

The unknowns of early human development and the impact on genome editing

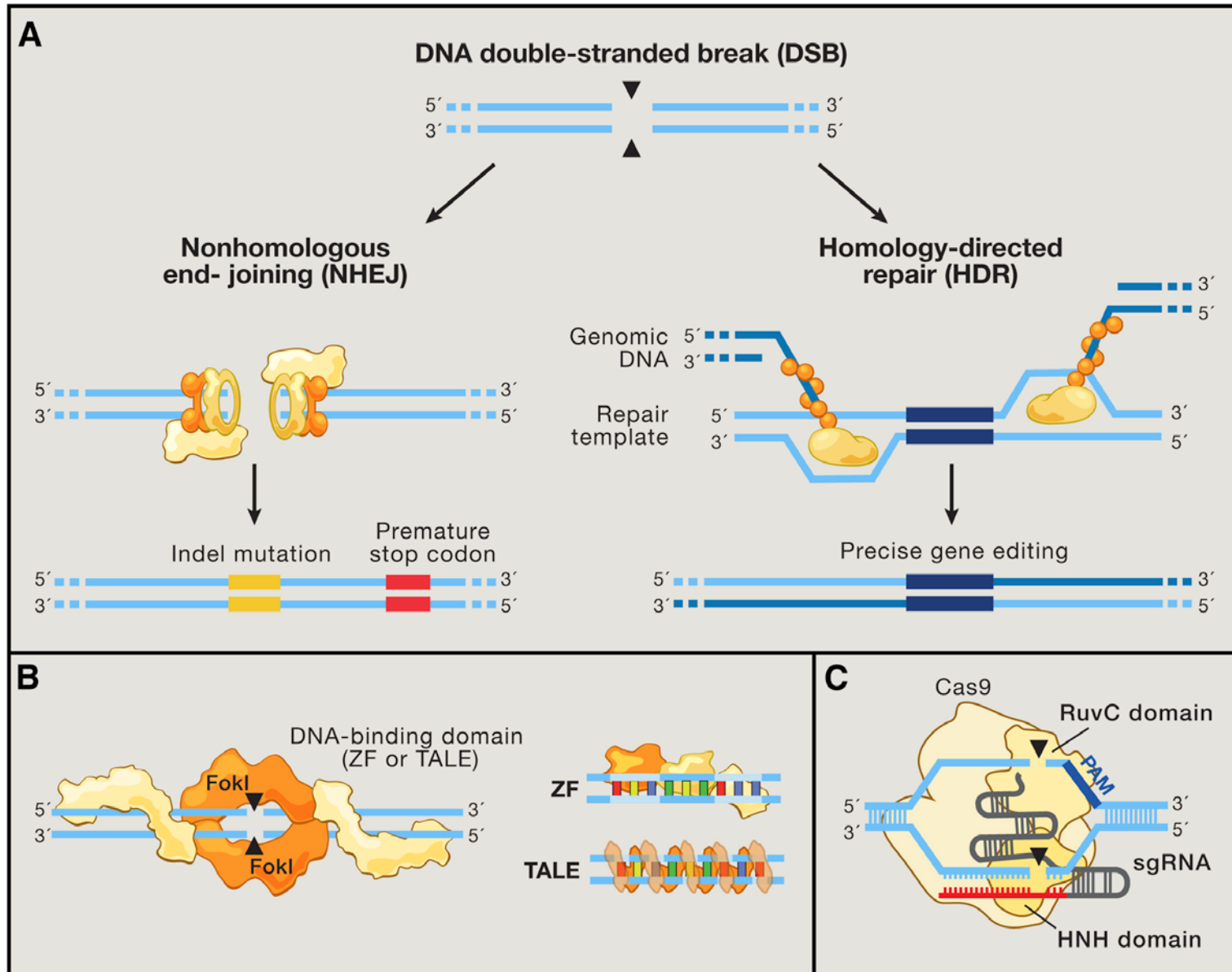


Kathy Niakan

Efficient genome editing of human embryos
depends on an understanding of:

