

Embryologie I OOGENESIS

autumn 2023

Oocyte meiosis

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Entry to meiosis

„MITOTIC-TO-MEIOTIC TRANSITION“

Retinoic acid (RA)

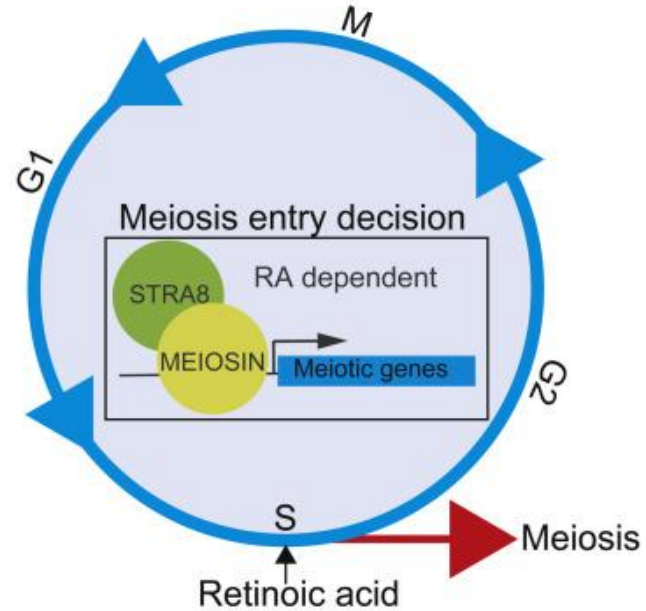


Stra8



meiosin

= meiosis initiator



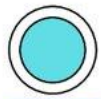
XX - prenatally
XY - postnatally

oogonium/spermatogonium

oocyt/spermatocyt

Mitóza

← Meiotic prophase → MI MII



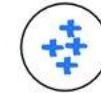
RA
STRA8

Pre-leptotene



RA
STRA8

MEIOSIN



AE formation

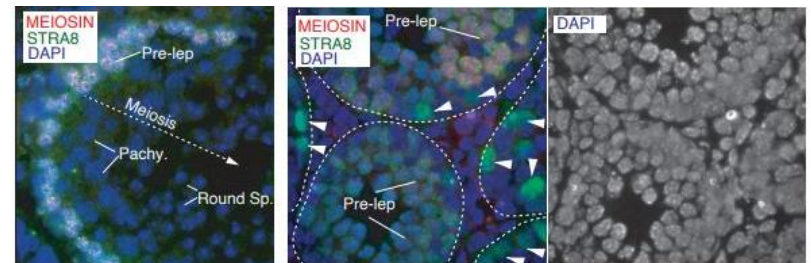
Meiotic recombination

Homolog synapsis

Telomere clustering

differentiation

Meiotic initiation
Pre-meiotic DNA
replication



Ishiguro et al 2020



Meiosis

(1) Premeiotic S phase

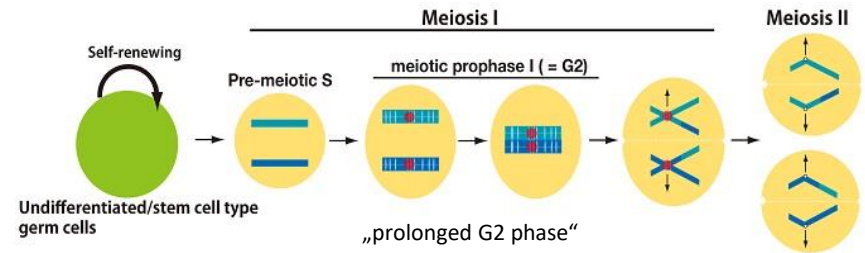
- DNA replication

(2) Meiosis I

- separation of homologous chromosomes

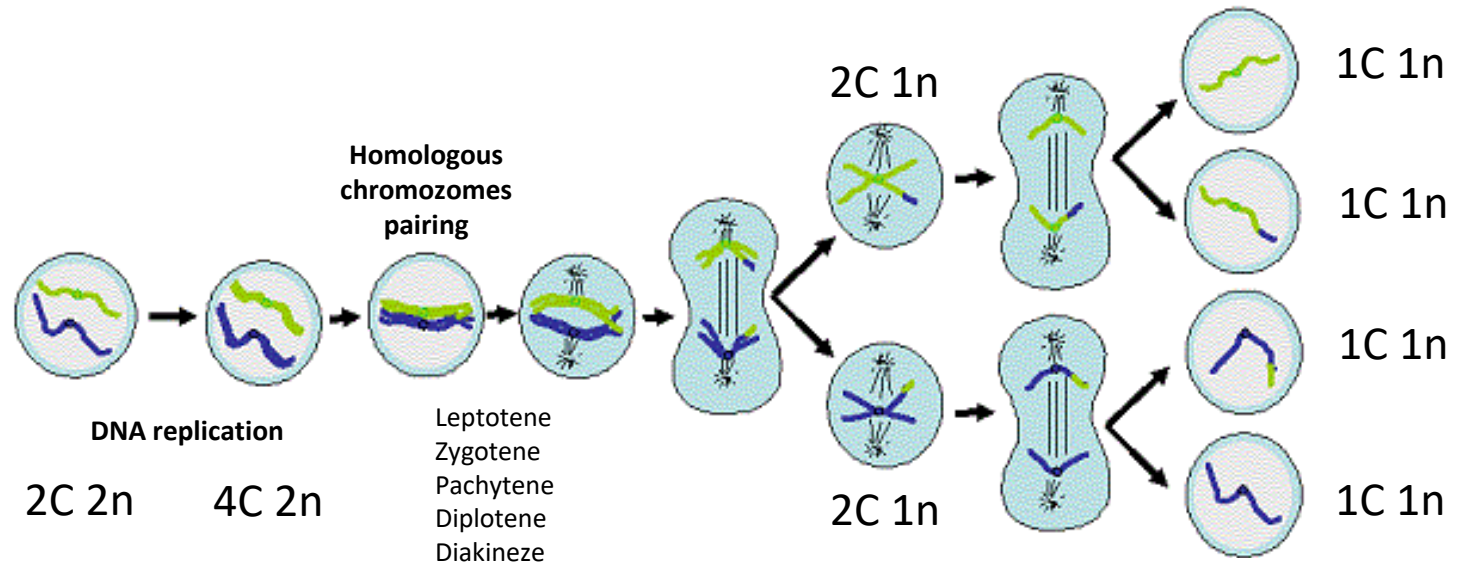
(3) Meiosis II

- separation of sister chromatids



Paternal chromosome

Maternal chromosome



I. Meiotic division

II. Meiotic division

Meióza



SPERMATOGENEZE

OOGENEZE



Mitóza

Spermatogonie
(2C 2n)

Meióza I

Primární
spermatocyt
(4C 2n)

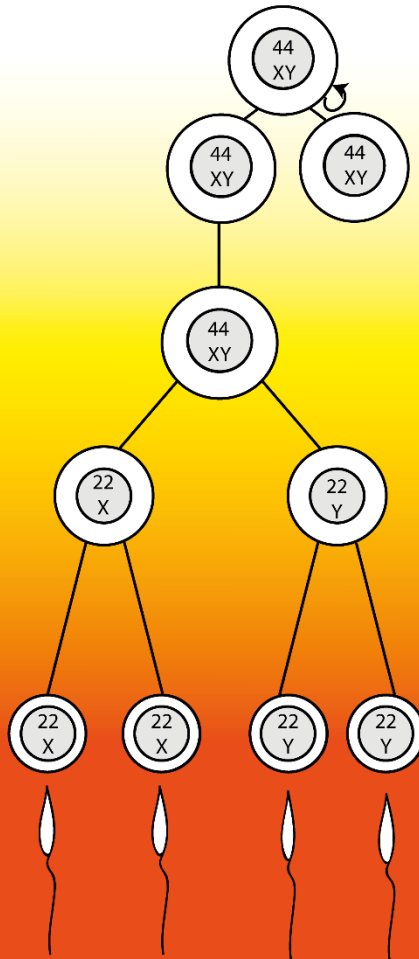
Sekundární
spermatocyt
(2C 1n)

Meióza II

Spermatidy
(1C 1n)

Spermiogeneze

Spermatozoa (1C 1n)



Primordiální zárodečná buňka
(2C 2n)

Mitóza

Oogonie
(2C 2n)

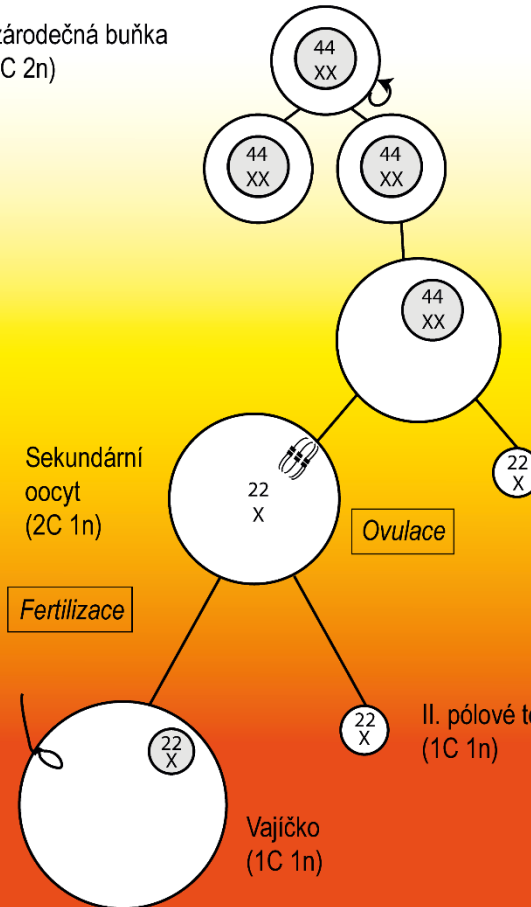
Meióza I

Primární
oocyt
(4C 2n)

I. pólóvé
tělísko
(2C 1n)

Meióza II

II. pólóvé tělísko
(1C 1n)



Sekundární
oocyt
(2C 1n)

Fertilizace

Ovulace

Vajíčko
(1C 1n)

n...počet sad chromozomů
C...počet kopií každého genu

Meiosis

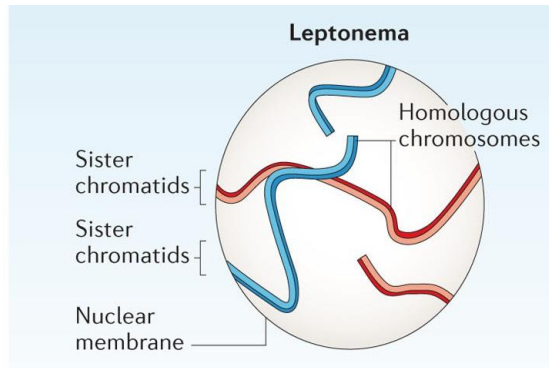
➤ Prophase I

(1) leptotene

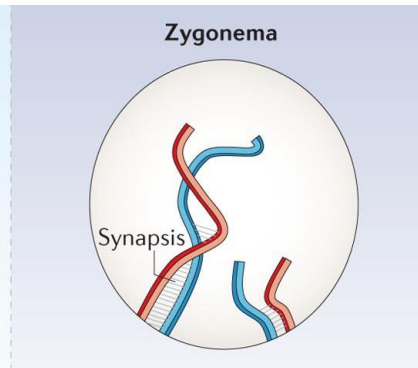
(2) zygotene

(3) pachytene

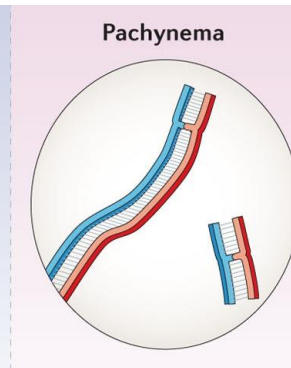
(4) diplotene



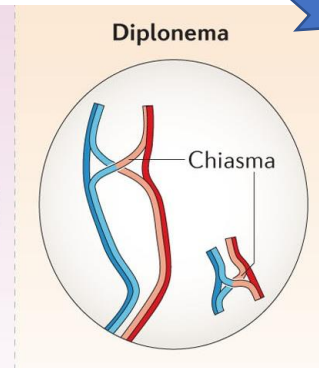
Condensation of replicated DNA



Synapsis of homologous chromosomes

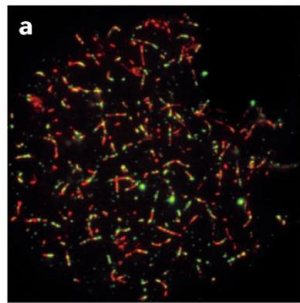


Recombination of homologous chromosomes
(crossing-over)

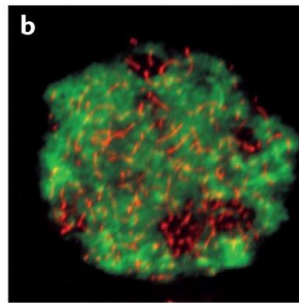


Synapsis disolution → bivalents with chiasmata

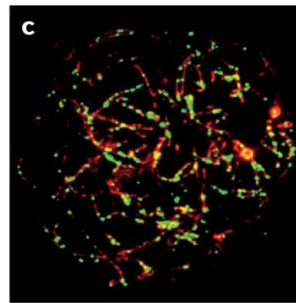
Prolonged arrest
= dictyotene



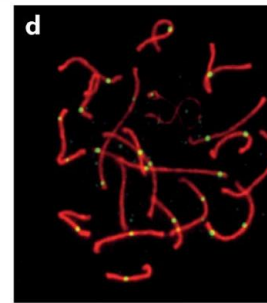
SYCP3 MEI4



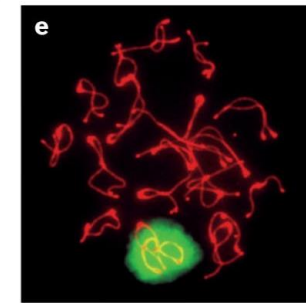
SYCP3 γ H2AX



SYCP3 DMC1 or RAD51



SYCP3 MLH1

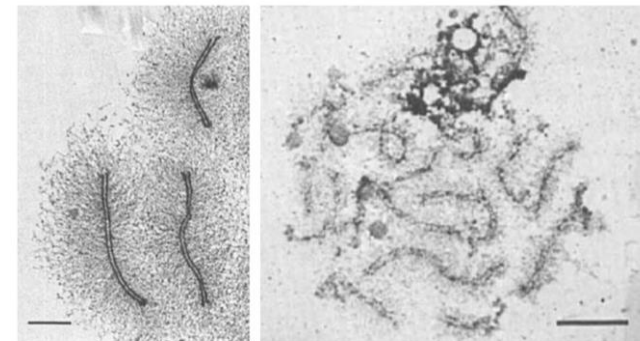
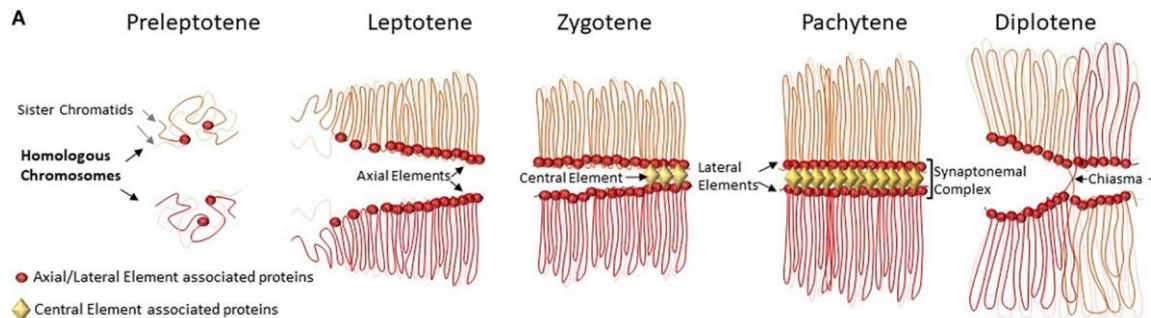
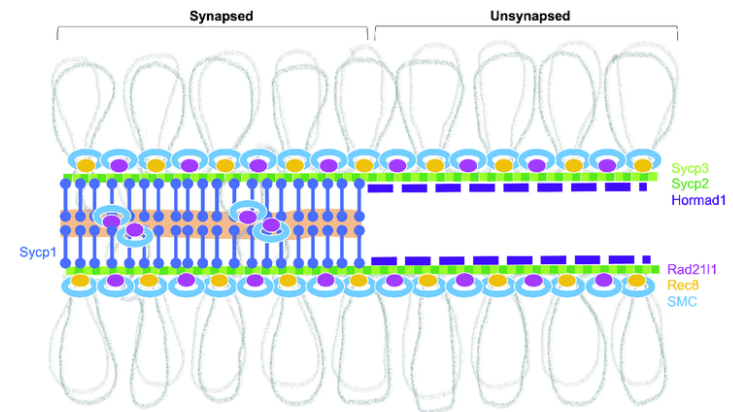
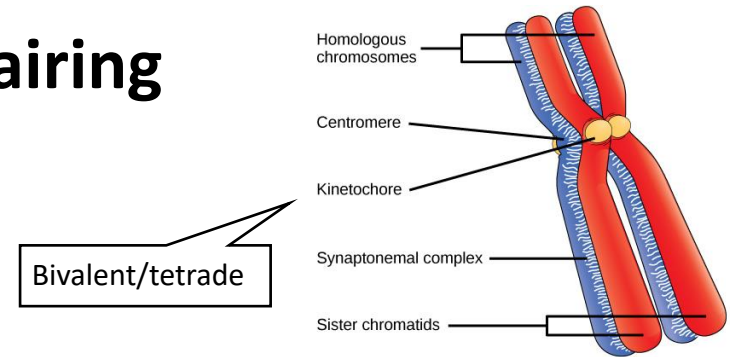


SYCP3 γ H2AX

Meiosis

➤ Homologous chromosomes pairing

- **synapsis** = physical association of **homologous chromosomes**
- homologous sequence pairing
- **Synaptonemal complex**
 - axial element - **SYCP3+SYCP2**
 - central element - dimer **SYCP1**
- gradual formation of SYCP3 and 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

