrol-treated, which inhibits kinesin-5, with a monopolar spindle. Microtubules are in green, DNA in white, and centromeres (CREST) in red. Scale bar is 10 μ m.



A) Wild-type *Drosophila* oocytes showing all kinetochore attachments in Drosophila and mouse oocytes A) Wild-type *Drosophila* oocytes showing all kinetochores making end-on attachments, with DNA in blue, microtubules in green, central spindle protein INCENP in red and centromeres (CENP-C) in white. B) Oocyte lacking the kinetochore protein NDC80, which is required for end-on attachments, shows evidence of lateral attachments. DNA is in blue, microtubules in green, and centromeres (CENP-C) in red. Inset is a 10 μ m region with the DNA removed showing most of the centromeres. C,D) Cold-treated mouse oocytes at metaphase I, showing examples of lateral (C) and end-on (D) kinetochore attachments. The boxed regions are shown at higher magnification. DNA is in blue, microtubules in green and centromeres (CREST) in red and the arrows point to lateral or end-on attachments. In all images, the scale bar is 10 μ m.