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► **To cite this version:**

Nora Bouftas, Katja Wassmann. Cycling through mammalian meiosis: B-type Cyclins in oocytes. Cell Cycle, In press, 10.1080/15384101.2019.1632139 . hal-02166962

HAL Id: hal-02166962

<https://hal.sorbonne-universite.fr/hal-02166962>

Submitted on 27 Jun 2019

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Cycling through mammalian meiosis: B-type Cyclins in oocytes

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Abstract

B-type cyclins in association with Cdk1 mediate key steps of mitosis and meiosis, by phosphorylating a plethora of substrates. Progression through the meiotic cell cycle requires the execution of two cell divisions named meiosis I and II without intervening S-phase, to obtain haploid gametes. These two divisions are highly asymmetric in the large oocyte. Chromosome segregation in meiosis I and sister chromatid segregation in meiosis II requires the sharp, switch-like inactivation of Cdk1 activity, which is brought about by degradation of B-type cyclins and counteracting phosphatases. Importantly and contrary to mitosis, inactivation of Cdk1 must not allow S-phase to take place at exit from meiosis I. Here, we describe recent studies on the regulation of translation and degradation of B-type cyclins in mouse oocytes, and how far their roles are redundant or specific, with a special focus on the recently discovered oocyte-specific role of cyclin B3.

Keywords:

Cyclin-dependent kinases; meiosis; oocytes; B-type cyclins; cyclin specificities; cell cycle control

Main Text

Introduction

The generation of healthy, haploid gametes depends on the correct execution of a specialized form of cell division named meiosis. Unlike mitosis, meiosis contains two steps of DNA division with only one round of DNA replication. These two divisions are called meiosis I and meiosis II. In oocytes, these two rounds of chromosome segregation are regulated temporally, are highly asymmetric and depending on the species, contain cell cycle arrest points that are crucial for female gametogenesis. In mammalian meiosis, after premeiotic S-phase and meiotic recombination, oocytes are arrested at prophase I from birth of the female onwards. Upon hormonal stimulation, this arrest is lifted and meiosis resumes with entry into the first meiotic division. Oocytes undergo the first metaphase-to-anaphase transition where chromosomes segregate and half the DNA is discarded in the smaller cell, the polar body. Oocytes enter meiosis II immediately following meiosis I exit. Oocytes must then arrest at metaphase II awaiting fertilization which triggers the second division with segregation of sister chromatids (Figure 1). Once the oocyte is fertilized, the embryo can begin its development (J. E. Holt, Lane, and Jones 2013; El Yakoubi and Wassmann 2017; Herbert et al. 2015).

It is primordial to understand the mechanisms that regulate the correct execution of female mammalian meiosis. Aneuploid embryos harbouring incorrect numbers of chromosomes, are mainly caused by maternal errors during female meiosis. 10-30% of all human oocytes are found to be aneuploid while only 1-2% of spermatozoa contain the wrong number of chromosomes (Templado, Uroz, and Estop 2013; Nagaoka, Hassold, and Hunt 2012; Hassold and Hunt 2001). This difference is increased in correlation with maternal age (Touati and Wassmann 2016; Hassold and Hunt 2001). By age 42, the incidence of aneuploidy may increase up to 80%, according to (Capalbo et al., « Human Female Meiosis Revised »). Not all segregation errors are due to aberrant meiotic divisions though, and additionally, a high rate of segregation errors is observed in the following mitotic divisions at later stages of embryogenesis, up to the blastocyst stage (Schneider and Ellenberg 2019; Vázquez-Diez and FitzHarris 2018). Today, with more commonly used medically assisted reproduction and the increase of maternal age at first pregnancies, it is indispensable to gain a better

comprehension of the regulatory mechanisms put in place to segregate the genetic material in meiosis and early mitotic divisions in the embryo. Insights into cell cycle regulation during meiotic maturation will help us understand what goes wrong so often in oocytes, which consequently leads to the generation of aneuploidies and hence, failure to generate healthy offspring.

Mammalian Meiosis

Although meiotic divisions are composed of the same steps as mitosis, the fact that two divisions without intervening S-phase are executed with entirely different segregation patterns requires additional regulatory mechanisms. Detailed reviews on different aspects of chromosome and sister chromatid segregation in meiosis I and II, respectively, have been published and are beyond the scope of this review (J. E. Holt, Lane, and Jones 2013; Wassmann 2013; Sanders and Jones 2018). Here, we will focus on very recent results concerning the role of B-type cyclins for cell cycle progression in mammalian oocytes.

In oocytes, the size of the cell, as well as the time frame of meiotic maturation to enter meiosis I and ensure fertilization takes place at the right moment of cell division demands distinct cell cycle regulation. Below we introduce progression through the meiotic divisions (meiotic maturation) from a cell cycle point of view, before describing the roles of the three existing mammalian B-type cyclins B1, B2 and B3 (Nieduszynski, Murray, and Carrington 2002; Chapman and Wolgemuth 1992; Pines and Hunter 1989), all three of which have now been studied by knock-out approaches in mouse oocytes (Karasu et al. 2019; Karasu and Keeney 2019; J. Li et al. 2018; Daldello et al. 2019; Y. Li et al. 2019; Brandeis et al. 1998).

