MUNI Med

Embryologie I OOGENESIS

autumn 2024

Oocyte meiosis

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Prenatal stage of oogenesis



Entry to meiosis



(1) Premeiotic S phase - DNA replication

(2) Meiosis I

- separation of homologous chromosomes

(3) Meiosis II

- separation of sister chromatids



1C 1n 2C 1n Homologous chromozomes 1C 1n pairing Patrenal chromosome 1C 1n Leptotene **DNA replication** Maternal Zygotene chromosome Pachytene 1C 1n 2C 2n 4C 2n 2C 1n Diplotene Diakineze I. Meiotic division II. Meiotic division

Meióza



Prophase I



Homologous chromosomes pairing

- synapsis = physical association of homologous chromosomes
- homolous sequence pairing
- Synaptonemal complex
 - axial elements SYCP3+SYCP2
 - central element dimer SYCP1
 - lateral elements cohesins (Rad21, Rec8)
- gradual formation of SYCP3 a SYCP2 foci
- axial element built by coalescence of SCP3 and SCP2 foci with cohesin complex proteins
- axial element stabilized by cross-linking with SYCP1 dimer, which forms central element









Homologous chromosomes pairing

 pairing of distant homologous chromosomes accomplished by congregation of telomeres attached to nuclear membrane

 \rightarrow tzv. "telomere bouqouet"



Non-specific telomere associations

Stable pairing near chromosome ends



Berrios et al 2013

Zhou et al 2012

Elkouby and Mullins 2017

Homologous recombination



Homologous recombination

- DNA double strand breaks (DSB) precede synapsis formation



Homologous recombination

- PRDM9
 - histon metyltransferase
 - recognition and epigenetic modification of specific DNA sequences (recombination hotspots)



- DNA endonuclease
- its dimer forms DSB

-/-

mouse Spo11-/- males sterile

Spo11 polymorfism in male infertility

- each monomer then binds 5' end of ssDNA
- Spo11+oligonucleotide cleaved away by exonuclase → free 3'end



PRDMS

Homologous recombination

- RPA (replication protein A)
 - binds free 3'end
 - recruits DMC1 and Rad51
 - during invasion binds DNA template strand and stabilizes D-loop
- DMC1 + Rad51
 - meiotic recombinases
 - bind and navigate free 3'end of ssDNA to invade dsDNA of homologous chromosome





RPA binds template in D-loop

DMC1 performs strand exchange

Hinch et al 2020



Homologous recombination

Holliday junction (HJ)

- named after Robin Holliday, who proposed its existence in 1964
- DNA duplex physical linkage of two DNA doublehelixes
- intermediate of homologous recombination and DSB repair mechanism
- visible in electron microscope
- HJ can move, double HJ can be resolved















Homologous recombination

non-crossover ~ 90

DSB

- no gene conversion

(a) D-loop resolution
 → DNA synthesis according to complementary strand

(b) convergent branch migration resulting in resolution of HJs

(c) strand exchange and HJ resolution without gene conversion

crossover ~10%

- gene conversion occurs at both chromatids
- HJ resultion produces new combination of genes



https://www.youtube.com/watch?v=3qgBKrAZCLg

Homologous recombination

- Chiasmata (chi-structure)
- physical contact sites of homologous chromosomes marking crossing-over regions
- visible after synaptonemal complex disolution during diplotene
- links homologous chromosomes together in the form of bivalents (tetrades)
- disapprear as homologous chromosomes separate during anaphase I
- sex differences in location and number of chiasmata (more distal in males)
- altered number of chiasmata and location of chiasmata associated with aneuploidy



Microtubules attach to the fused kinetochore:

of the sister chromatids

emain attached a

the centromer

Regulation of meiotic prophase overview



Chromatin configuration during diplotene arrest

- dictyate stage = prolonged diplotene arrest
- chromosomes become dispersed, less distinct and form faint network
- germinal vesicle (GV)
 - = prophase nucleus
 - ~30-40 μm
- nucleoli*
 - = "nucleolar-like body (NLB)", "nuclear remnant"
 - structure containing electron dense fibrilar/granular material
- during oocyte growth phase, chromosomes decondense and chromatin becomes transcriptionally active allowing for accumulation of cellular mass