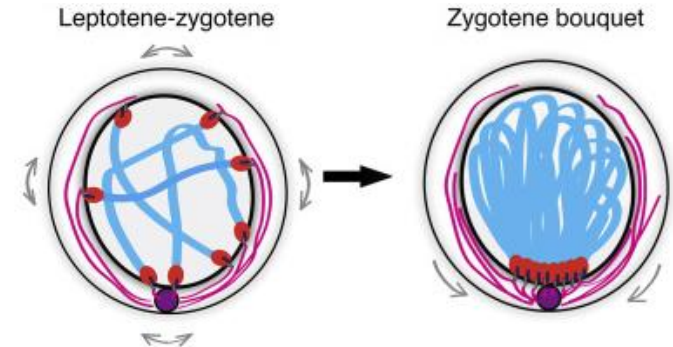


Meiosis

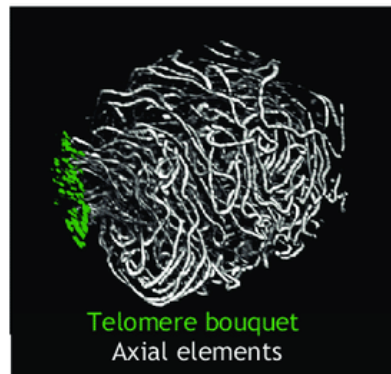
➤ Homologous chromosomes pairing

- pairing of distant homologous chromosomes accomplished by congregation of telomeres attached to nuclear membrane
→ tzv. „telomere bouquet“

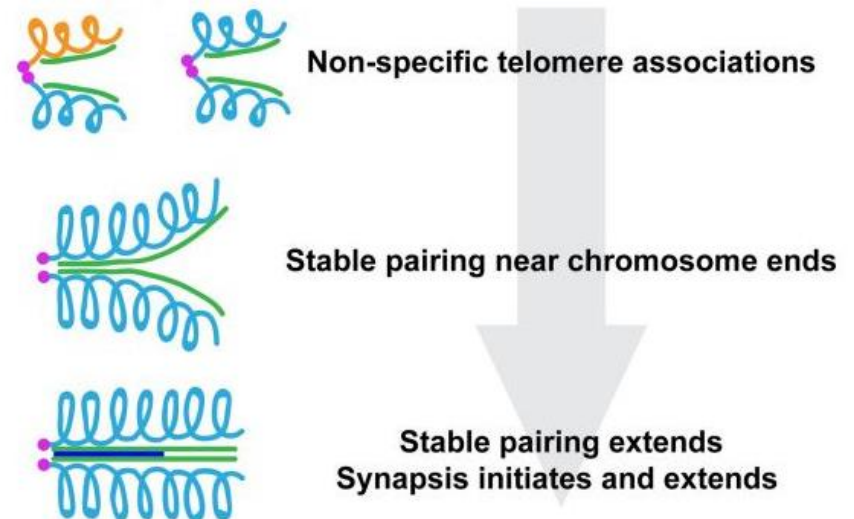
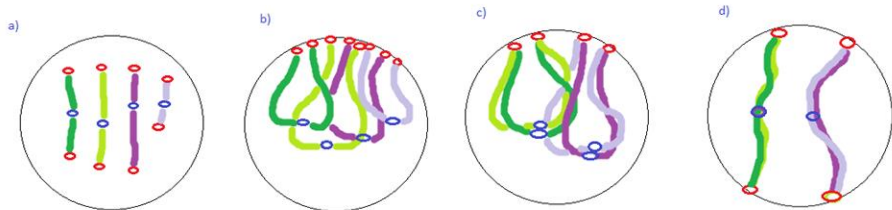
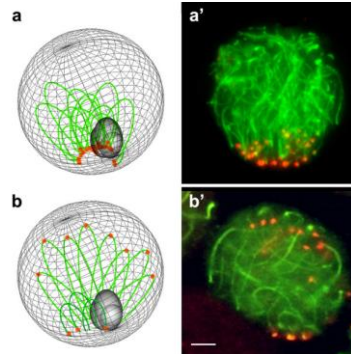
Elkouby and Mullins 2017



Unc84a in mammals *Zhou et al 2012*

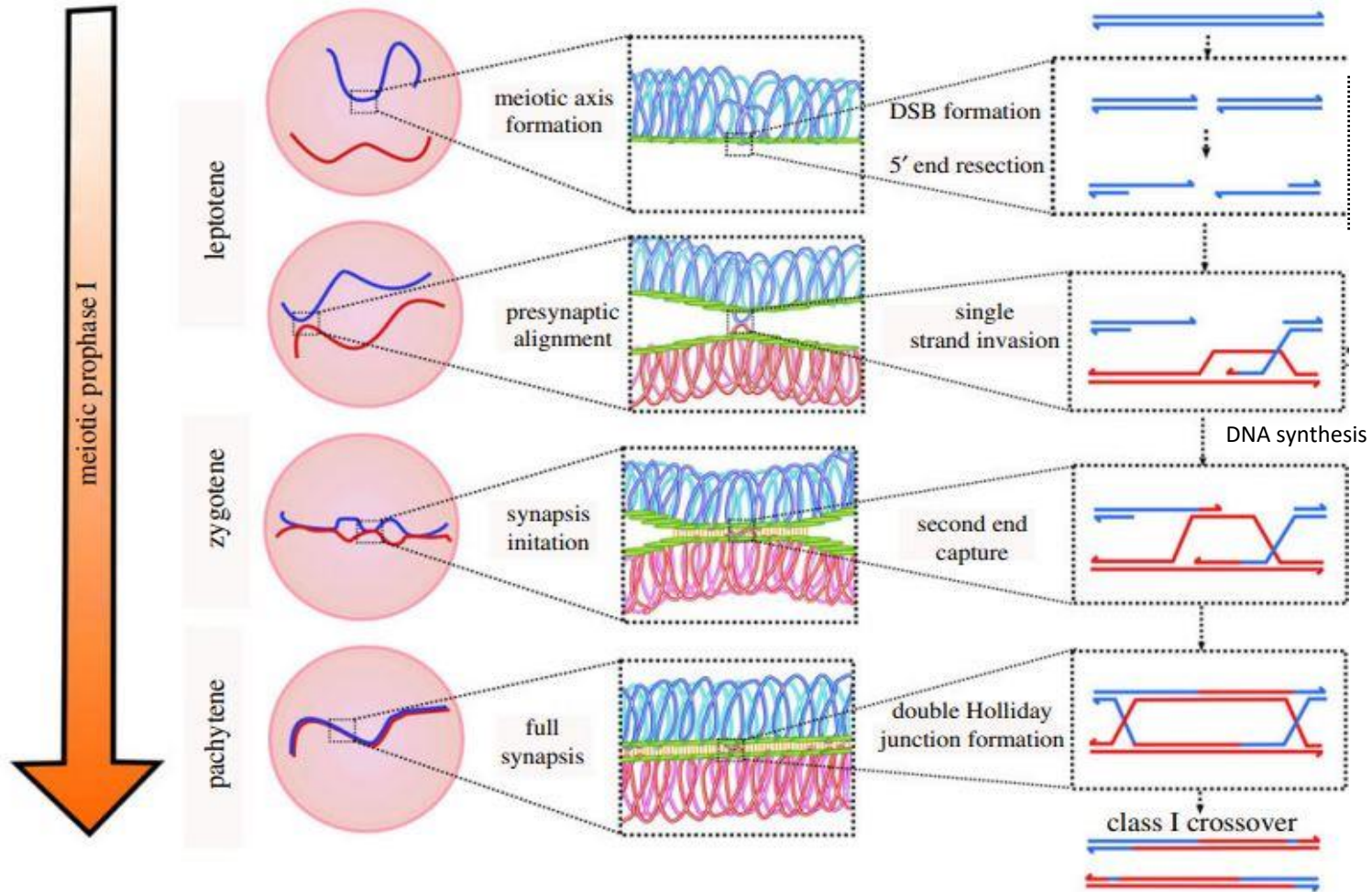


Berrios et al 2013



Meiosis

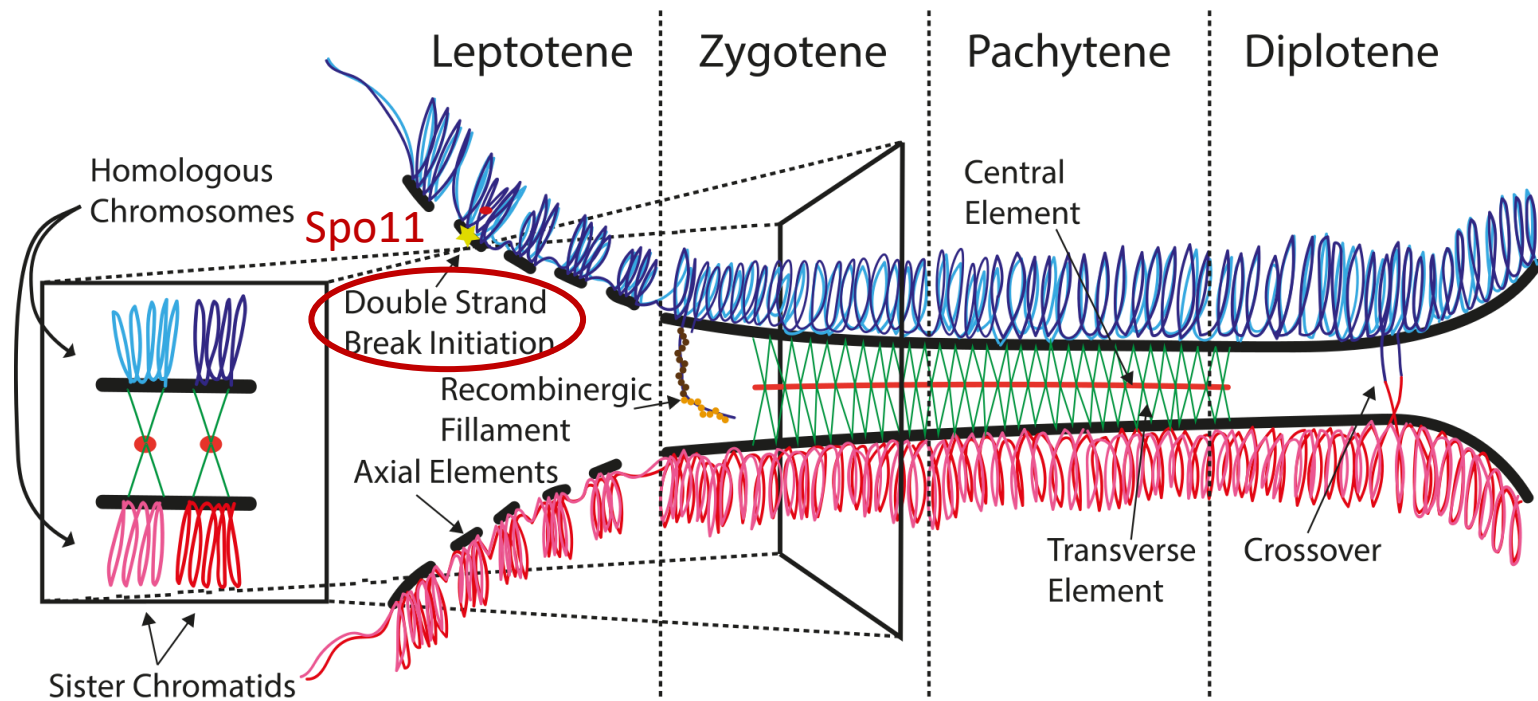
➤ Homologous recombination



Meiosis

➤ Homologous recombination

- DNA **double strand breaks (DSB)** precede synapsis formation



Meiosis

➤ Homologous recombination

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

- **Spo11**
 - DNA endonuklease
 - its dimer forms DSB
 - each monomer then binds 5' end of ssDNA
 - Spo11+oligonucleotide cleaved away by exonuclease → free 3' end



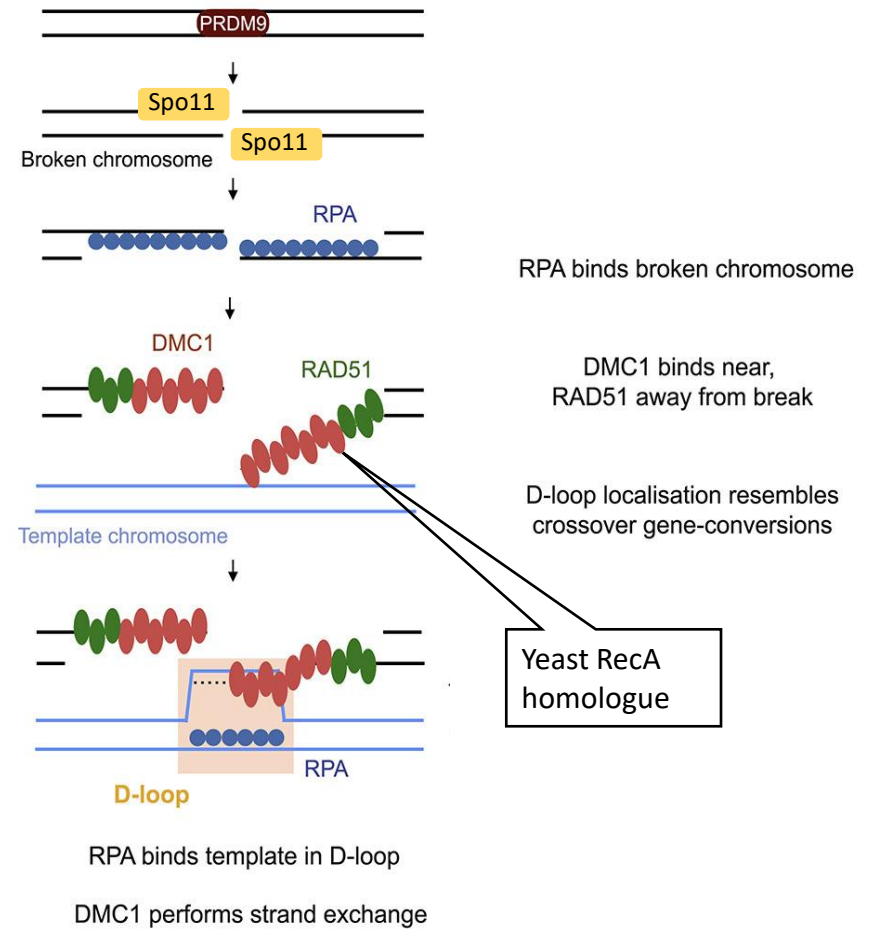
- mouse Spo11^{-/-} males sterile
- Spo11 polymorphism in male infertility



Meiosis

➤ 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



Meiosis

➤ Homologous recombination

- DNA synthesis of DNA 3' end using non-sister chromatid as a template

+ strand ligation



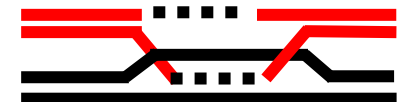
Synthesis dependent strand annealing (SDSA)



Single end invasion (SEI)



Double strand break repair (DSBR)

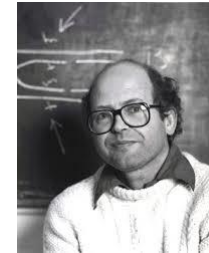


Meiosis

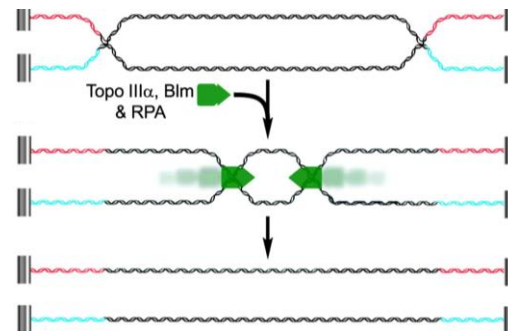
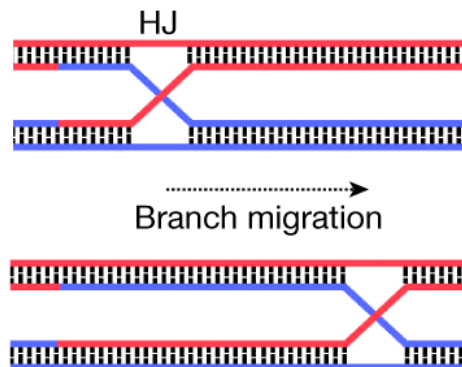
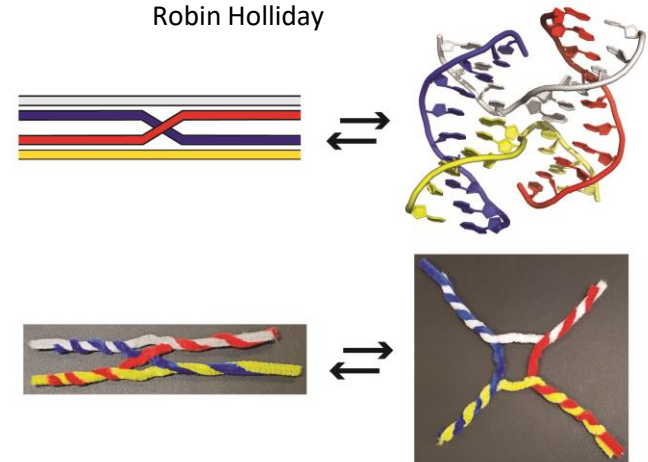
➤ Homologous recombination

▪ 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



Robin Holliday



Meiosis

➤ Homologous recombination

DSB



non-crossover ~ 90

- no gene conversion

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

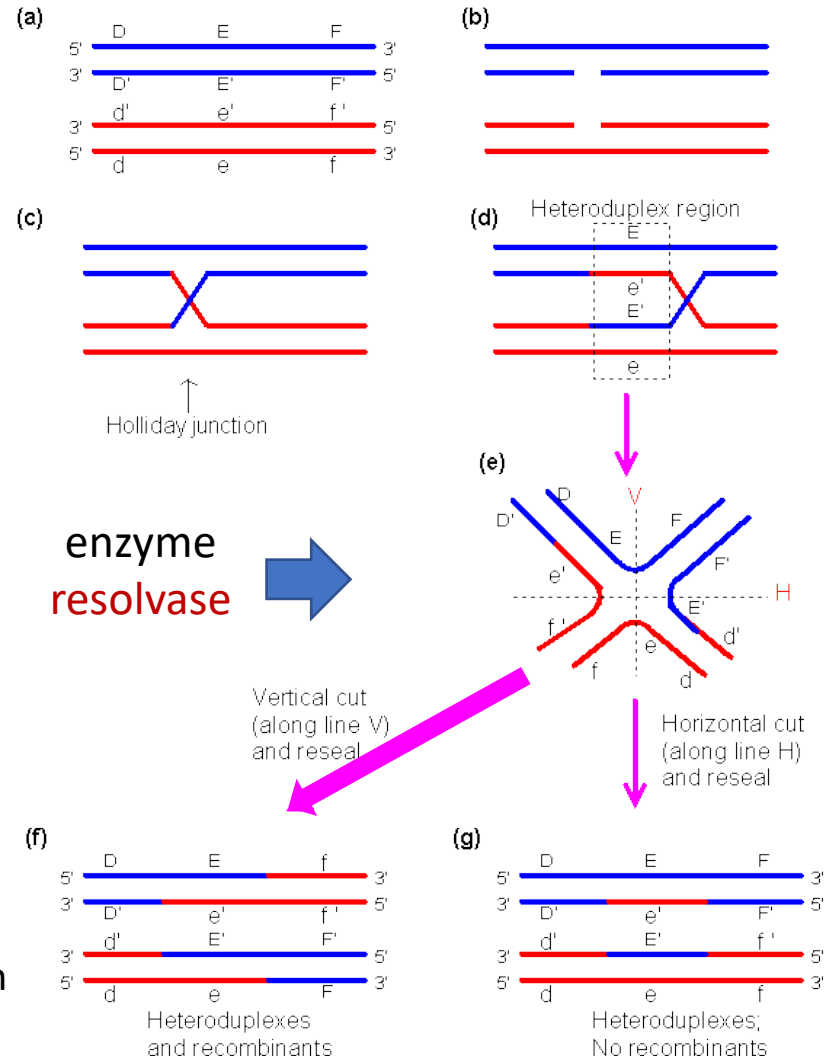
(b) convergent branch migration resulting in resolution of HJs