Mammalian oocytes are formed during embryonic development and following DNA replication and recombination, oocytes arrest at prophase I. This is known as GV stage due to the presence of a germinal vesicle. This signifies that oocytes must stay arrested for extended periods of time (months for mice, decades for humans) before the meiotic divisions. Before resuming meiosis, selected oocytes undergo a short growth phase in the adult female. Upon hormonal stimulation, the GV breaks down (GVBD) followed by chromosome condensation and spindle formation (J. E. Holt, Lane, and Jones 2013; Herbert et al. 2015; El Yakoubi and Wassmann 2017). In mammals, a prolonged prometaphase in meiosis I allows the spindle to

fully form and microtubules (MT) to attach correctly to kinetochores (KT), hence chromosomes can align at the metaphase plate. Similarly to mitosis, the Spindle Assembly Checkpoint (SAC) is present and active during meiotic cell division. It imposes a delay in anaphase I onset by inhibiting the Anaphase Promoting Complex/ Cyclosome (APC/C), an E3 ubiquitin ligase essential for this cell cycle transition, to allow time for establishing correct KT-MT attachments (Touati and Wassmann 2016; Gorbsky 2015; Homer 2013). In order to allow metaphase-to-anaphase I transition to take place, the APC/C must be activated in order to ubiquitinate proteins that block separase activity, namely securin and cyclin B1. Once ubiquitinated, these proteins are degraded and separase becomes active leading to chromosome segregation (Herbert et al. 2003; Terret et al. 2003; Kudo et al. 2006). Oocytes then enter meiosis II immediately, and crucially, enter meiosis II without carrying out a second S-phase. Chromosomes then align and arrest at metaphase II. APC/C activation is inhibited to maintain this second meiotic arrest (also called Cystostatic factor (CSF-) arrest) until anaphase II is triggered by fertilization (Perry and Verlhac 2008; Runft, Jaffe, and Mehlmann 2002; Sanders and Jones 2018).

M-phase promoting factor (MPF)

Upon hormonal stimulation, the protracted GV arrest is lifted and this depends on the increase in MPF (M-phase promoting factor) kinase activity, the main driver of mitosis and meiosis. MPF phosphorylates a large number of substrates, and is required for entry into and progression through mitosis and meiosis, chromosome condensation, dissolution of the nuclear membrane, spindle formation, and chromosome alignment. It has to be inactivated for anaphase onset and exit from cell division. What makes up this factor has been extensively debated, but in both mitosis and meiosis MPF is considered as corresponding to the complex cyclin B-Cdk1 (Kishimoto 2015). However, MPF activity is not only due to cyclin B-Cdk1 complexes but also the kinase Greatwall (Mastl in mammals) that is essential to protect the phosphorylated substrates of cyclin B-Cdk1 from the counteracting phosphatase PP2A, allowing appropriate entry and progression through mitosis and meiosis (Vigneron et al. 2016; Kishimoto 2015; Adhikari and Liu 2014).

During the prolonged GV arrest in oocytes, Cdk1 is kept inactive through phosphorylation established by the Wee1 and Myt1 kinases, and APC/C dependent degradation of cyclin B

(mouse oocytes). Although the molecular mechanism is not completely known in mouse oocytes, depending on hormonal stimulation and release from surrounding follicle cells, activation of Cdk1 is triggered by the phosphatase Cdc25 that removes the inhibitory phosphorylation on Cdk1. To be active, Cdk1 needs to be associated with cyclins, in this case cyclin B1 or B2, to induce GVBD. Once Cdk1 is dephosphorylated and sufficient MPF activity is generated, GVBD occurs with resumption of meiosis (Han and Conti 2006; Solc, Schultz, and Motlik 2010).

Overview of M-phase B type cyclins

Most of the components needed for meiosis are also present during mitosis. This is the case for the main regulators of the cell cycle, cyclins. Several cyclins are expressed depending on the species and cell cycle phase. In order to fulfill their purpose, cyclins bind to their catalytic partners, Cdks (cyclin-dependent kinases). Progression through mitotic cell division is thought to require just one B-type cyclin together with Cdk1, as in both budding and fission yeast, Clb2 or Cdc13 respectively, is the only cyclin required to be paired up with Cdk1 to drive cells through mitosis. In *S. pombe*, it was even shown that Cdc13-Cdc2 (which corresponds to Cdk1) can -in principle- drive progression through all stages of the cell cycle, indicating that cyclin specificity is not determinant for ordering cell cycle events (Gutiérrez-Escribano and Nurse 2015; Fisher and Nurse 1996; Richardson et al. 1992). This has led to a model proposing that threshold levels of Cdk1-associated activity determines substrate specificities, and not individual cyclins. Importantly though, this model does not seem to apply to mitosis and meiosis in other organisms. At least in mammalian oocytes, specific B-type cyclins harbour distinct roles. Individual knock-outs of all three B-type cyclins have been described in the mouse and provide useful insights to the specificities of each cyclin during the meiotic divisions. In this review, we will discuss the redundant and distinct roles that B-type cyclins occupy in oocyte meiosis, highlighting the importance of each cyclin in the correct timing of cell cycle events during this specialized cell division.