Large scale chromatin remodelling



Large scale chromatin remodelling





 bidirectional GCs-oocyte comunication is necessary for timely coordination of chromatin decondensation and onset of transcriptional silencing





release from GV arrest



Kordowitzki et al 2021

Control of prophase arrest



* Oocyte incubation in cAMP analogs or PDE inhibitors prevents spontaneus maturation in vitro





He et al 2021





Visible sign of meiotic reactivation is

nuclear envelope breakdown (NEBD)
also known as GV breakdown (GVBD)
= meiotic diakinesis

Preceded by:

- chromatin congression
- relocation of nucleus towards oolema
- GV belt disapperance





GVBD



- GV collaps
- arrays of membrane fragments (annulate lamelae)
- organelles populate subcortical region

GV belt – subcortical region depleted from cellular organelles



Trebichalska et al 2021

Oocyte maturation



time

 hormonally-primed oocytes
 spontaneously mature in vitro when denuded from cumulus cells



Oocyte maturation



Nuclear maturation

- chromosomal segregation
- polar body (PB) extrusion
- Mll arrest
- 2n 4C oocyte \rightarrow 1n 2C egg

Cytoplasmic maturation

- structural and functional modification of organelles
- global changes in organelle arrangement
- mRNA translation and posttranslational modifications
- synthesis/degradation of maternal factors

Acquisition of fertilization and developmental competence

Nuclear maturation in mammalian oocytes

- (1) Nuclear envelope break-down
- Chromatin condensation and chromosome individualisation (2)
- Chromosome alignment at spindle (3)
- Homologous chromosome segregation (asymmetric cytokinesis) (4)
- MII spindle formation MII arrest (5)













10 µm





Schuh and Ellenberg, Cell 2007

~ 12 hours

 \sim 24 hours

Holubcova et al Science 2015

Nuclear maturation in human oocyte



Holubcova et al Science 2015

Nuclear maturation in human oocyte

Chromosome clustering

- transient stage of chromatin aggregation
- after NEBD and before onset of spindle assembly



Chromosomes





Melina Schuh





Chromosomes F-actin (EGFP-UtrCH) - actin cables invade disassembling nucleus and drive chromosome coalescence

Harasimov et al, Nature Cell Biology 2023

Nuclear maturation in human oocyte

Chromosome clustering

- promotes rapid capture of chromosomes by acentrosomal spindle and prevents chromosome losses in the long gap phase between nuclear envelope breakdown and the onset of spindle assembly





Melina Schuh



After NEBD

Microtubule "cage"

Chromosome segregation



- haploid set of chromosomes eliminates to 1st PB
- $2n 4C \rightarrow 1n 2C$

 $1n 2C \rightarrow 1n 1C (+ 1n 1C from sperm)$

Kinetochore-microtubule attachment

MITOSIS/ MEIOSIS II



attachment: kinetochores in relation to spindle poles

orientation: chromosome in relation to spindle poles



monotelic

amphitelic

sister kinetochores

bi-orientation

(monotelic) mono-orientation

both sister kinetochores one of the attached to the same pole



syntelic

(syntelic) mono-orientation





merotelic

(merotelic) bi-orientation











Loss of cohesion in anaphase I



Shugoshin (Sgo2)

recruits PP2A, which removes Rec8 phosphorylation, making it a poor substrate for separase-dependent cleavage.

Metaphase I

Anaphase I



- Stabilizes location of Sgo
- Ensures chromosome alignment in MII







Marson and Wassmann 2017





Chromosomes (H2B-mRFP) Microtubules MAP4-EGFP)

SAC in human oocytes is permissive

Bi-directional, no lagging



Bi-directional, with chromatin mass separation



B J

Bi-directional, with lagging

Tri-directional, with or without lagging



- misaligned chromosomes do not block anaphase I onset
 - Tripolar anaphase may result in
 - reunion of 2 chromosomas masses in the aneuploid oocyte, 3rd mass extruded to the PB
 - (1) re-joining of chromatin and cytokinesis failure (seemingly non-maturing MI !)