(c) strand exchange and HJ resolution without gene conversion



crossover ~10%

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

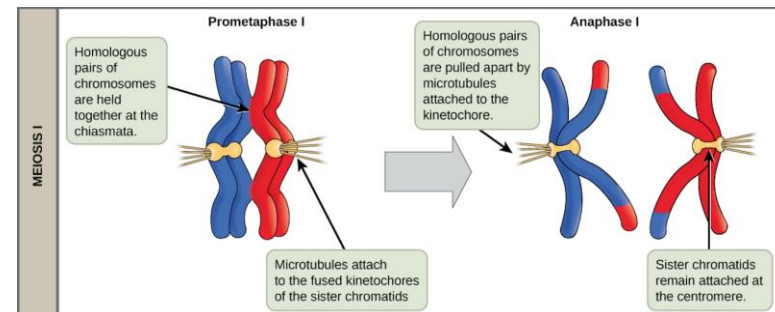
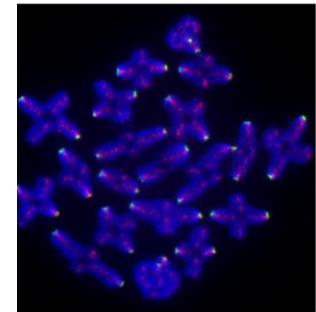
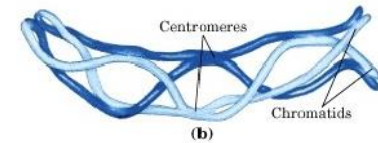
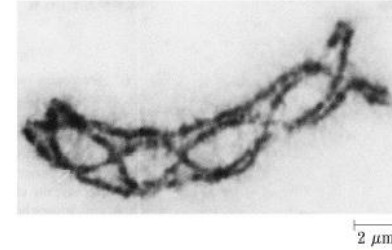
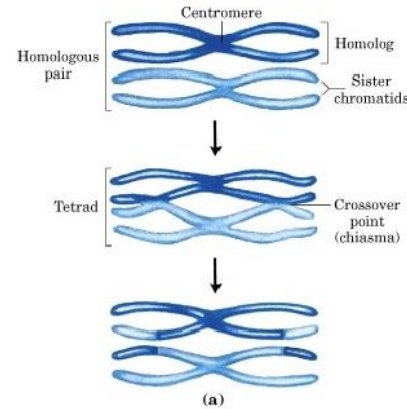


Meiosis

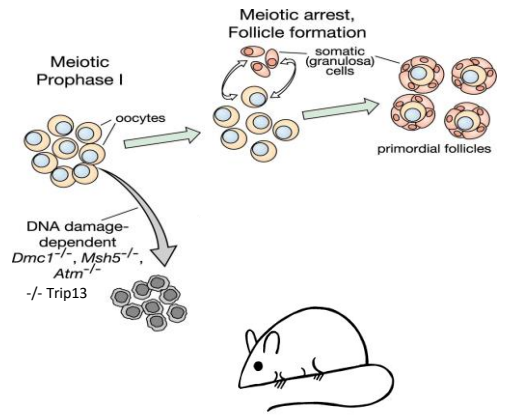
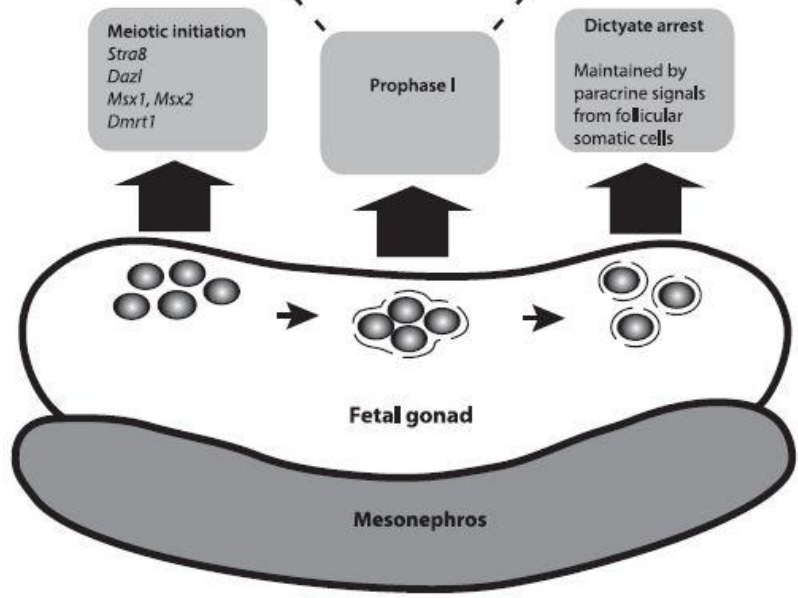
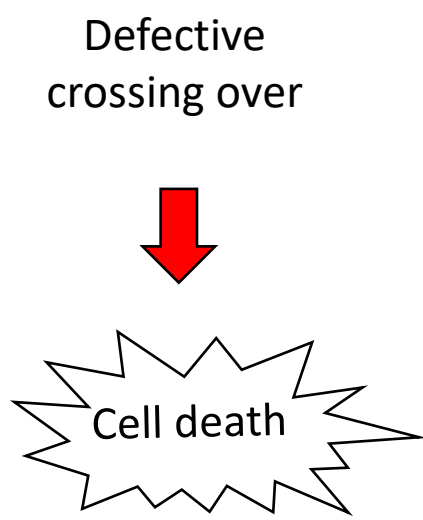
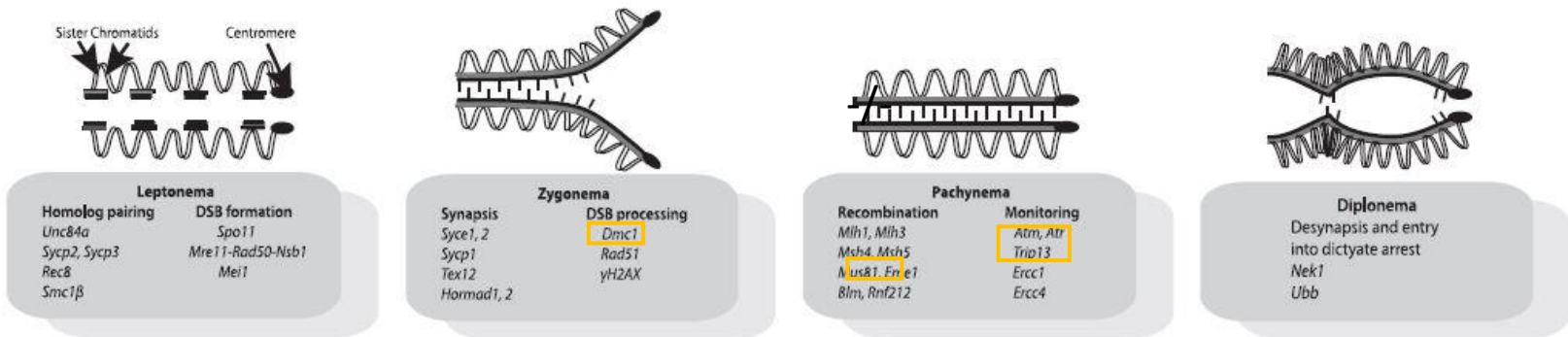
➤ Homologous recombination

■ Chiasmata (chi-structure)

- physical contact sites of homologous chromosomes marking crossing-over regions
- visible after synaptonemal complex dissolution during diplotene
- links homologous chromosomes together in the form of bivalents (tetrads)
- disappear 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



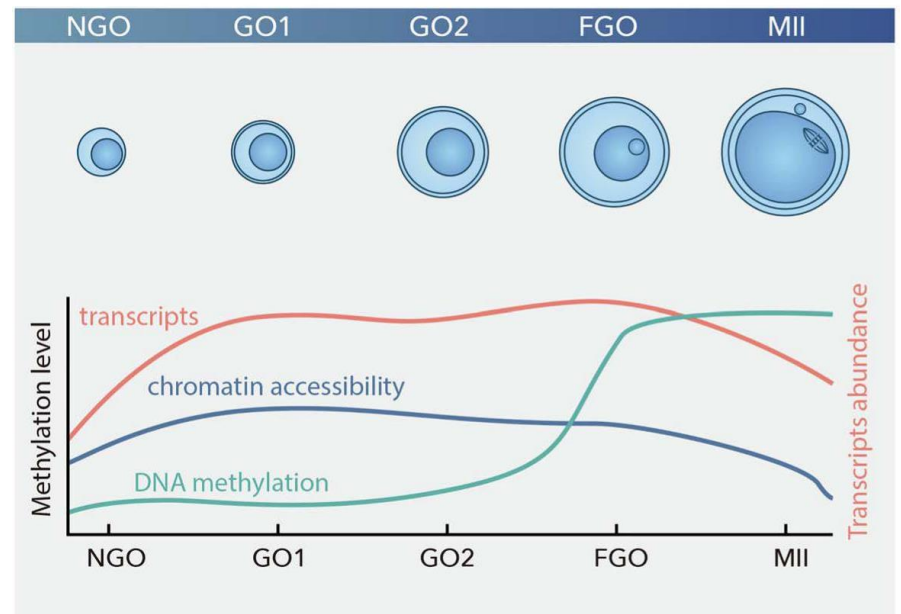
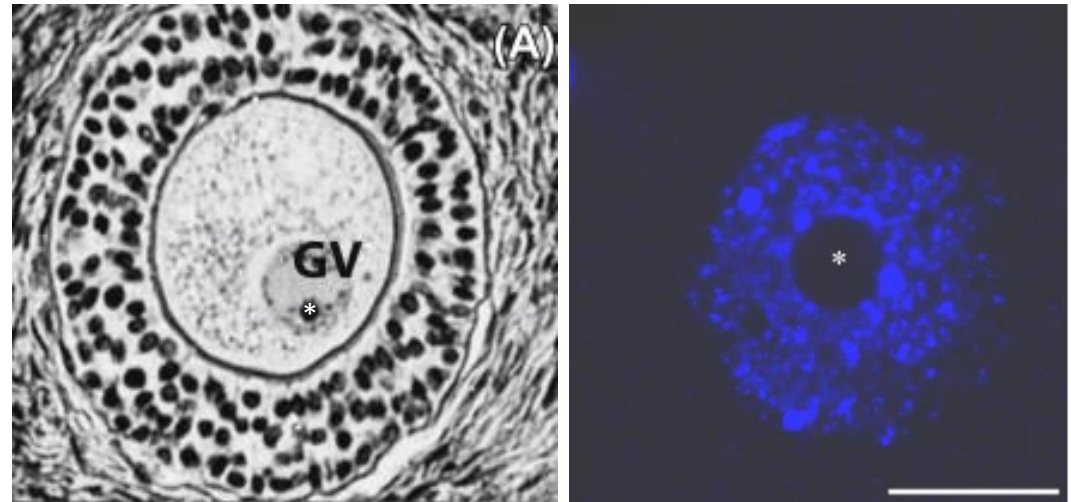
Regulation of meiotic prophase overview



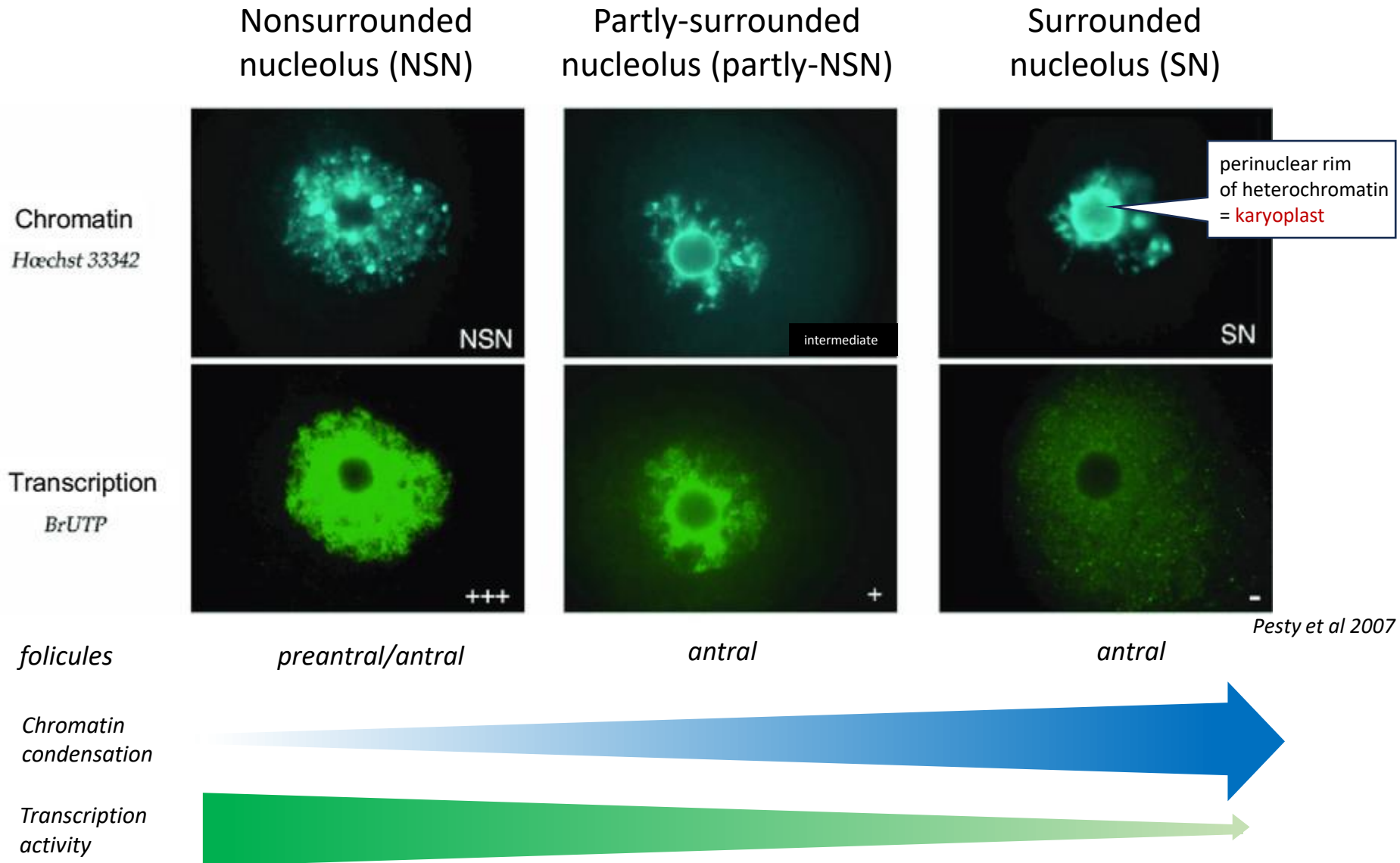
- devoid of primordial follicles
- infertile

Chromatin configuration during diplotene arrest

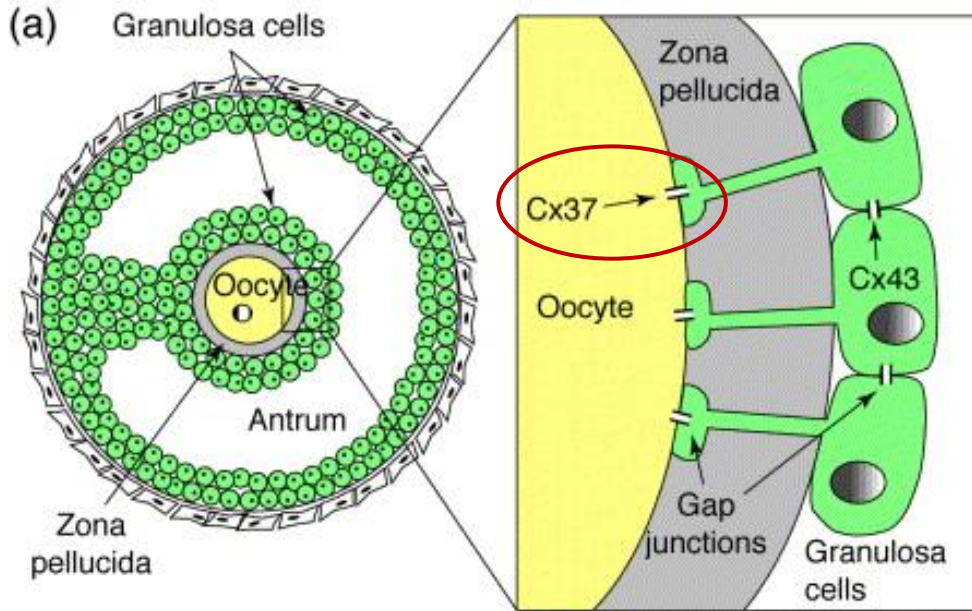
- **dictyate** stage = prolonged diplotene arrest
- chromosomes become dispersed, less distinct and form faint network
- **germinal vesicle (GV)**
 - = prophase nucleus
 - ~30-40 μm
- **nucleoli***
 - = „nucleolar-like body (NLB)“, „nuclear remnant“
 - structure containing electron dense fibrillar/granular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



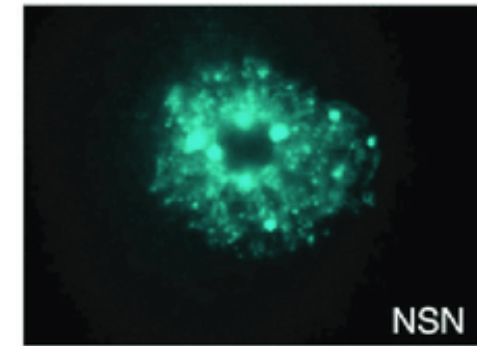
Large scale chromatin remodelling



Large scale chromatin remodelling

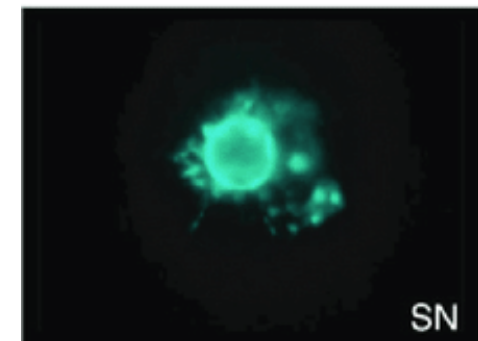


- bidirectional GCs-oocyte communication is necessary for **timely** coordination of chromatin decondensation and onset of transcriptional silencing