Cyclin B1

In mammals, cyclin B1 is the main cyclin driving entry into and progression through mitosis and meiosis. Apart from phosphorylations impinging on Cdk1, protein levels of cyclin B1 directly correlate with Cdk1-associated kinase activity, regulating key steps of mitotic and

meiotic cell cycle progression. In mouse oocytes, cyclin B1-Cdk1 activity gradually increases for the first hours after GVBD, as oocytes progress through prometaphase, to reach its highest levels in metaphase I. This is due to inactivation of the APC/C activator Cdh1, and concomitant SAC activation, preventing activation of the APC/C in a Cdc20-dependent manner (McGuinness et al. 2009; Reis et al. 2007; Wassmann, Niaux, and Maro 2003; Herbert et al. 2003; Ledan et al. 2001). Slow progression through prometaphase allows the establishment of KT-MT attachments and correct alignment of chromosomes at the metaphase plate. For anaphase I onset, cyclin B1-Cdk1 activity must decrease drastically for a switch-like transition from metaphase to anaphase, and this is brought about by inactivation of the SAC and cyclin B1 degradation (Touati and Wassmann 2016; Gorbsky 2015; Sanders and Jones 2018). Cyclin B1-Cdk1 activity has to rise again quickly to inhibit S-phase onset, and instead promote entry into meiosis II with metaphase II or CSF-arrest. Upon fertilization, cyclin B1-Cdk1 activity must drop again for anaphase II onset with sister chromatid segregation (Perry and Verlhac 2008; Sanders and Jones 2018; Jones 2005).

Although protein synthesis is not necessary for GVBD, Cyclin B synthesis is essential for correct progression through meiosis in mouse oocytes (de Vantéry et al. 1996; Hashimoto and Kishimoto 1988). Following GVBD, the increase of cyclin B1 synthesis is responsible for the rise in Cdk1 activity, inducing chromosome condensation and congression as well as alignment during metaphase I. Another specificity of meiosis in mouse oocytes is that -unlike in cultured human cells - cyclin B1 is expressed in excess compared to its partner Cdk1 (Levasseur et al. 2019; Arooz et al. 2000), indicating that there is a distinct regulation of MPF activity in meiosis compared to mitosis. The large size of the cell and the extended time needed for prometaphase and metaphase might be the reason for this difference. Similarly to mitotic securin degradation which occurs preferentially when securin is not found in association with separase (Hellmuth et al. 2014; L. J. Holt, Krutchinsky, and Morgan 2008), Levasseur et al demonstrated in a very recent study that there are different pools of cyclin B1; free cyclin B1 and Cdk1-bound cyclin B1, and that association with Cdk1 determines temporal control of cyclin B1 degradation. Free cyclin B1 is degraded before all chromosomes are correctly attached and the SAC is turned off, in an APC/C-dependent manner. It continues to be degraded afterwards, through only its destruction box (Dbox), which is a motif recognized by the APC/C in association with Cdc20. The first degradation wave of free cyclin B1 depends on

a newly identified motif within the N-terminal helix of cyclin B1. However, when cyclin B1 is bound to Cdk1, it is somewhat protected from the first wave of degradation that depends on this new motif. The authors show that this newly identified degron-like motif in cyclin B1 is recognized by the APC/C leading to prometaphase degradation in mouse oocytes, and that the sequence is masked through binding of Cdk1. This would allow the cell to eliminate progressively excess cyclin B1 in prometaphase I in a SAC-independent manner, to allow the rapid inactivation of cyclin B1-Cdk1 needed for the anaphase switch to occur on time(Levasseur et al. 2019). At the same time, this would provide a stock of cyclin B1 available for association with Cdk1, to make sure that cyclin B1 does not become a limiting factor for Cdk1 activity in oocytes (Figure 2). Indeed, reducing the amount of free cyclin B1 in oocytes leads to defects in chromosome alignment at the metaphase I plate and hence, missegregation events(Levasseur et al. 2019). The authors propose that excess cyclin B1 compensates for weak SAC control in oocytes to ensure correct chromosome segregation(Levasseur et al. 2019). It is interesting to note in this context that endogenous cyclin A2 degradation occurs also in two waves in oocytes, with most of cyclin A2 being degraded in prometaphase I, and a small fraction of cyclin A2 localized to kinetochores escaping degradation until anaphase I onset(Touati et al. 2012). Future studies will show whether a similar motif as the one identified in cyclin B1 can be identified in the N-terminal helix of cyclin A2, or other B-type cyclins.

Additionally, the location of cyclin B1 and binding to separase may also play a role in regulating its ubiquitination by the APC/C. *In vitro* cultured mitotic cells seem to harbour a pool of cyclin B1 that preferentially inhibits separase near chromosomes, compared to cytoplasmic cyclin B1. The degradation of this pool of cyclin B1 inhibiting separase is slower compared to the cytoplasmic pool(Konishi et al. 2018), indicating that APC/C dependent degradation is determined by the partners cyclin B1 is associated with. Whether location of cyclin B1 and binding to separase regulate cyclin B1 ubiquitination also in oocytes is currently unknown.