ORIGINAL ARTICLE Reproductive biology

Tri-directional anaphases as a novel chromosome segregation defect in human oocytes

Jenna Haverfield^{1,2}, Nicola L. Dean^{1,3}, Diana Nöel³, Gaudeline Rémillard-Labrosse¹, Veronique Paradis³, Isaac-Jacques Kadoch^{2,3}, and Greg FitzHarris^{1,2,8}

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- Bi-directional, no lagging
- Bi-directional, with lagging
- Bi-directional, with chromatin separation
- Tri-directional

A

Chromosomes re-joining after cytokinesis



Chromosomes re-joining in the absence of cytokinesis





SAC in human oocytes is permissive

>20-hour exposure



Induced DNA damage does not prevent anaphase entry in human oocytes

- Human oocyte with DNA damage harbour abnormal spindles but exhibit apparently normal morphology!

Human oocytes harboring damaged DNA can complete meiosis I

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Greg FitzHarris



Modes of SAC functionality

- in somatic cells, one unattached kinetochore capble to activate SAC¹
- DNA damage prevents entry to mitosis but not anaphase

- mouse oocytes with DNA damage undergo GVBD but arest in MI and fail to extrude PB²
- lack of response to unaligned chromosomes and lack of tension³

- in human oocytes, neither misaligned chromosomes nor DNA damage activate robust SAC response^{4,2}
- only severe spindle disruption can prevent PB
- SAC signalling machinery is present⁵

BUT inefficient to prevent anaphase onset when one or a few chromosomes are not congressed and attached to kinetochores

¹ Kuhn & Dumont, 2019; Rieder et al., 1994

² Remillard-Labrosse et al. 2019

³ Gui & Homer, 2012; Kolano et al., 2012; Lane et al., 2012; Nagaoka et al., 2011; Sebestova et al., 2012

⁴ Zielinska et al 2015, Havrfield et al. 2017

⁵ Lagirand-Cantaloube et al, 2017

Cytoplasmic volume affects SAC stringency

- cytoplasm scales the spindle and affects the timing of anaphase onset and efficiency of chromosome alignment
- oocytes with decreased cytoplasmic size have spindles with better-focused poles and higher SAC stringency
- large cytoplasmic volume dilutes the nuclear factors, including anaphase inhibitors, thus resulting in the failure of the spindle to induce a checkpoint arrest in response to a small number of misaligned chromosomes

Large Cytoplasm Is Linked to the Error-Prone Nature of Oocytes

Chromosome segregation erro

Chromosome segregation errors

Chromosome lagging

 delayed chromosome/chromatid movement during anaphase

 risk of inaccurate segregation, chromosome loss/gain and aneuploidy

No lagging

Class I laggards

Class II laggards

Transition from meosis I to meiosis II

Metaphase II arrest

Emi2 (Early Mitotic Inhibitor)
 meiosis specific inhibitor of APC (=,,cytostatic factor" - CSF)
 required for establishment and maintanance of MII arrest in mammalian oocytes
 phosphorylation needed to keep Emi2 stable
 Magwick et al 2006

Btg4

- contributes to APC/C inhibition by controling protein expression during MII arrest
- expression of Emi2 is perturbed when BTG is absent (RNAi depletion)

Pasternak et al 2016

- MII arrest reached hours before ovulation and maitained for ~ 24 hours

MII arrest release

Calmodulin dependent protein kinase II (CAMKII)

- activated by Ca2+ signal at fertilization
- phosphorylates Emi2 causing its degradation
- activates Week1B kinase that phosphorylates CDK1 contributing to its inactivation

Fertilization stage in different species

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e-Atlas klinické embryologie

- CZE/ENG e-learningová pomůcka
- veřejně dostupné po zadání hesla

= datum narození Luise Brown (DDMMYYYY)

Úvod

Tento atlas je e-learningovým materiálem primárně určený studentům výukového programu Embryolog ve zdravotnictví, Lékařské fakulty, Masarykovy univerzity.

Představuje kolekci dříve nepublikovaných fotografií a videí lidských oocytů a ranných embryí vzniklých v rámci procesu umělého oplození. Cílem této práce je seznámení s morfologickou variabilitou ženských pohlavních buněk a embryí a zachycení dynamiky preimplantačního vývoje.

Poděkování patří klinikám Reprofit International (Brno) a IVF clinic (Olomouc) za poskytnutí použitého obrazového materiálu.

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