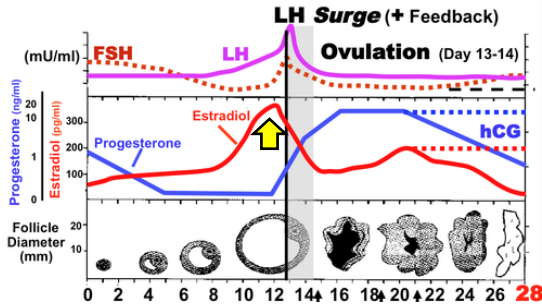


FSH →
(+COC in IVM)

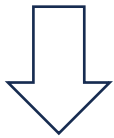
⊥  -/- Connexin37



Resumption of meiosis



↑ estrogen



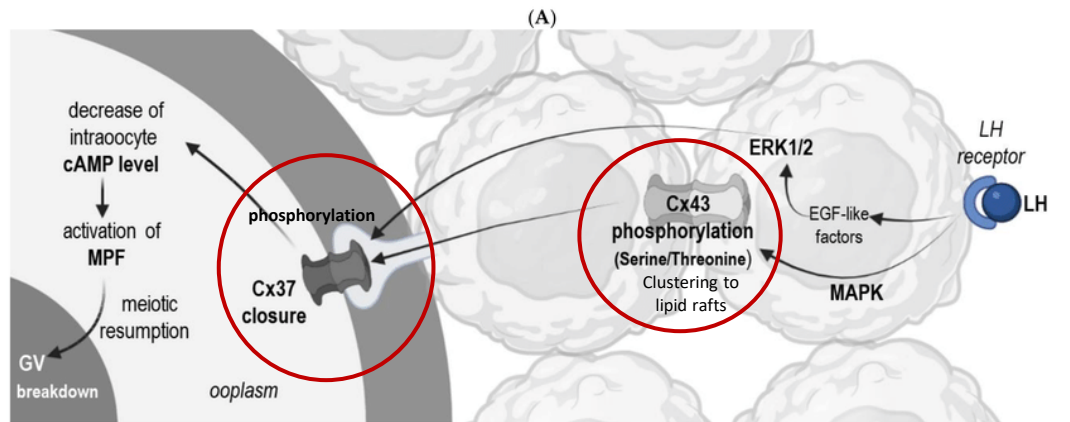
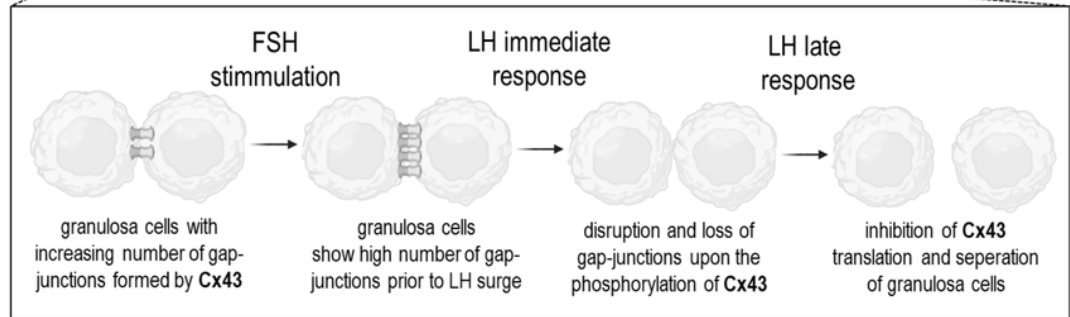
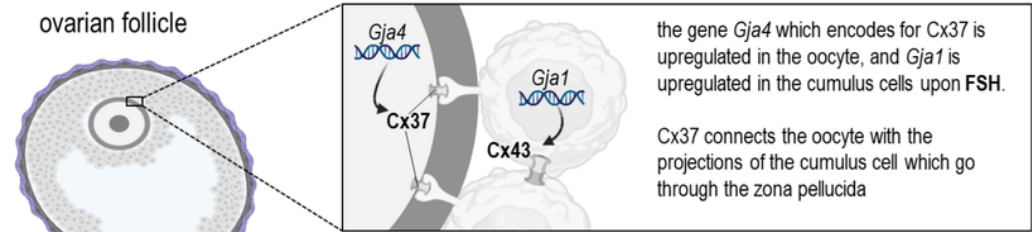
LH surge



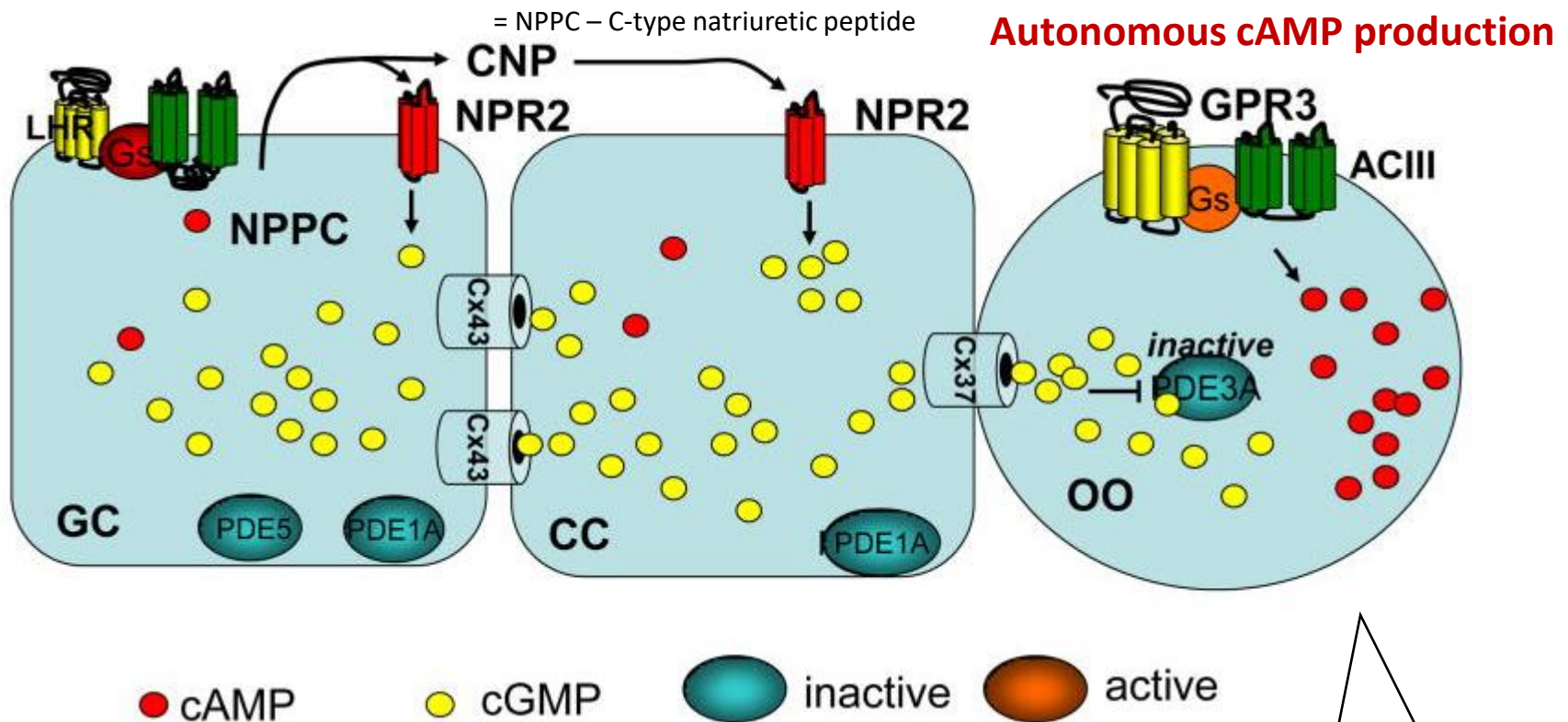
oocyte-cGCs uncoupling



release from GV arrest



Control of prophase arrest



Conti et al 2011

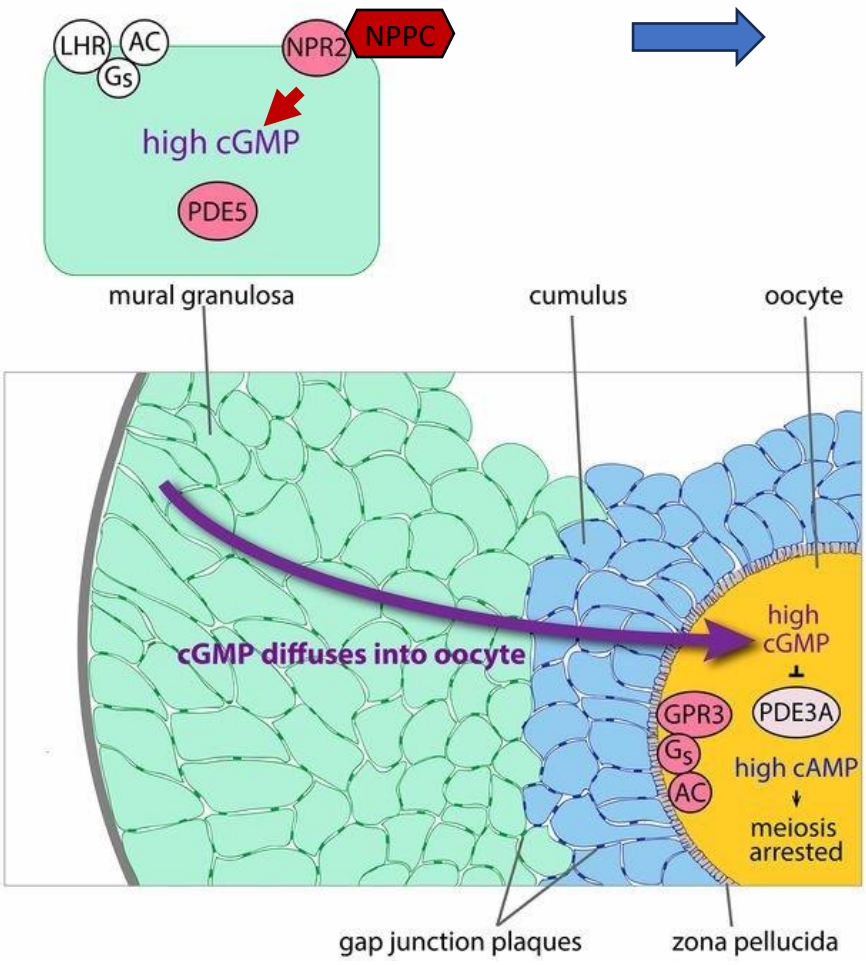
Supplementation of cGMP by cGCs balances cAMP degradation by PDE

Prophase arrest is dependent on high levels of cAMP in ooplasm

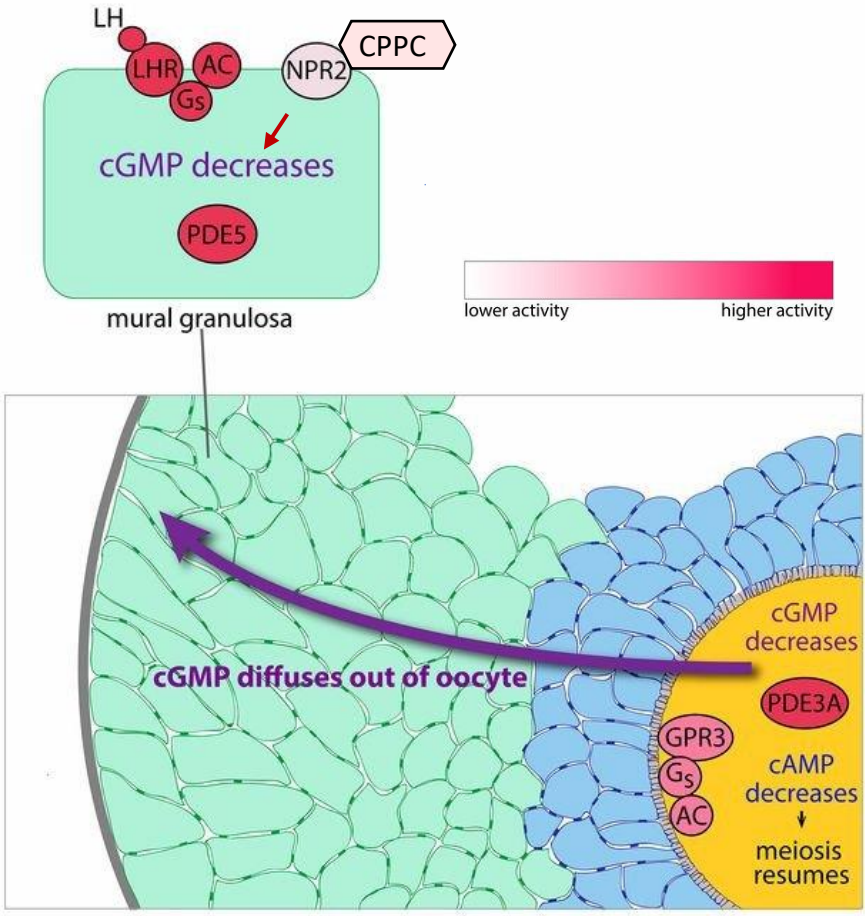
* Oocyte incubation in cAMP analogs or PDE inhibitors prevents spontaneous maturation in vitro

Resumption of meiosis

A Before Luteinizing Hormone

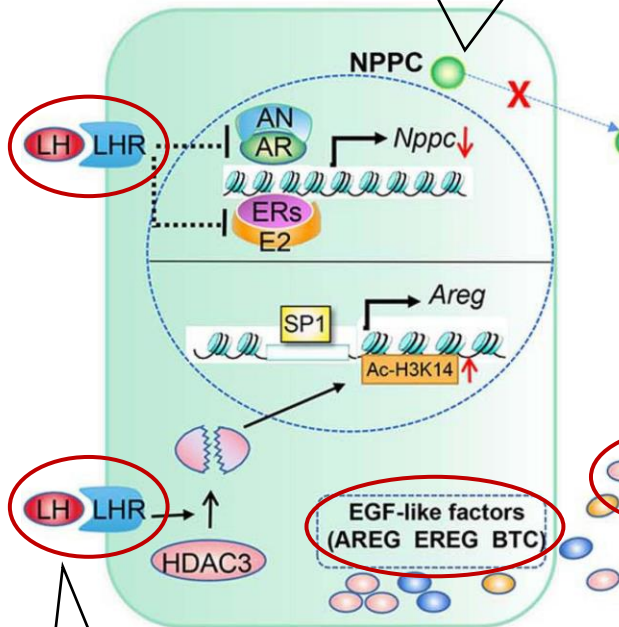


B After Luteinizing Hormone



Resumption of meiosis

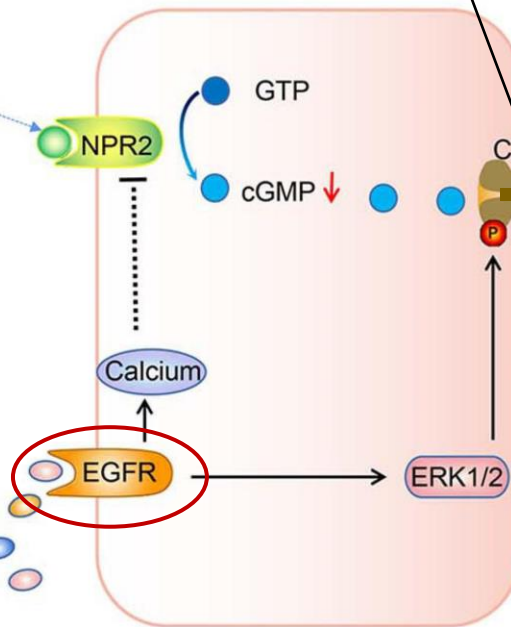
Decrease of NPPC production (1)



Mural granulosa cells

Oocyte and cumulus GCs in preovulatory follicles lack LHR!