Altogether, these studies indicate that additional regulation of cyclin B1 in meiosis and mitosis is accomplished through differential APC/C affinity for the separate pools present in the cell in prometaphase. When the SAC is satisfied, APC/C with its co-activator Cdc20, targets cyclin B1 for degradation through ubiquitination. This brings about a sharp decrease of Cdk1 activity

and exit from meiosis I. Once vertebrate oocytes exit meiosis I, they continue into meiosis II and arrest at metaphase II awaiting fertilization. In order for this arrest to be maintained, Cdk1 activity must be kept at high enough levels. Cyclin B1 is necessary for this very dynamic arrest, with an equilibrium of cyclin B1 degradation and synthesis to keep Cdk1 activity at an appropriate threshold(Kubiak et al. 1993; Nixon et al. 2002). In *Xenopus* oocytes, it was found that cyclin B1 and B2 synthesis continues during CSF arrest while Cdk1 activity level seems to be constant. It is thought that once a high amount of B-type cyclins is synthesized this leads to an increase in Cdk1 activity past a specific threshold. The APC/C with its co-activator Cdc20 becomes active inducing degradation of B-type cyclins. As a consequence, Cdk1 activity drops until another regulatory pathway, the Mos-MAPK pathway, counteracts APC/C activation allowing once again synthesis of cyclin B1 and B2. Therefore, CSF-arrest is dynamic, brought about by an equilibrium between cyclin B1 and B2 synthesis and APC/C activation(Yamamoto et al. 2005). Additionally, Cdc25 phosphatase is required during the arrest to remove inhibitory phosphorylation on Cdk1. Upon fertilization, the Mos-MAPK pathway is inhibited, allowing full APC/C activation and cyclin B1 and B2 degradation, while Cdk1 is additionally downregulated by the kinase Wee1B, which phosphorylates Cdk1 for further inhibition(J. S. Oh et al. 2013; Jeong Su Oh, Susor, and Conti 2011; M.-H. Verlhac et al. 2000; M. H. Verlhac et al. 1994). Oocytes exit meiosis II and form a female pronucleus, which fuses with the male pronucleus provided by the sperm, to re-establish the diploid chromosome content in the zygote(Clift and Schuh 2013).

Cyclin B1 vs cyclin B2

Overall, cyclin B1 expression seemed to be necessary for development and fertility in mice while cyclin B2 is dispensable(Brandeis et al. 1998). Cyclin B1 null mice are not viable while cyclin B2 null mice are viable and fertile, although litters are smaller in size and numbers(Brandeis et al. 1998). Lack of cyclin B1 in embryos induces an arrest after only two divisions at the 4-cell stage, underlining its significance in embryonic development(Strauss et al. 2018). Both cyclins are expressed during mitosis although they are localized differently, next to microtubules (cyclin B1) and the Golgi region (cyclin B2), respectively, and Cyclin B1 is translocated to the nucleus during prophase unlike cyclin B2(Jackman, Firth, and Pines 1995). In mouse oocytes, the distribution of exogenous cyclin B1 evolves before GVBD with a cytoplasmic localization at first followed by its import into the nucleus(Marangos and Carroll

2004). Nuclear targeting of exogenous cyclin B1 significantly increased its capacity to induce entry into meiosis I (J. E. Holt, Weaver, and Jones 2010), indicating that localization is an important factor regulating cyclin B1's capacity to induce GVBD. As the importance of cyclin B1 and B2 subcellular localization for its role during meiotic maturation is largely unknown in mouse oocytes, it will not be subject of further discussion here.

Translational regulation is specific to each cyclin in mouse oocytes. The translation of cyclin B1 is tightly regulated in mouse oocytes since at different time points, cyclin B1 has to be rapidly synthesized. mRNA granule formation plays a role in inhibiting translation while their dispersion allows quick synthesis of cyclin B1 in mouse oocyte meiosis I (Kotani et al. 2013). Additionally, timing of cyclin B1 translation is regulated through polyadenylation mediated by specific lengths of the 3'UTR. Long and intermediate 3'UTR length of cyclin B1 mRNA leads to repression of translation in GV arrested oocytes, and to their translation after GVBD while cyclin B1 mRNA with short 3'UTR is constitutively translated already in GV oocytes (Yang et al. 2017). Following GVBD, the synthesis and amount of cyclin B1 increases while cyclin B2 remains at stable levels (Figure 3). Unlike cyclin B2, cyclin B1 transcripts are regulated through ribosome loading, dependent on the RNA-binding protein CPEB1. This loading of ribosomes on cyclin B1 increases after GVBD while ribosome loading on cyclin B2 does not change after resumption of meiosis. Intriguingly, Cdk1-associated activity was found to play a role in the phosphorylation of CPEB1 and therefore, to regulate cyclin B1 translation (Han et al. 2017). It is attractive to speculate that cyclin A2, a strong inducer of GVBD when expressed exogenously (Touati et al. 2012), may act as a trigger for entry into meiosis I by mediating phosphorylation of CPEB1, but this has not been addressed yet.

Complete loss of either Cyclin B1 or B2 in mouse oocytes

To elucidate the contributions of cyclin B1 and B2 for meiotic progression in oocytes, two recent studies examined meiotic maturation in mice harbouring oocytes devoid of either cyclin with knock-out approaches targeting the gene encoding cyclin B1 (*Ccnb1*) or cyclin B2 (*Ccnb2*). In mitotic tissues, only cyclin B1 seems to be essential, and a conditional knock-out approach using a *Gdf9*-dependent deletion strategy was necessary to determine its role in oocytes. Surprisingly though, oocytes without cyclin B1 are still able to enter meiosis I and extrude a polar body. No abnormalities as far as chromosome alignment and spindle

formation is concerned, were observed in oocytes devoid of cyclin B1. However, once these oocytes exit meiosis I, they fail to enter meiosis II and to establish a CSF arrest. At entry into meiosis I, cyclin B2 is upregulated and promotes GVBD in these *Ccnb1*^{-/-} oocytes, but after meiosis I, except when over-expressed, cyclin B2 cannot substitute for cyclin B1. Because oocytes fail to re-activate Cdk1 after meiosis I, *Ccnb1*^{-/-} oocytes decondense chromosomes and seem to enter an interphase-like state. Importantly, injection of mRNA coding for cyclin B2 can rescue meiosis II in *Ccnb1*^{-/-} oocytes, indicating that the failure of cyclin B1 oocytes to progress beyond meiosis I may be due to a failure to re-accumulate sufficient levels of cyclin B2 to compensate for loss of cyclin B1 such as in meiosis I, and not due to a specific role of cyclin B1 in meiosis II. Collectively these data indicate that cyclin B2 can substitute for cyclin B1 in oocyte meiosis (Brandeis et al. 1998; J. Li et al. 2018).

Ccnb2 knock-out mice are viable, allowing the analysis of meiosis in oocytes devoid of cyclin B2. Without cyclin B2, entry into meiosis I is significantly delayed (Daldello et al. 2019; J. Li et al. 2018; Gui and Homer 2013), because oocytes fail to activate Cdk1 efficiently for GVBD. Unlike cyclin B2 in *Ccnb1*^{-/-} oocytes, no upregulation of cyclin B1 to compensate for loss of cyclin B2 was observed. Quite the opposite, translation of mRNA coding for cyclin B1 was affected in the absence of cyclin B2 in a proportion of oocytes, indicating that this cyclin has a specific role in translational control of cyclin B1 and other, oocyte-specific mRNAs such as coding for Mos (Daldello et al. 2019). How cyclin B2 brings about translational control of meiotic transcripts is currently unknown, but may involve phosphorylation of CPEB1, which occurs in a Cdk1-dependent manner (Han et al. 2017). If cyclin B2 together with Cdk1 is able to phosphorylate CPEB1, it will be important to determine whether this is due to specific substrate specificity of cyclin B2, or due to the fact that cyclin B2 translation occurs independently of the CPEB1. Because of missing cyclin B2 protein in GV oocytes and failure to accumulate sufficiently high levels of cyclin B1, overall MPF activity is lower in *Ccnb2*^{-/-} oocytes, resulting in defects in spindle formation, delayed and inefficient APC/C activation, delay in anaphase I onset and reduced number of oocytes that succeed to extrude a polar body and exit meiosis I. These defects were rescued by overexpression of cyclin B1 (Daldello et al. 2019). The authors suggest that the metaphase I arrest observed in a significant proportion of *Ccnb2*^{-/-} oocytes is due to SAC activation because inhibition of the essential SAC kinase Mps1 can rescue polar body extrusion. *Ccnb2*^{-/-} oocytes that fail to extrude a polar

body show kinetochore recruitment of the SAC component Mad2 indicating prolonged SAC activation, even though under normal conditions the SAC is activated only very transiently in meiosis I (Daldello et al. 2019). A small caveat of this assay is the fact that without a functional SAC, oocytes undergo metaphase-to-anaphase transition precociously and with much lower Cdk1 activity (Touati et al. 2015; Rattani et al. 2014). Therefore oocytes devoid of cyclin B2 may harbour just enough cyclin B1-Cdk1 activity to undergo metaphase-to-anaphase transition in the absence of a functional SAC, explaining the rescue. Overall, the data obtained from complete loss of either cyclin indicates that cyclin B1 and B2 harbour specific roles during meiotic maturation mainly because of their expression profiles brought about in part by a specific role of cyclin B2 for translational control of cyclin B1. They can substitute for each other when expressed at the required times during meiotic maturation (Daldello et al. 2019; J. Li et al. 2018), indicating that they can phosphorylate the same substrates throughout meiosis I and II.

Cyclin B3

Evidence from the nucleotide sequence, cyclin box conservation, expression pattern and Cdk binding site pointed to cyclin B3 being a distinct and unique cyclin. Cyclin B3 was first described in chicken and found to exhibit 33% identity with cyclin B2 and 30% identity with A-type cyclins showing that it resembles both types of cyclins. Cyclin B3 was found to be located in the nucleus in interphase, similarly to A-type cyclins, while cyclin B1 is located in the cytoplasm (J.-C. Lozano et al. 2002; Gallant and Nigg 1994; Sigrist et al. 1995). In *C. elegans*, cyclin B3 was found to be close to chicken cyclin B3 with 50% identity in the cyclin box. Cyclin B3 is mainly expressed in maternal germ cells and conserved in other worms (Kreutzer et al. 1995). In humans, cyclin B3 mRNA was detected by RT-PCR in testis and 8 other tissues, but only in testis by northern blot analysis, suggesting low abundance in tissues other than germ cells. Protein levels and mRNA expression was found at highest levels in prepachytene spermatocytes and in oocytes during embryonic development (J.-C. Lozano et al. 2002). This particular expression hinted at a role in early meiosis I (prophase) in both male and female meiosis.