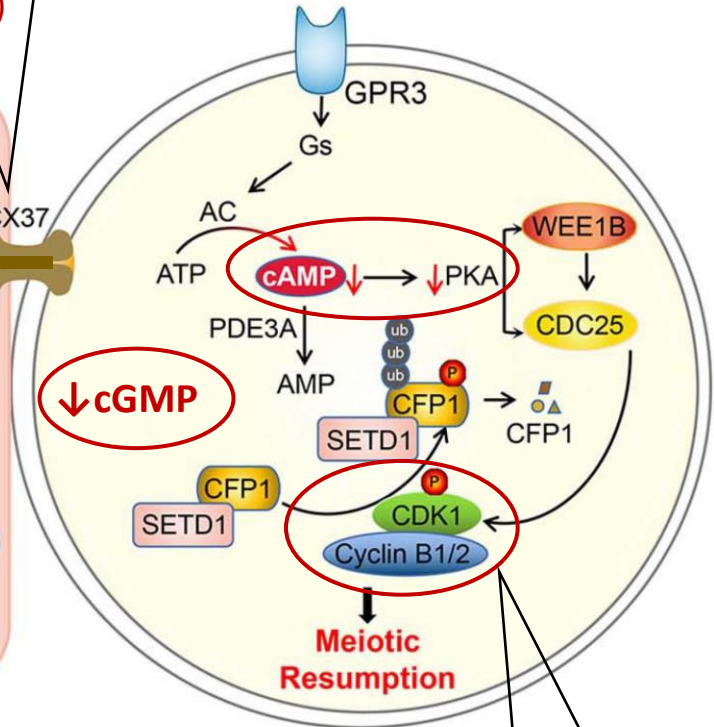
Gap junction closure (2)



Cumulus granulosa cells

LH surge (/hCG trigger)

- 1) ↓ cGMP synthesis in cGC
- 2) ↓ diffusion of cGMP from cGC to the oocyte



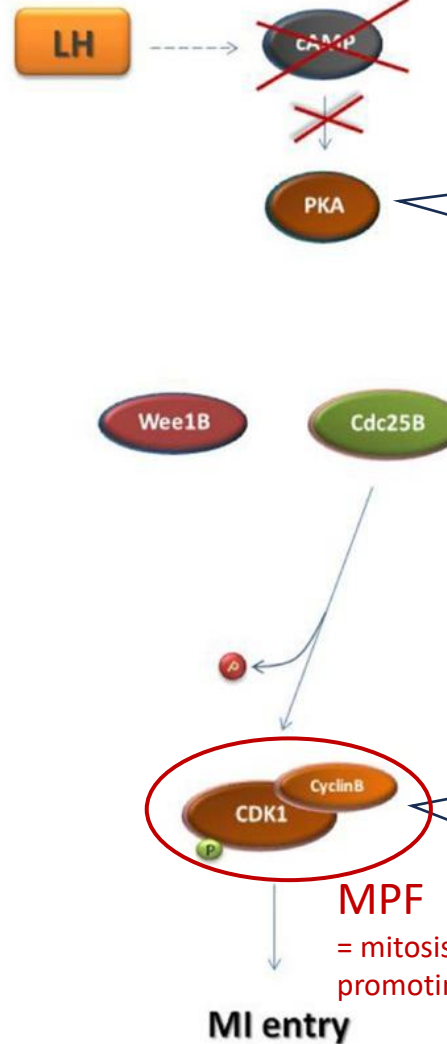
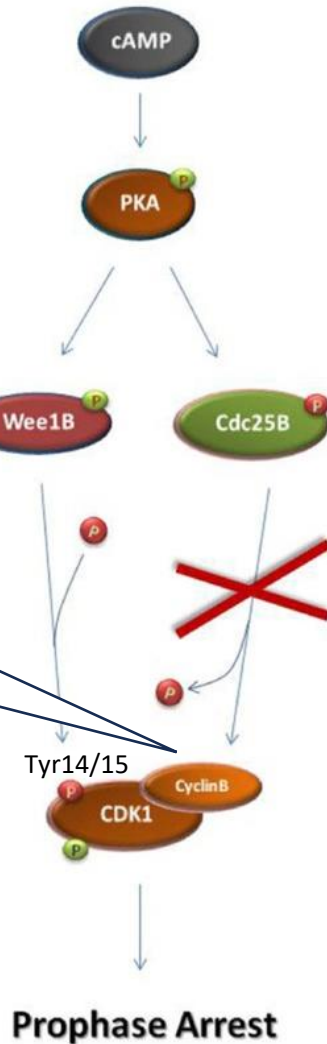
Oocyte

Decrease of cAMP and PKA activity leads to activation of meiosis promoting complex (MPF) and resumption of meiosis

Resumption of meiosis

Active Wee2 and inactive Cdc25B maintain prophase I arrest

- cycB accumulated during oocyte growth
- mouse oocytes with 80% of full size (60-65µm) have sufficient stockpile to resume meiosis



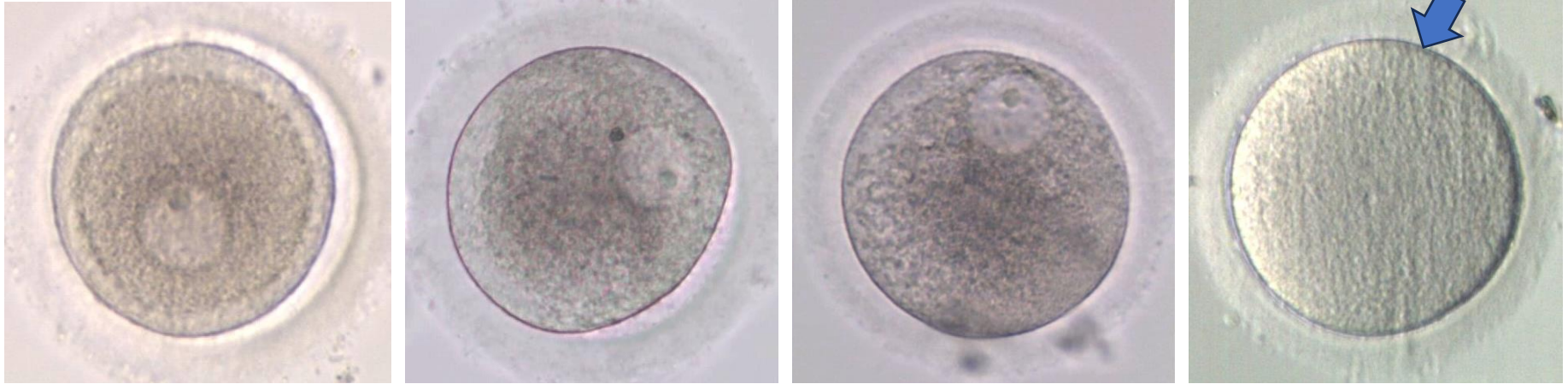
LH-induced drop of cAMP level leads to PKA inactivation

Active cdc25B removes inhibitory phosphate from CDK1

MPF
= mitosis/meiosis promoting factor

- in 3-4 hours in mouse oocytes
- In 12 hours in human oocytes

Resumption of meiosis



Visible sign of meiotic reactivation is

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

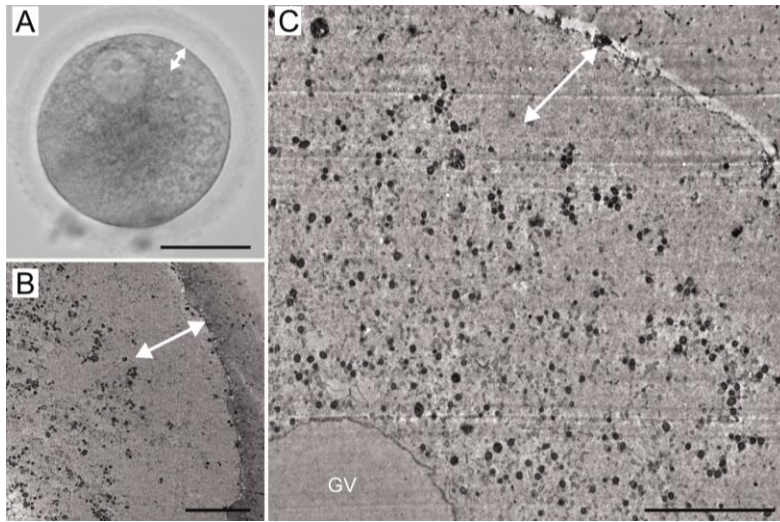
Preceded by:

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



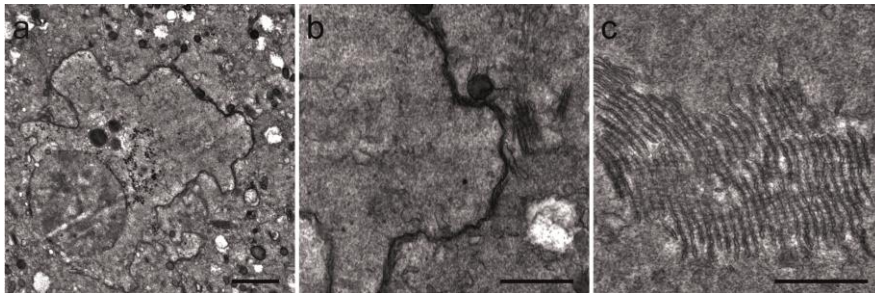
Resumption of meiosis

GV

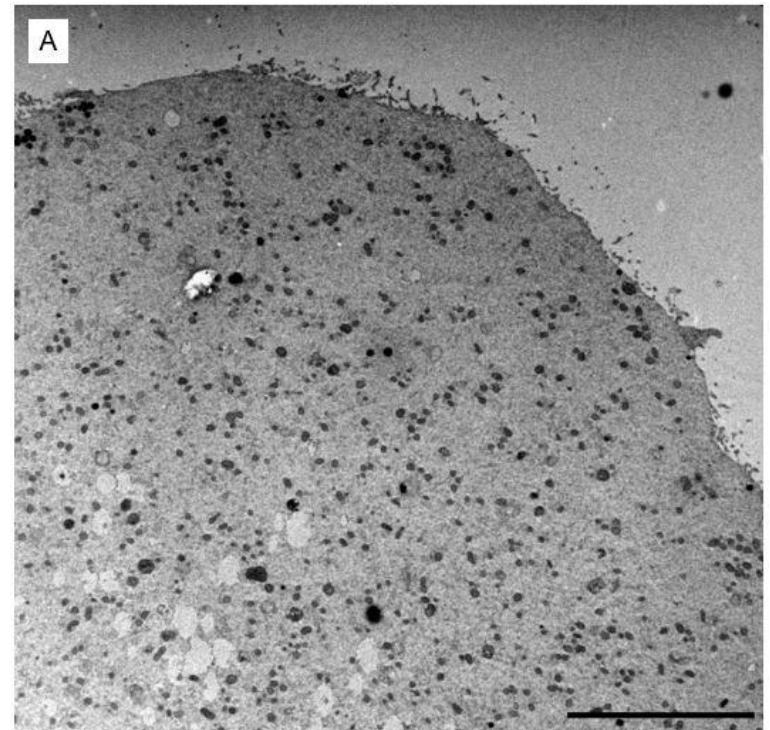


GV belt – subcortical region depleted from cellular organelles

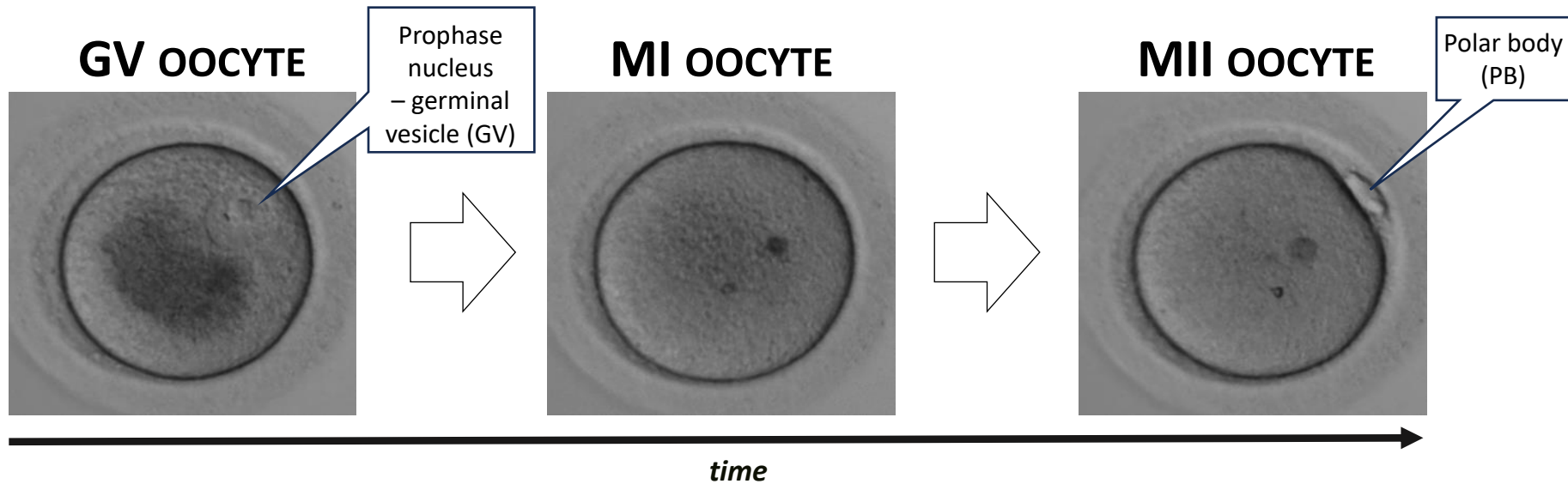
GVBD



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



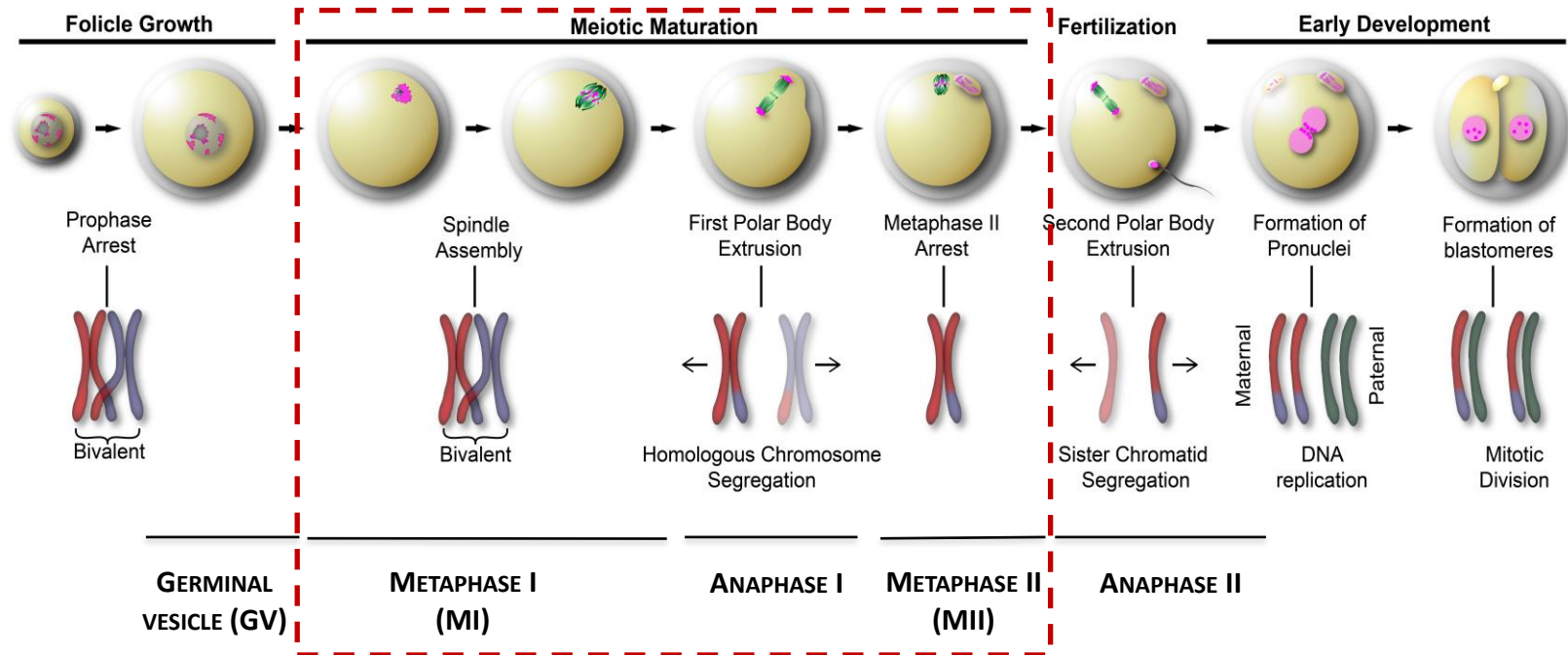
Oocyte maturation



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



Oocyte maturation

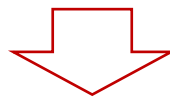


➤ Nuclear maturation

- chromosomal segregation
- polar body (PB) extrusion
- MII 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

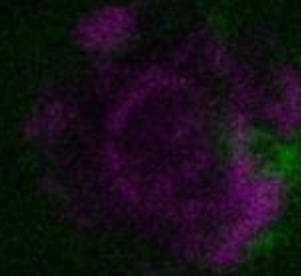
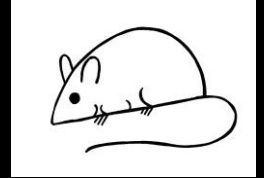


Acquisition of fertilization and developmental competence

Nuclear maturation in mouse oocytes

- (1) Initiation of microtubule nucleation
- (2) Nuclear envelope break-down
- (3) Chromatin condensation and chromosome individualisation
- (4) Spindle build-up and bipolarization
- (5) Chromosome alignment
- (6) Spindle relocation to cortex
- (7) Homologous chromosome segregation (asymmetric cytokinesis)
- (8) MII spindle formation – MII arrest

MAP4 – microtubules
H2B - DNA



~ 12 hours

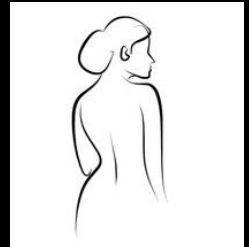


0.00 h

Nuclear maturation in human oocyte

- (1) Nuclear envelope break-down
- (2) Chromatin condensation and clustering
- (3) Initiation of microtubule nucleation
- (4) Chromosome individualisation
- (5) Spindle build-up and remodelling
- (6) Spindle bipolarization and chromosome alignment
- (7) Homologous chromosome segregation (asymmetric cytokinesis)
- (8) MII spindle formation – MII arrest