Concerning Cyclin B3's sequence encoded by *Ccnb3* in mice, an important extension of exon 8 is observed in placental mammals and not in other species, suggesting that the resulting 3-

fold increase in the size of the protein occurred during evolution of placental mammals. The extension and increase in size makes placental mammals' cyclin B3 of substantial size compared to other species; human cyclin B3 is 1395 aa long while chicken and *Xenopus* cyclin B3 are 403 aa and 416 aa long, respectively. This extended sequence could mediate new protein-protein interactions that may be important for its function specifically in these species(J. C. Lozano et al. 2012). Contrary to human and mouse cyclin B3, *Drosophila* and *C.elegans* cyclin B3 is expressed in both meiotic and mitotic cells during embryogenesis(Jacobs, Knoblich, and Lehner 1998; Bourouh et al. 2016; Deyter et al. 2010; Kreutzer et al. 1995; Michael 2016; Yuan and O'Farrell 2015). Although it is expressed in mitotic *Drosophila* cells, cyclin B3 was found to be dispensable for mitosis but essential for female meiosis(Jacobs, Knoblich, and Lehner 1998). This differs from *C. elegans* cyclin B3 which is essential for embryonic development. In humans, aberrant expression of cyclin B3 during mitosis has been observed in a specific type of cancer, Ewing-like sarcoma, where a fusion of cyclin B3 and a ubiquitously expressed protein has been found(Cohen-Gogo et al. 2014; W.-S. Li et al. 2016; Pierron et al. 2012), indicating that untimely expression of this cyclin can have serious consequences on cell fate.

The importance and role of cyclin B3 has been established in several species although the molecular mechanism is not fully understood. In *Drosophila* and *C. elegans* oocytes and embryos, cyclin B3 was found to be a partner of Cdk1 although a possible interaction with Cdk2 was not examined(Jacobs, Knoblich, and Lehner 1998; van der Voet et al. 2009). In chicken mitotic cells, cyclin B3 was able to bind to both Cdk1 and Cdk2(Gallant and Nigg 1994). Surprisingly, human cyclin B3 did not co-immunoprecipitate with Cdk1 but was found to interact with Cdk2 resulting in very low *in vitro* kinase activity(Nguyen et al. 2002). Mouse cyclin B3 purified from insect cells is able to be partnered up with Cdk1 and associated *in vitro* kinase activity was detected(Karasu et al. 2019).

In *C. elegans*, cyclin B3 knockdown is lethal and detrimental to development. Cyclin B3 deficient embryos show several phenotypic defects: In meiosis, cyclin B3 depleted oocytes are not able to segregate sister chromatids during anaphase II although other cell cycle steps occur. Once both male and female pronuclei are formed, their migration is slower in mutant embryos. During the first mitosis, metaphase was found to be delayed and chromosomes do

not segregate. This phenotype is partially rescued by simultaneously knocking down SAC components indicating that cyclin B3's role in *C. elegans* is SAC dependent and essential for embryonic development(Deyter et al. 2010). As previously discussed, cyclin B3 in *Drosophila* is dispensable for viability and male meiosis but essential for female meiosis. Cyclin B3 deficient *Drosophila* females are sterile(Jacobs, Knoblich, and Lehner 1998). In *Drosophila* oocytes, cyclin B3 shares some functions with other cyclins, cyclin A and cyclin B, but also harbours a unique role in promoting anaphase onset in both oocytes and embryos, contrary to cyclin B1 that inhibits anaphase onset(Sigrist et al. 1995; Yuan and O'Farrell 2015; Jacobs, Knoblich, and Lehner 1998; Bourouh et al. 2016). Furthermore, together with cyclin A, cyclin B3 inhibits entry into S phase between meiosis I and II(Bourouh et al. 2016). Although cells progress slower through the cell cycle, cyclin B1 is dispensable for mitosis in *Drosophila*. Only double knockdown of cyclin B3 and B1 leads to embryonic lethality once maternal reserves are depleted(Jacobs, Knoblich, and Lehner 1998; Bourouh et al. 2016). Surprisingly, a different role of cyclin B3 was uncovered in the ascidian *Ciona intestinalis*. *Ciona intestinalis* cyclin B3 was found to be essential for maternal to zygotic transition, where zygotic genome activation (ZGA) occurs. The authors described a correlation between the decrease of cyclin B3 mRNA and start of ZGA. Knockdown of cyclin B3 led to advanced initiation of ZGA(Treen et al. 2018) demonstrating yet another role for this cyclin.

Through the analysis of *ccnb3* knock-out male mice it was found that although expressed in testis in mice, its role is dispensable during spermatogenesis(Karasu et al. 2019; Karasu and Keeney 2019). Furthermore, its protein and mRNA levels are low in adult ovaries; nevertheless, its role is essential for correct meiotic maturation, as oocytes cannot progress beyond metaphase I without cyclin B3(Karasu et al. 2019; Y. Li et al. 2019; Zhang et al. 2015). Complete loss of cyclin B3 leads to a SAC-independent metaphase I arrest, with high Cdk1 activity and a failure to correctly degrade endogenous APC/C substrates(Karasu et al. 2019; Y. Li et al. 2019), even though exogenously expressed substrates are degraded(Karasu et al. 2019). This may indicate that cyclin B3 is required for targeting specific pools of cyclin B1 and securin for APC/C dependent ubiquitination, such as cyclin B1 associated with Cdk1, and securin in association with separase(Konishi et al. 2018; Hellmuth et al. 2014; Levasseur et al. 2019). Alternatively, cyclin B3 may be involved in regulating substrate specificity of the APC/C by phosphorylating one of its subunits, or by activating phosphatases that determine