MAP4 – microtubules
H2B - DNA

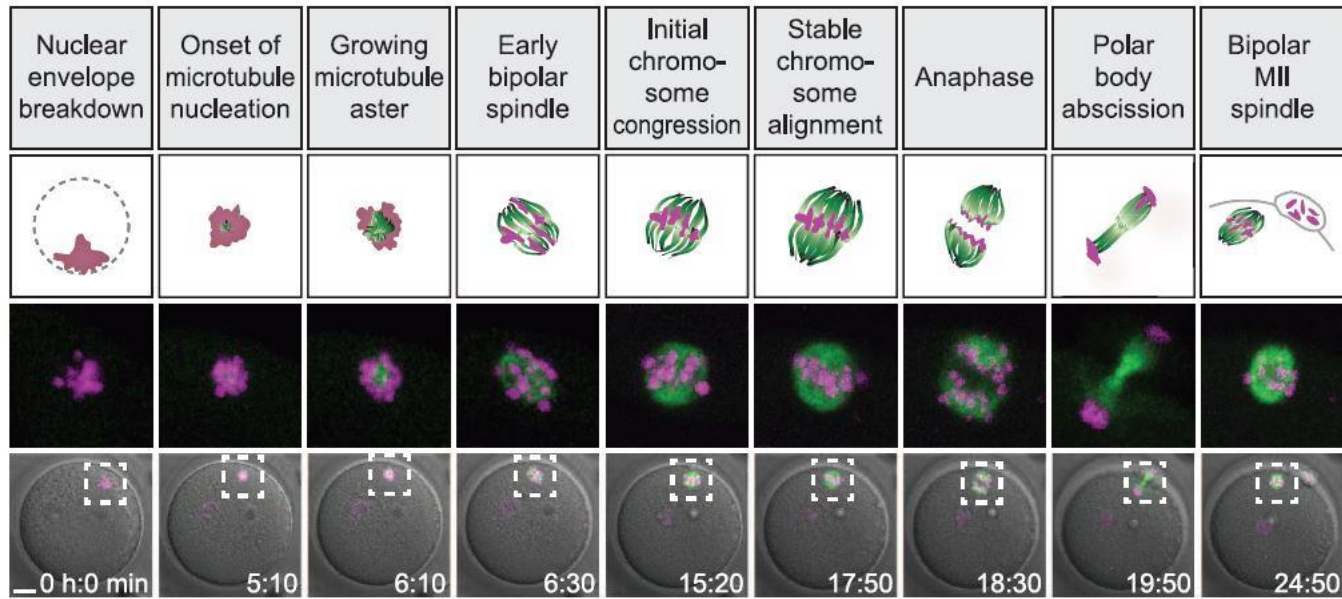


~ 24 hours

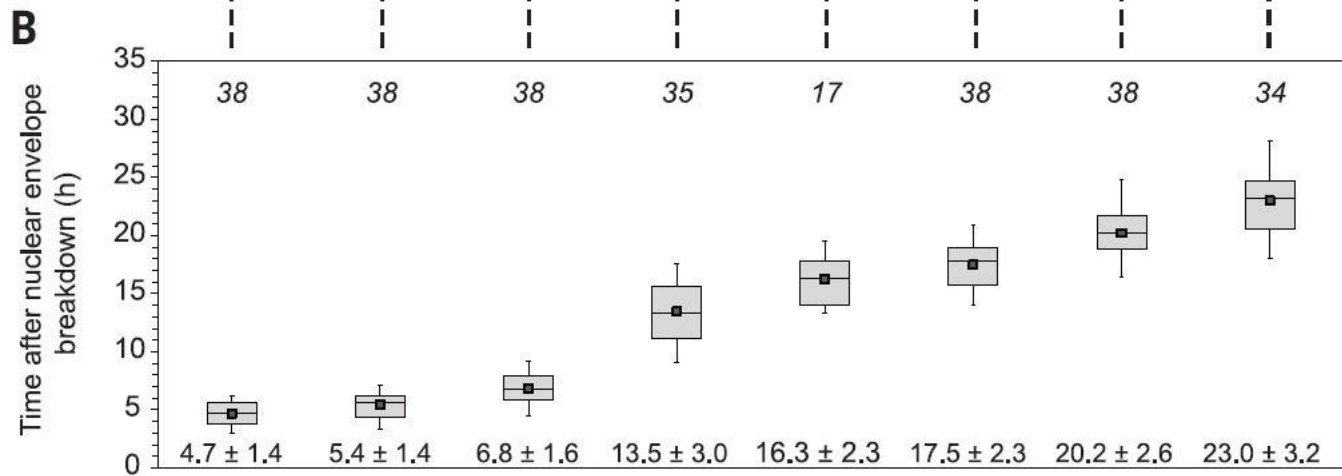


0.00 h

Nuclear maturation in human oocyte



■ Chromosomes
■ Microtubules



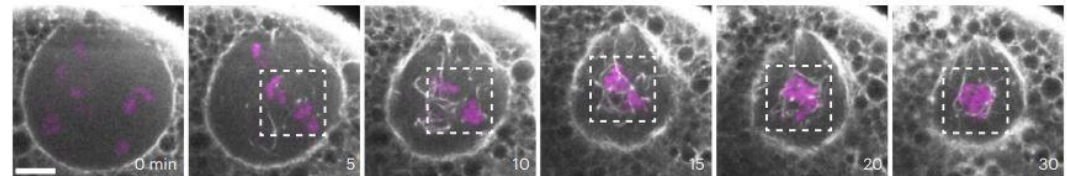
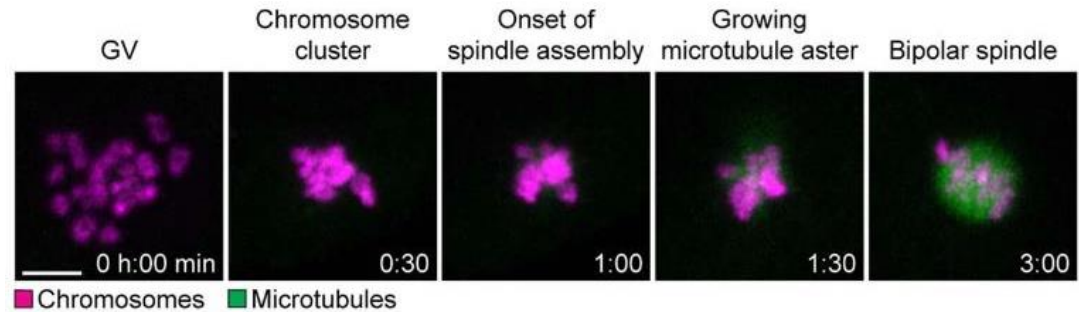
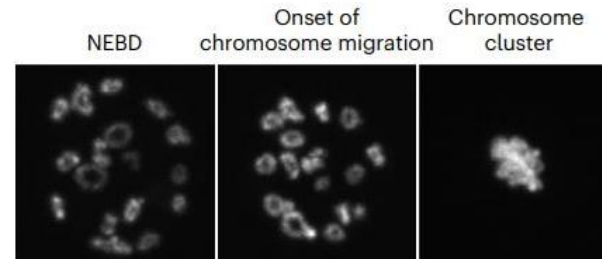
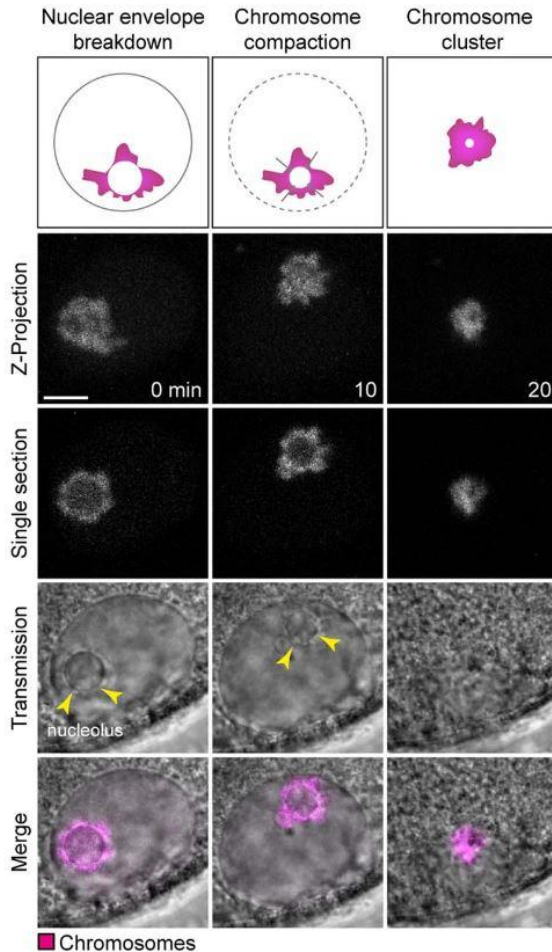
Nuclear maturation in human oocyte

❖ Chromosome clustering

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



Melina Schuh



- actin cables invade disassembling nucleus and drive chromosome coalescence

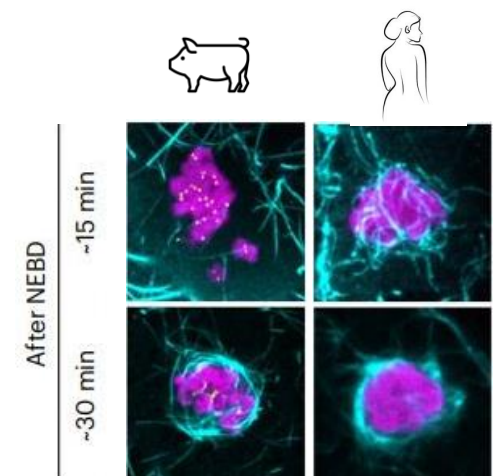
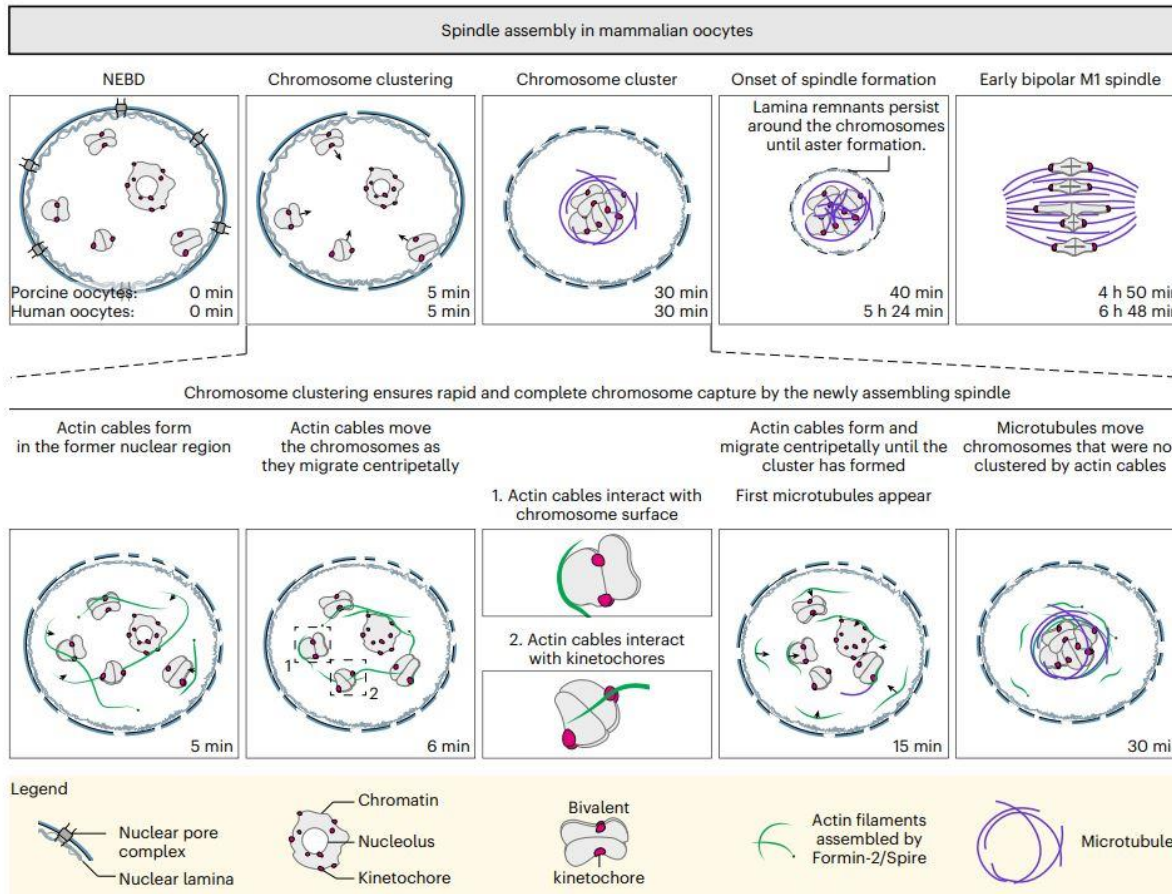
Nuclear maturation in human oocyte

❖ 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

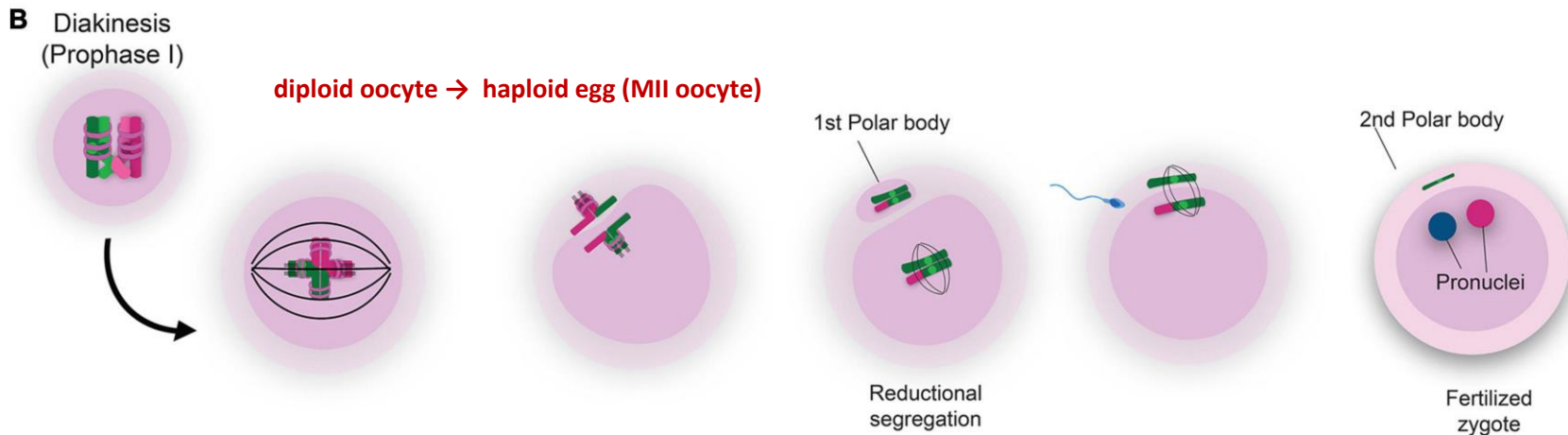
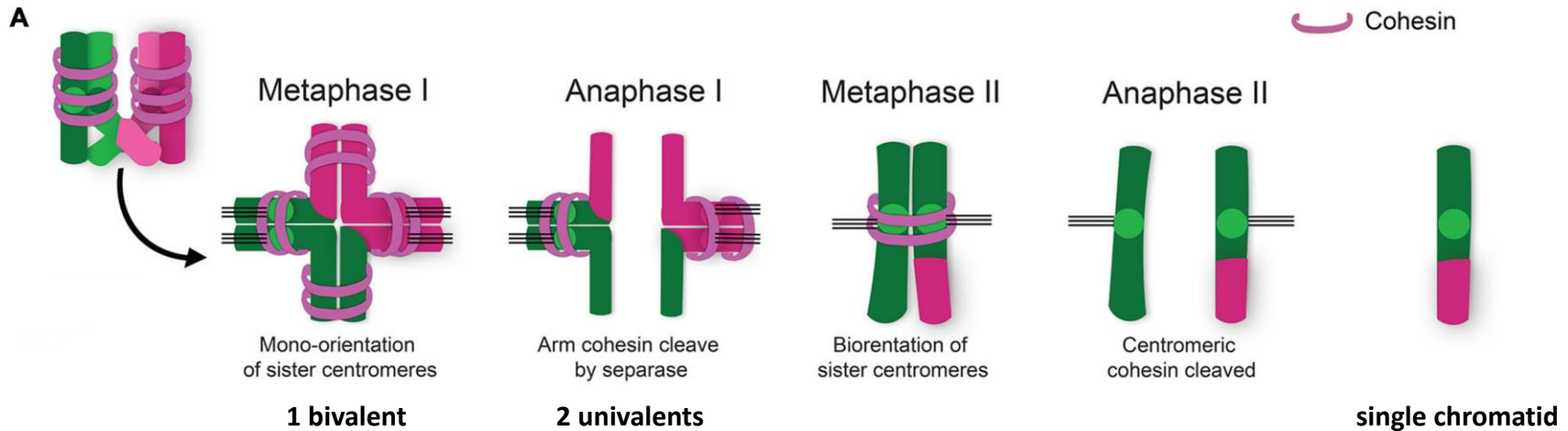
~15 min

~30 min

Microtubules DNA

Microtubule „cage“

Chromosome segregation



➤ Meiosis I

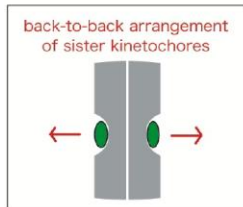
- separation of homologous chromosomes
- haploid set of chromosomes eliminates to 1st PB
- $2n\ 4C \rightarrow 1n\ 2C$

➤ Meiosis II

- separation of sister chromatids
- one set of chromatids eliminated to 2nd PB
- $1n\ 2C \rightarrow 1n\ 1C$ (+ $1n\ 1C$ from sperm)

Kinetochores-microtubule attachment

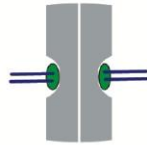
➤ MITOSIS/ MEIOSIS II



attachment:
kinetochores in relation to spindle poles

orientation:
chromosome in relation to spindle poles

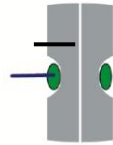
sister kinetochores attached to opposite poles



amphitelic

bi-orientation

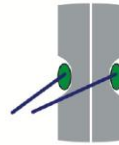
only one of the sister kinetochores attached to one pole



monotelic

(monotelic) mono-orientation

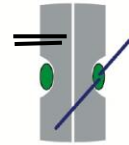
both sister kinetochores attached to the same pole



syntelic

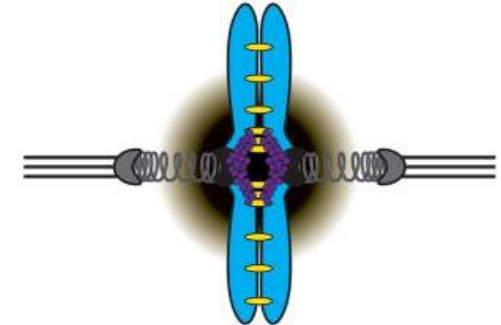
(syntelic) mono-orientation

one of the sister kinetochores attached to both poles



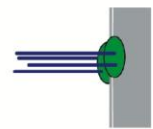
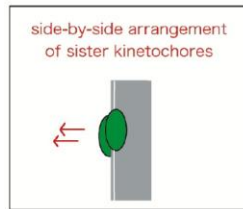
merotelic

(merotelic) bi-orientation

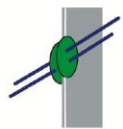


amphitelic attachment (biorientation) during Mitosis & Meiosis II

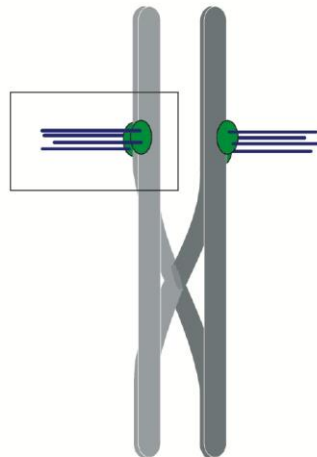
➤ MEIOSIS I



mono-orientation of sister kinetochores



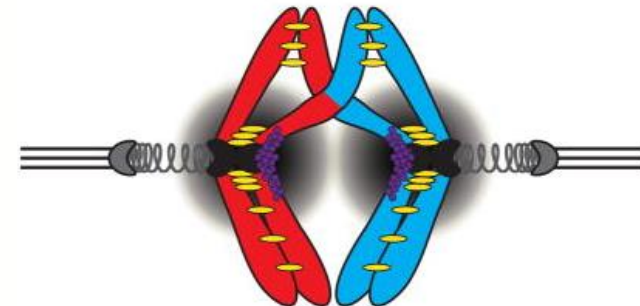
bi-orientation of sister kinetochores (side-by-side arrangement can be preserved or not)



bi-orientation of bivalent



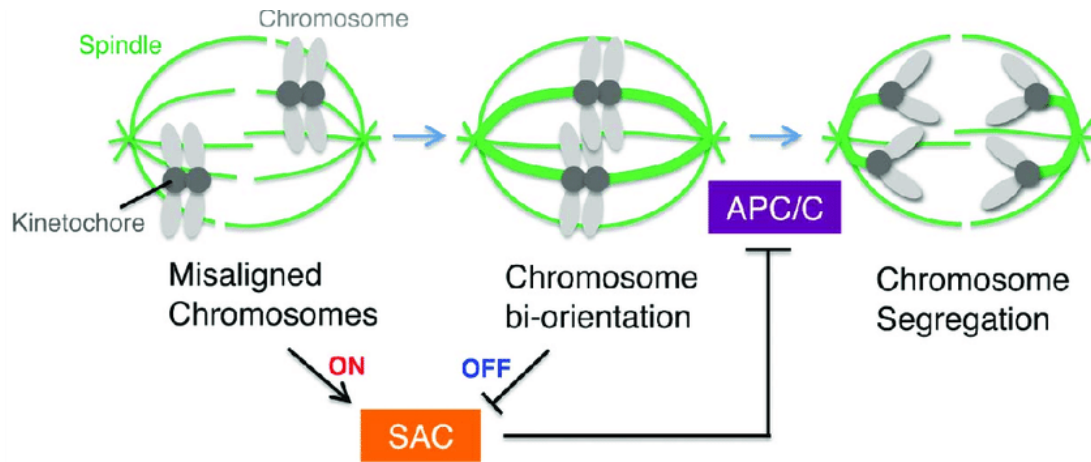
mono-orientation of bivalent



amphitelic attachment (biorientation) during Meiosis I

- sister chromatid monoorientation (behave like functional unit !)