substrate specificity through some yet unknown mechanism (Figure 4). Lastly, cyclin B3 may be required for translational regulation of cyclin B1 and securin mRNAs, and in its absence, translation is upregulated leading to the observed failure to degrade endogenous substrates and metaphase I arrest. A cyclin B3 mutant that does not yield *in vitro* kinase activity with Cdk1 was also not able to rescue oocytes devoid of cyclin B3(Karasu et al. 2019), strongly suggesting that Cdk1 activity associated with cyclin B3 is essential for its role in female meiosis. Most importantly, neither cyclin A2 nor cyclin B1 can substitute for cyclin B3, showing that cyclin B3 has a non-redundant and specific role in oocytes, indicating that it confers distinct substrate specificity to Cdk1 than cyclin A2, B1, or B2(Karasu et al. 2019). As mentioned above, cyclin B3 in placental mammals is three times bigger than from other vertebrates(J. C. Lozano et al. 2012). It came therefore as a surprise that the function of cyclin B3 seems to be conserved between species, because expression of cyclin B3 derived from *Drosophila*, zebrafish and *Xenopus* was able to rescue the meiotic defects of *Ccnb3* knock-out mouse oocytes(Karasu et al. 2019). The extended exon 8 in placental mammals seems therefore not absolutely required for cyclin B3 function in meiosis.

Conclusion

Recent studies using knock-out mouse models and sophisticated imaging approaches have provided important insights on the roles of B-type cyclins during female mammalian meiosis. Cyclin B2 occupies a specific role in oocytes, regulating translation of meiotic mRNAs, and this role of cyclin B2 is essential for accumulating sufficient levels of MPF activity for meiotic cell cycle progression. Cyclin B1 and B2 seem to be largely redundant as far as MPF substrate phosphorylation for progression through meiosis I and meiosis II are concerned, and their individual roles in oocyte meiosis are most likely due to differences in expression and localization. Cyclin B3 clearly occupies a distinct role in meiotic cell cycle progression, specific to oocytes. Cyclin B3 is required for metaphase-to-anaphase transition in meiosis I, whereas cyclin B1 and B2 inhibit anaphase I onset. Recent data indicate that cyclin B3-Cdk1 substrates are not shared with other A- and B-type cyclins, because none of them can substitute for cyclin B3, putting this cyclin apart. Future work will aim at identifying the targets of cyclin B3 to elucidate why this cyclin is required specifically in oocytes. It is attractive to speculate that the size of the oocyte requires the additional contribution of cyclin B3 for ensuring switch-like transitions in meiosis.

Acknowledgements

We thank members of the Wassmann lab for discussion, and S.A. Touati (Francis Crick Institute, London) for valuable comments. NB received a 3-year PhD fellowship from the French Ministère de la Recherche, followed by a 1-year PhD fellowship by the Fondation ARC pour la Recherche sur le Cancer. The lab of K.W. is financed through a grant "Equipe FRM" by the Fondation de la Recherche Médicale (Equipe DEQ20160334921, to KW), an international ANR grant (ANR-16-CE92-0007-01 to KW), and core funding from Paris Sorbonne Université and the CNRS.

The authors report no conflict of interest.

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Figure Legends

Figure 1: **Mammalian oocyte maturation.** Oocytes are arrested at prophase I, also known as germinal vesicle (GV) stage. Upon hormonal stimulation, fully-grown oocytes resume meiosis with germinal vesicle breakdown (GVBD). Chromosomes (in blue) condense, the bipolar spindle forms and microtubules attach to the chromosomes throughout prometaphase. In metaphase I chromosomes that are properly attached to the spindle and aligned at the metaphase plate migrate to the cortex. Chromosomes segregate during anaphase I with the extrusion of a first polar body. Oocytes immediately enter meiosis II and arrest at metaphase II awaiting fertilization.

Figure 2: **Regulation of cyclin B1 degradation during meiosis I.** At early prometaphase, different pools of cyclin B1 are present in the oocyte: free cyclin B1 and cyclin B1 bound to Cdk1. During prometaphase, free cyclin B1 starts to be degraded while the SAC is on, whereas Cdk1-bound cyclin B1 is protected from APC/C-dependent ubiquitination and degradation through its binding. Most of the free cyclin B1 is degraded once oocytes progress into metaphase I, while bound cyclin B1 begins to be degraded only once the SAC is satisfied and Cdc20-APC/C under SAC control becomes active. Once the APC/C is fully active, cyclin B1 bound to Cdk1 is degraded leading to a sharp decrease of Cdk1 activity and anaphase I onset.

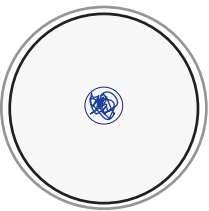
Figure 3: **Protein levels of cyclin B1 vs. cyclin B2 in meiosis I.** According to Han et al. 2017, preceding resumption of meiosis, cyclin B2 (in red) is translated at higher levels than cyclin B1 (in blue) while their degradation rates are similar. Therefore, at GV stage, cyclin B2 protein is more abundant than cyclin B1. After entry into meiosis I, protein levels of cyclin B2 remain similar with only a slight increase in translation. However, cyclin B1 is actively translated leading to an increase of protein levels following GVBD reaching its maximum at metaphase I. Cyclin B1 translation depends on cyclin B2 protein at entry into meiosis I. Hence, protein levels of both cyclins are regulated differentially and temporally in mouse oocytes.