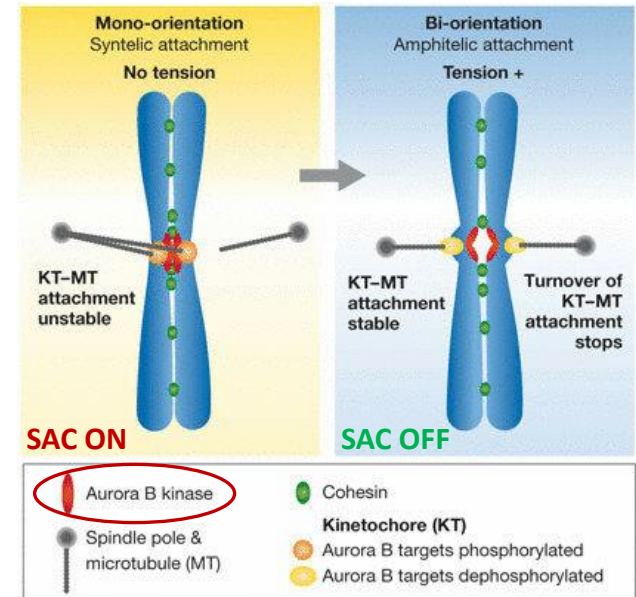
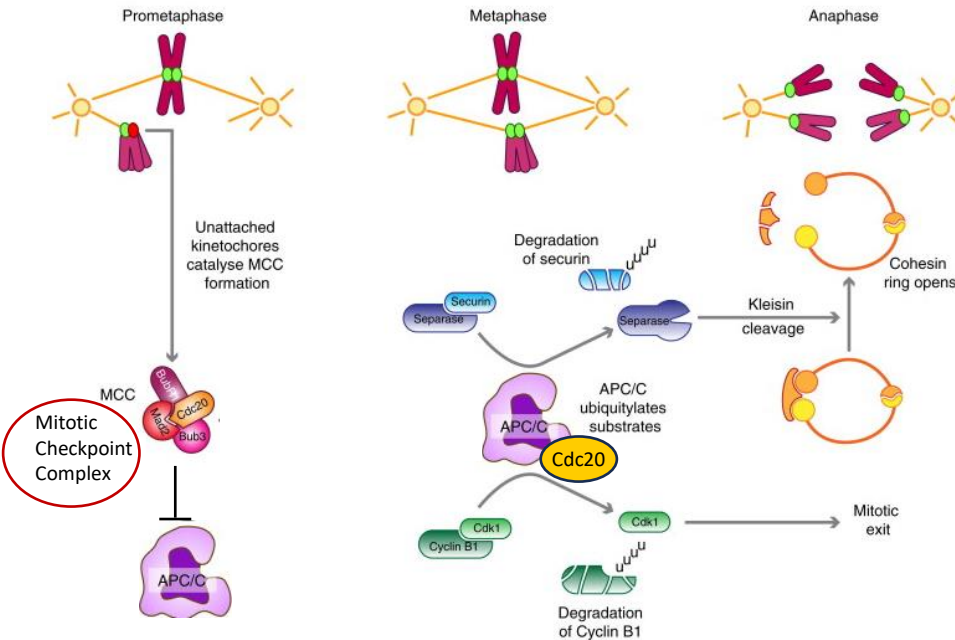
Spindle assembly checkpoint



- the pathway that delays mitosis until **all kinetochores** are attached to microtubules

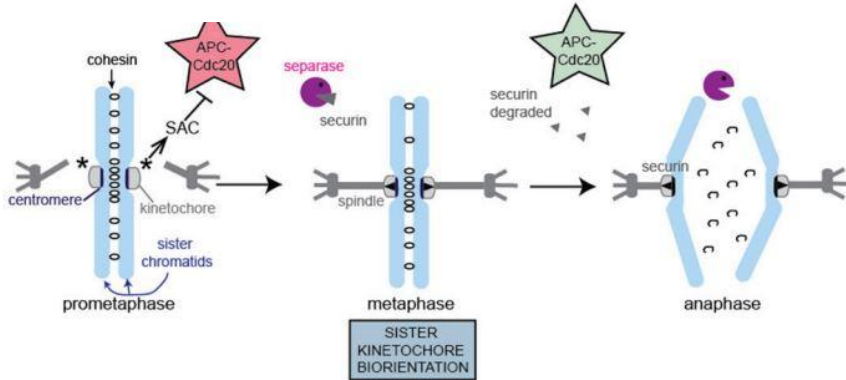
- MCC on unattached kinetochores prevent Anaphase Promoting Complex (APC) from cyclin B and securin destruction

- SAC responds to interkinetochore tension



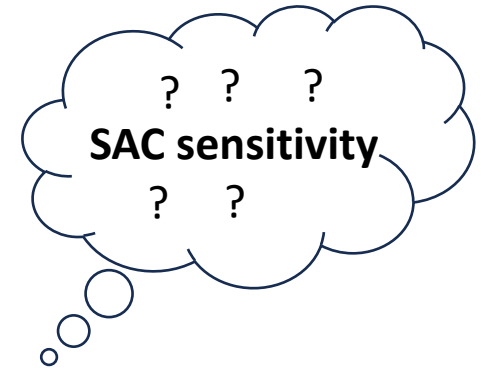
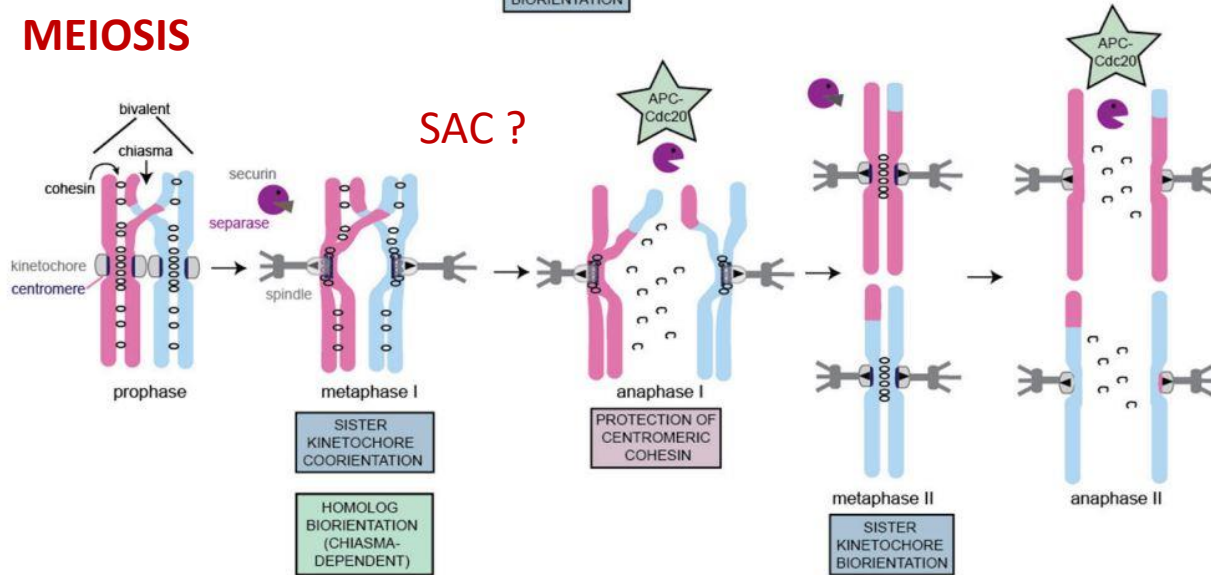
Spindle assembly checkpoint

➤ MITOSIS



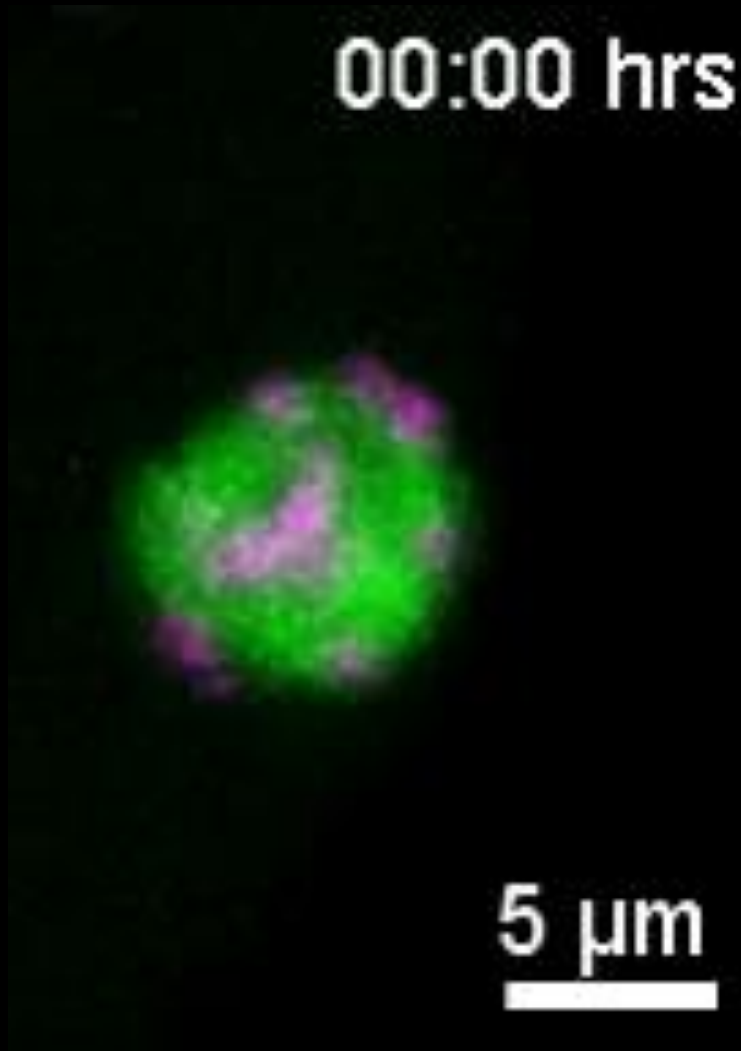
- all kinetochores must be occupied
- APC/C activation in minutes
- low aneuploidy rate

➤ MEIOSIS



- only a majority of kinetochores must be occupied
- APC/C activation in hours
- high rate of aneuploidy

Spindle assembly checkpoint



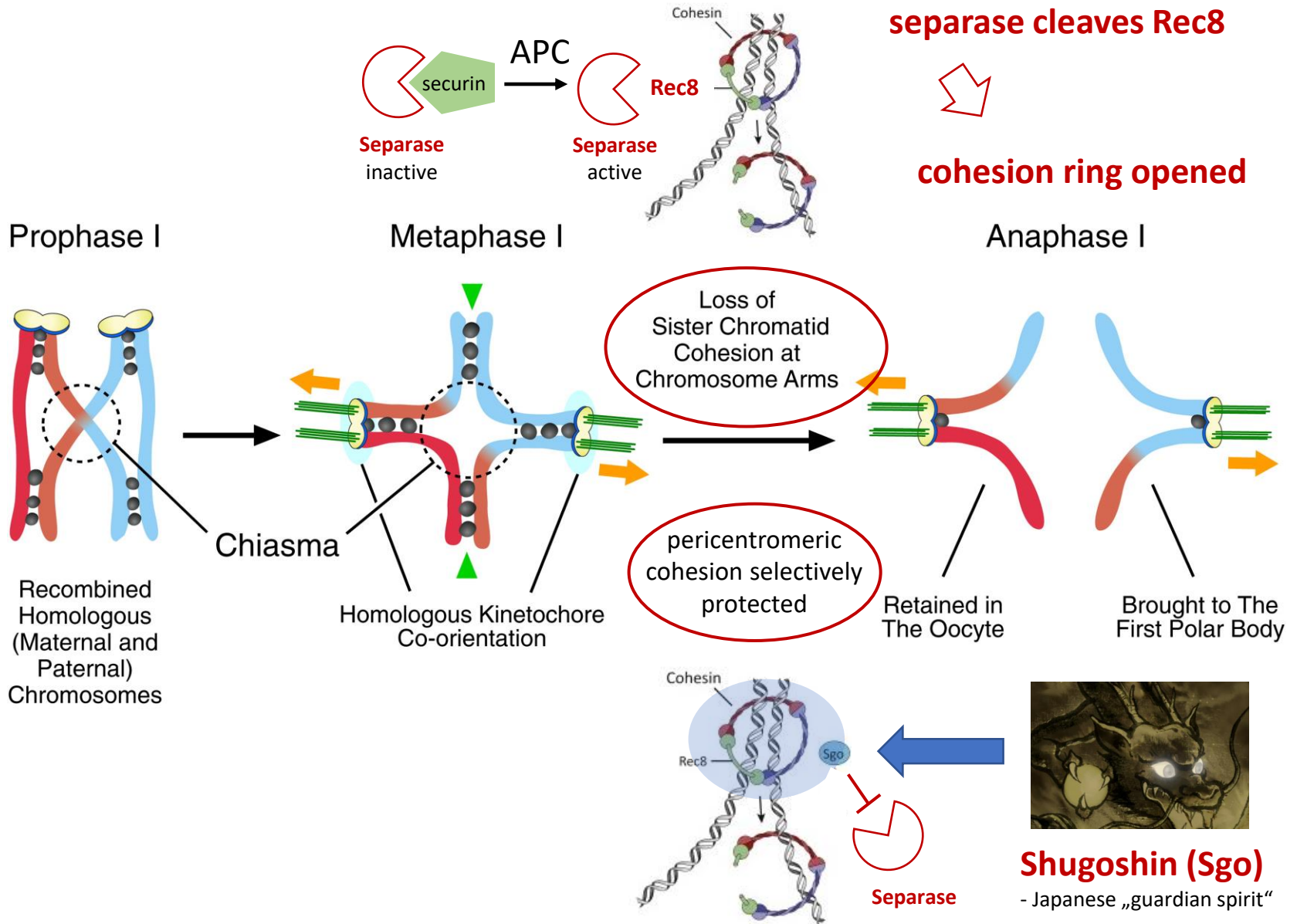
00:00 hrs

5 μm

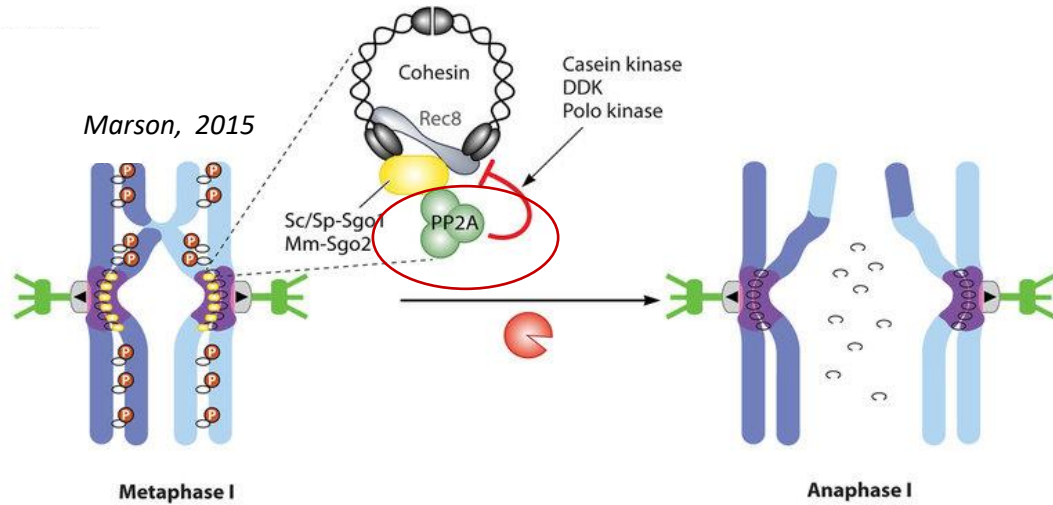


MAP4 – microtubules
H2B - DNA

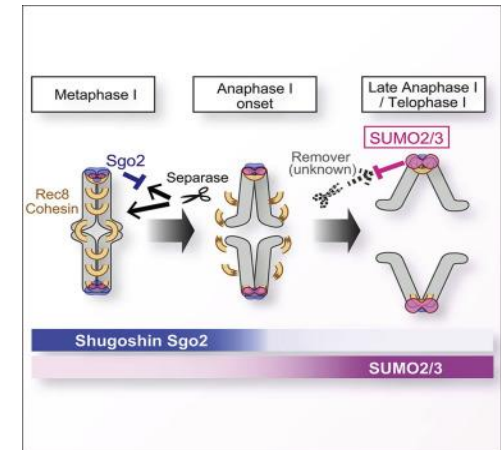
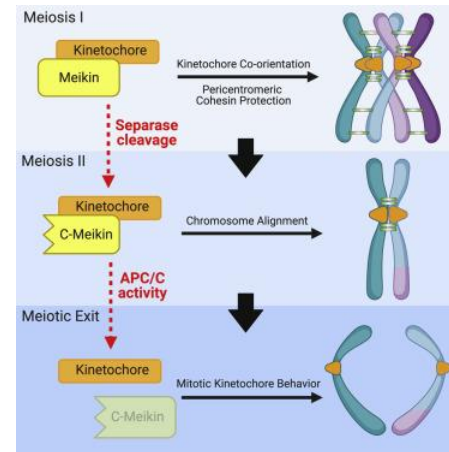
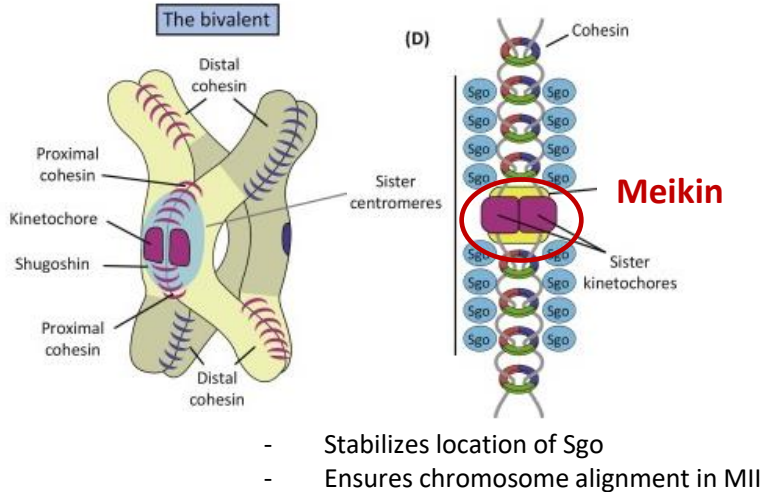
Loss of cohesion in anaphase I



Loss of cohesion in anaphase I

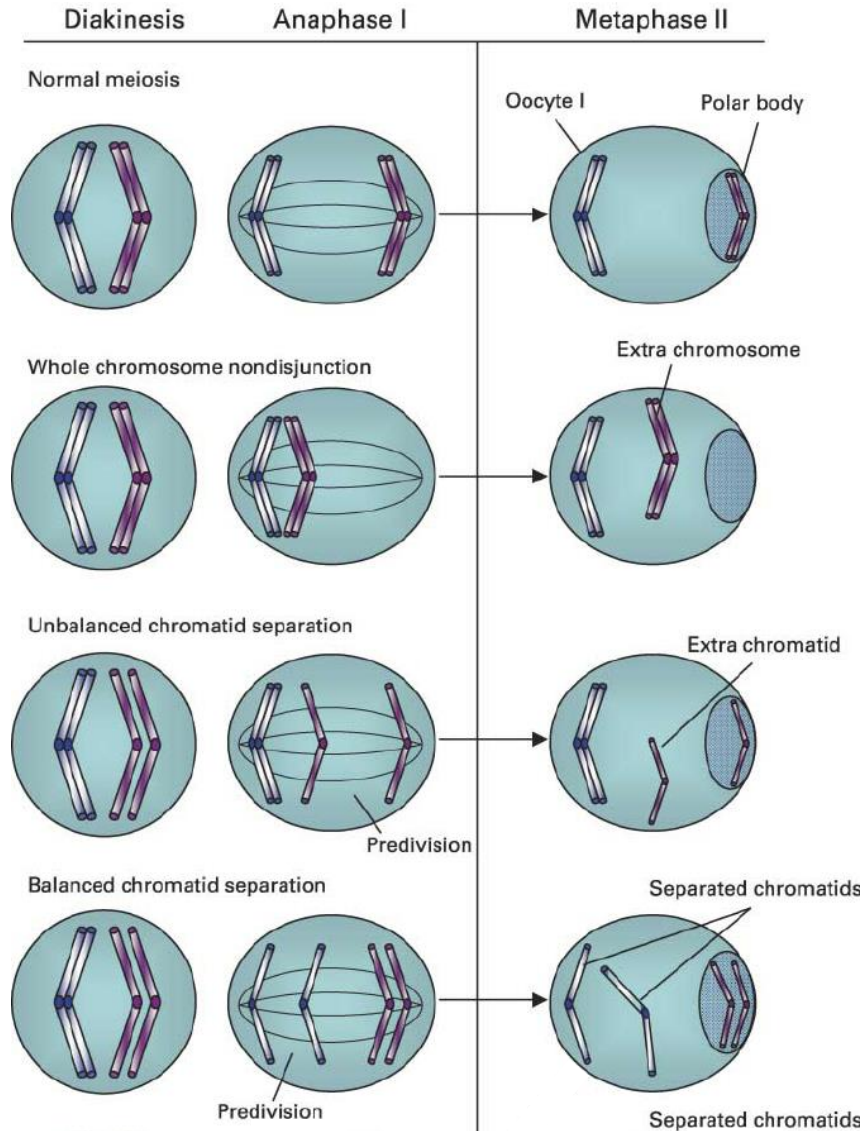


Sgo recruits **PP2A**, which removes Rec8 phosphorylation, making it a poor substrate for separase-dependent cleavage.



Chromosome segregation errors

**Oocyte
aneuploidy**




**Chromosome
nondisjunction (NDJ)**

**Precautious
separation of sister
chromatids (PSSC)**

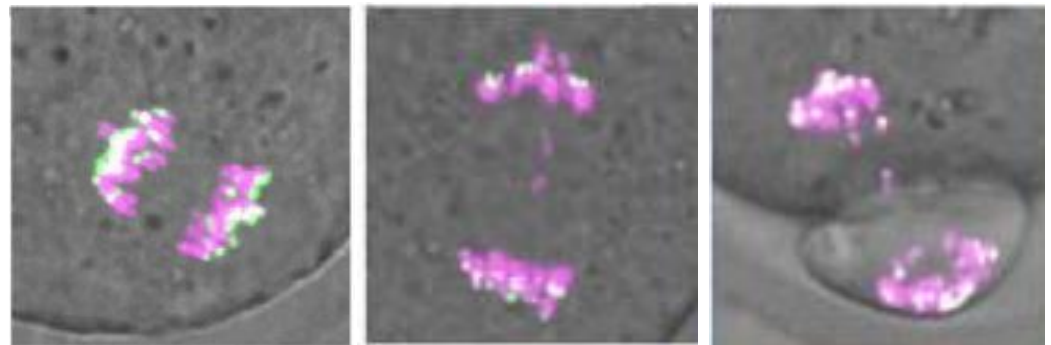
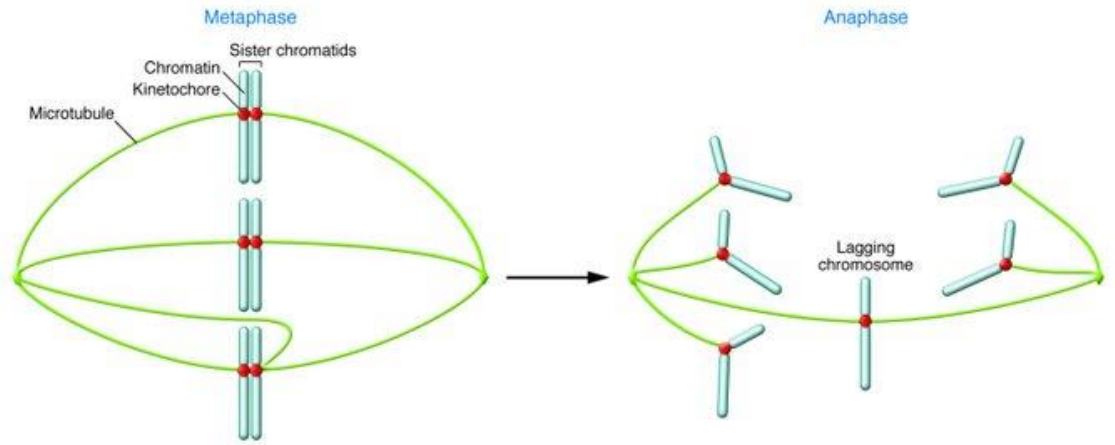
Chromosome lagging

- delayed chromosome/chromatid movement during anaphase

- risk of inaccurate segregation, chromosome loss/gain and aneuploidy

 oocytes

- class I – laggards
 - aligned at anaphase
 - reduced velocity
 - aneuploidy producing
- class II – laggards
 - misaligned at anaphase
 - normal velocity
 - benign

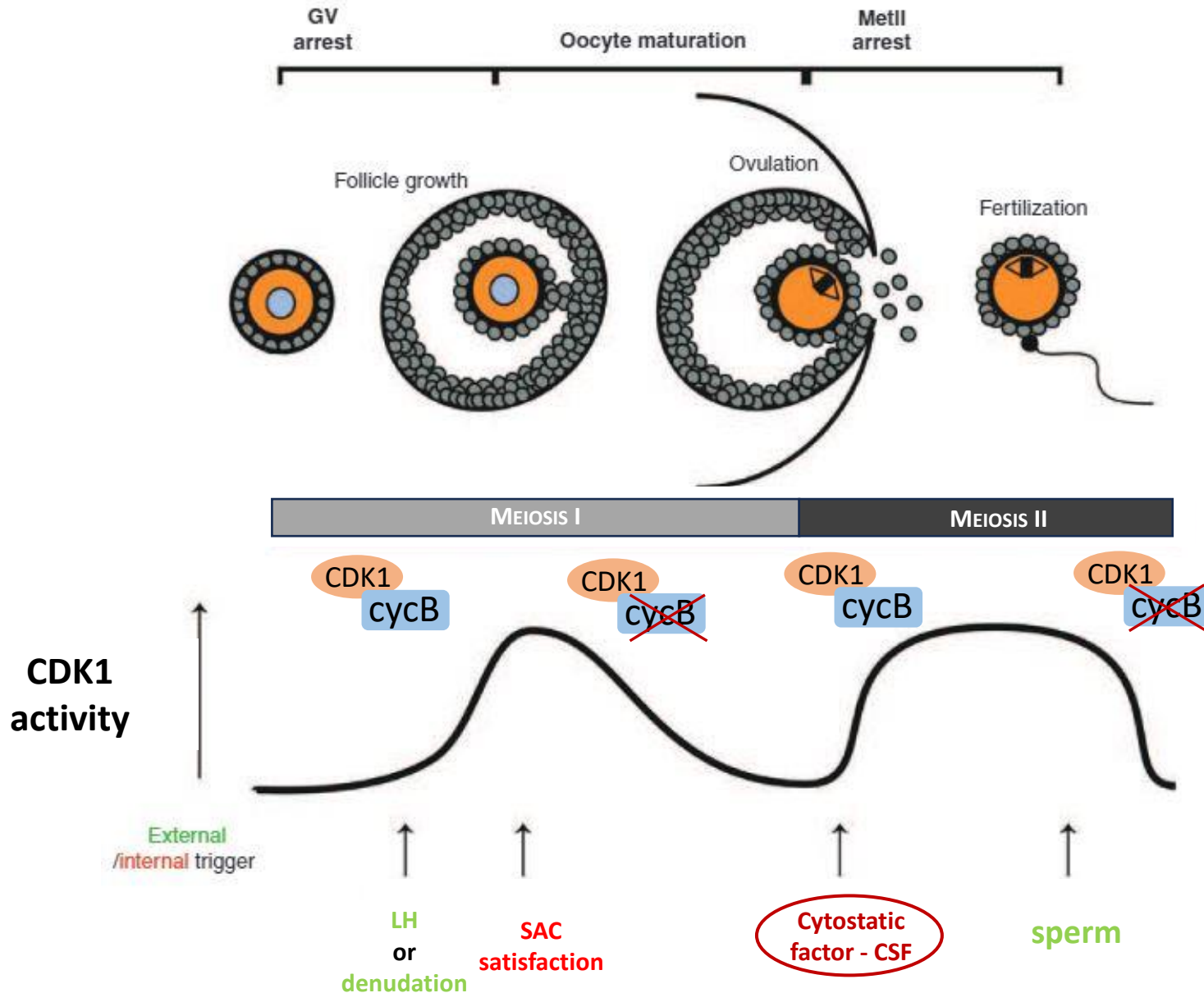


No lagging

Class I laggards

Class II laggards

Transition from meiosis I to meiosis II



Metaphase II arrest

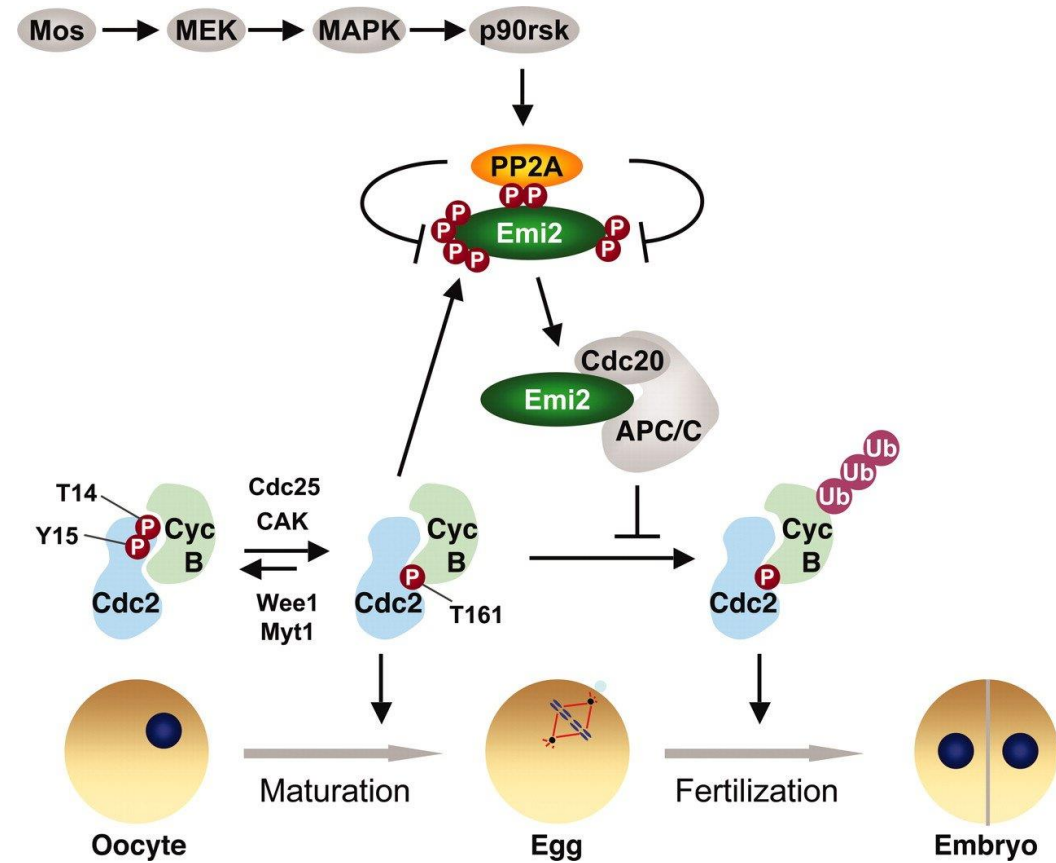
➤ **Emi2** (Early Mitotic Inhibitor)

- meiosis specific inhibitor of APC (= „cytostatic factor“ - CSF)
- required for establishment and maintenance of MII arrest in mammalian oocytes
- phosphorylation needed to keep Emi2 stable

Magwick et al 2006

➤ **Btg4**

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

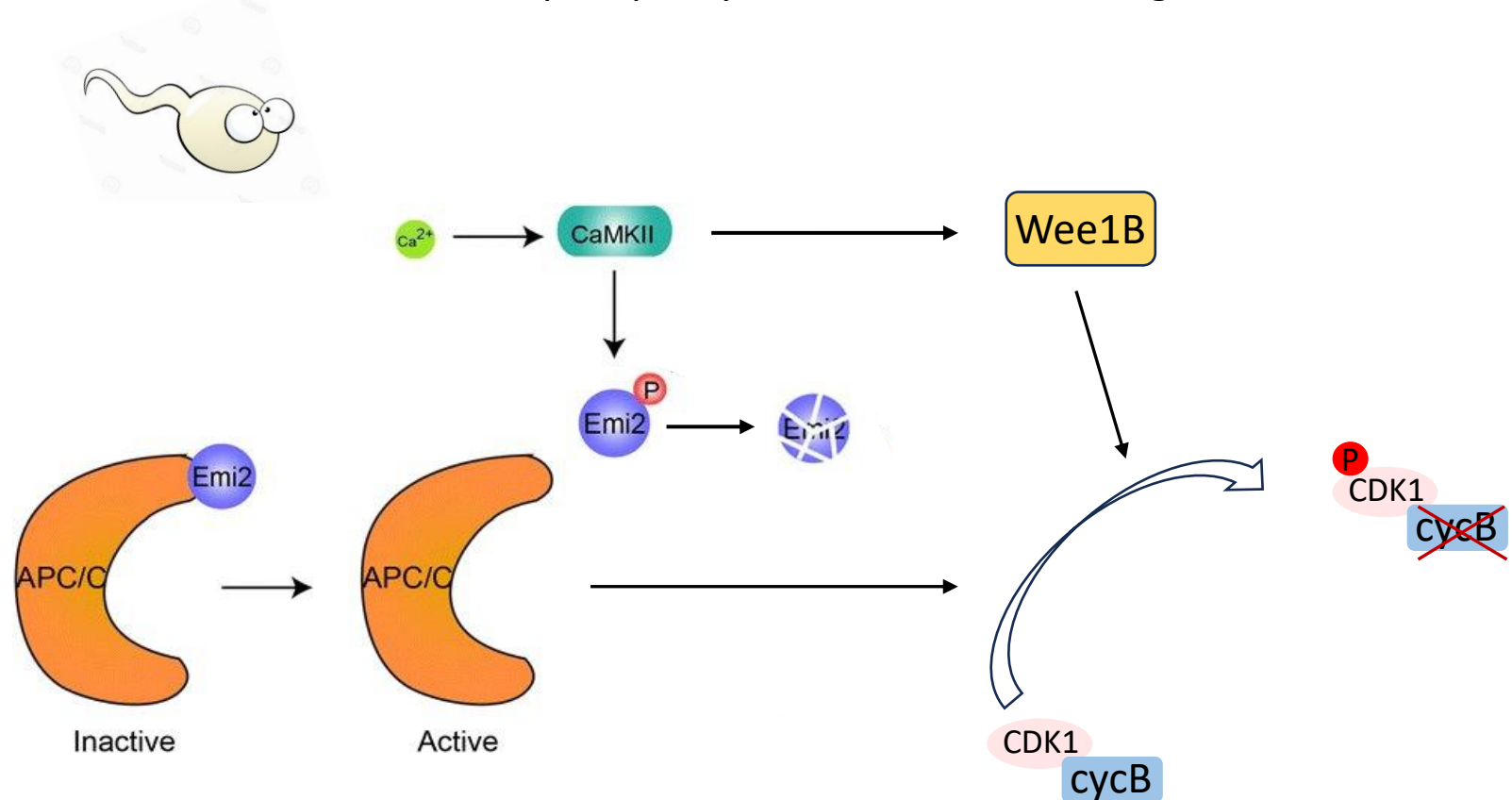


- MII arrest reached hours before ovulation and maintained for ~ 24 hours

MII arrest release

➤ Calmodulin dependent protein kinase II (CAMKII)

- activated by Ca^{2+} signal at fertilization
- phosphorylates Emi2 causing its degradation
- activates Wee1B kinase that phosphorylates CDK1 contributing to its inactivation



Fertilization stage in different species