Figure 4: **Model of how cyclin B3 may promote anaphase I onset in mouse oocytes.** Cyclin B1/B2-Cdk1 (in blue) activity increases as oocytes progress through prometaphase into metaphase, because the APC/C (light blue) in association with its activator Cdc20 (brown) is

kept in check by the SAC. At the metaphase-to-anaphase transition, the SAC is satisfied and inactivated, allowing full APC/C activity, ubiquitination and hence degradation of cyclin B1 and B2 in association with Cdk1. In oocytes, full APC/C activity requires the function of cyclin B3-Cdk1. Cyclin B3-Cdk1 (in green) promotes APC/C activity leading to chromosome segregation and exit from meiosis I. Therefore cyclin B3, a late substrate of the APC/C, shows an opposing role to cyclin B1 and B2 during mouse oocyte meiosis I.

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Figure 1

GV



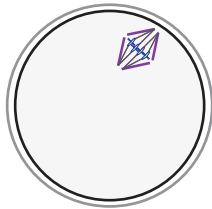
GVBD



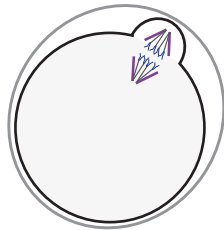
Prometaphase I



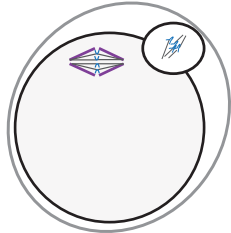
Metaphase I



Anaphase I

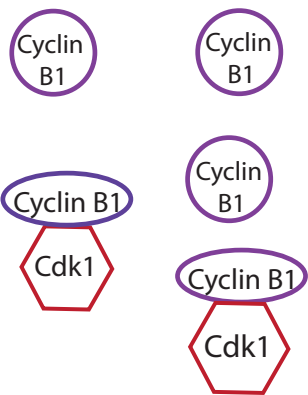


Metaphase II

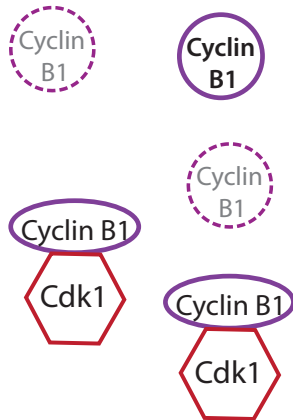


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Figure 2

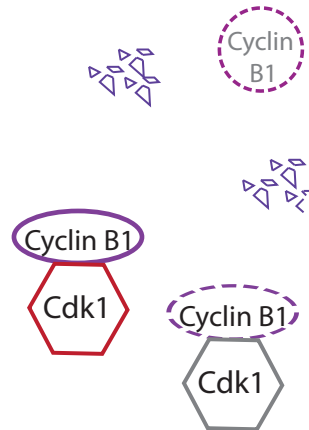
Early Prometaphase



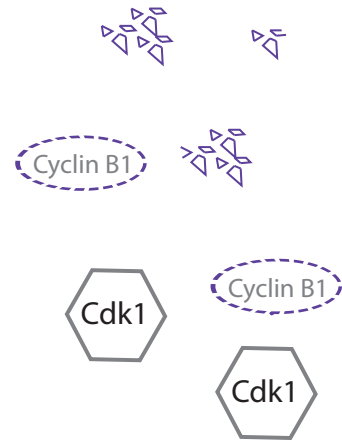
Prometaphase I



Metaphase I

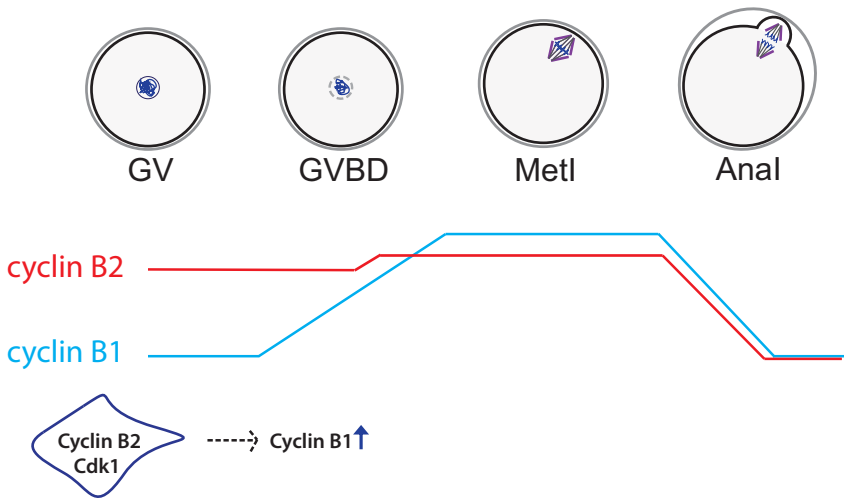


Anaphase I



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Figure 3



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Figure 4