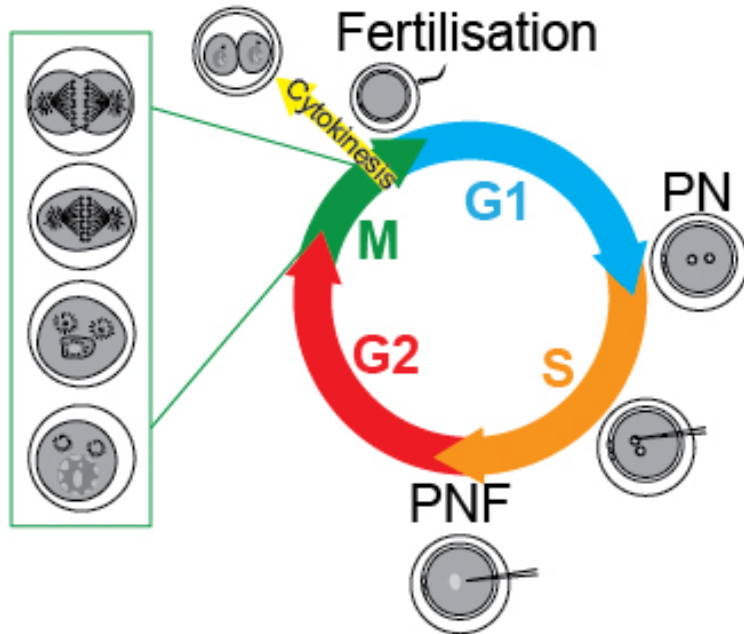
1. Cell cycle
2. Chromosome segregation
3. DNA-damage repair

Methods of genome editing

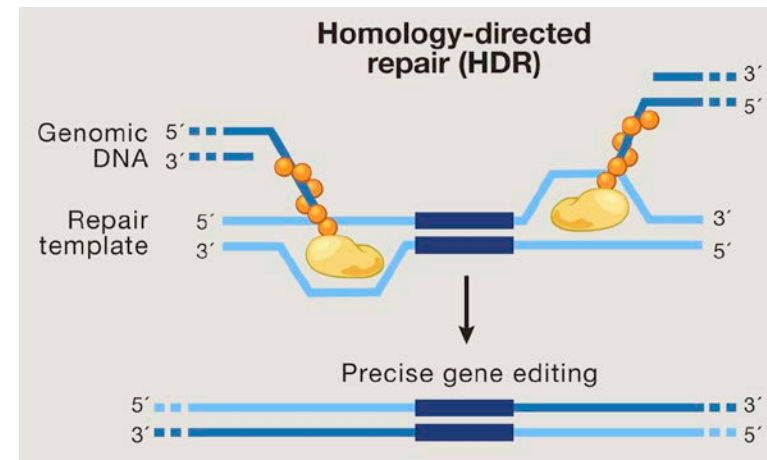


DSB repair pathway choice depends on cell cycle

G1 phase

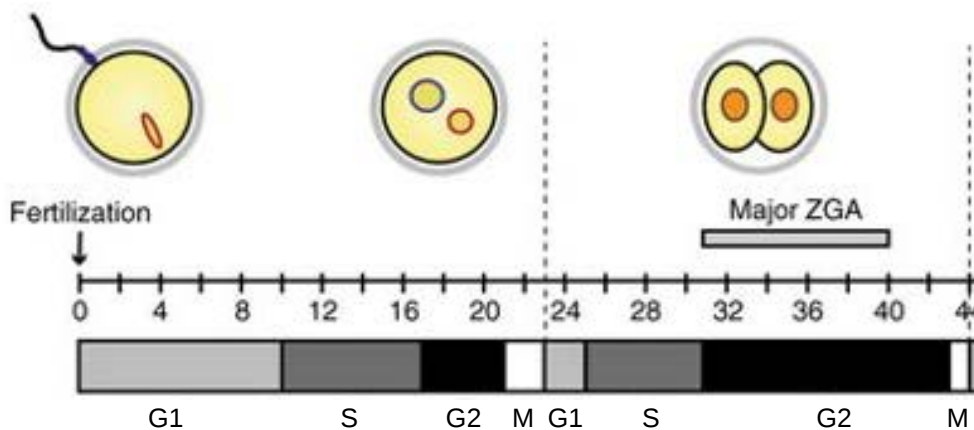


G2/S phase

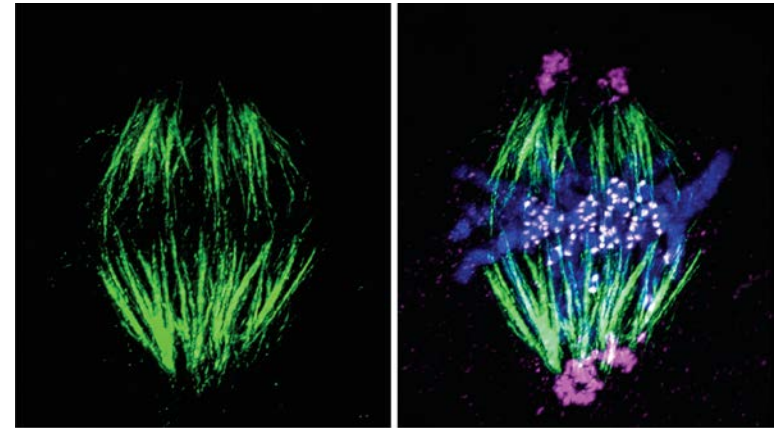
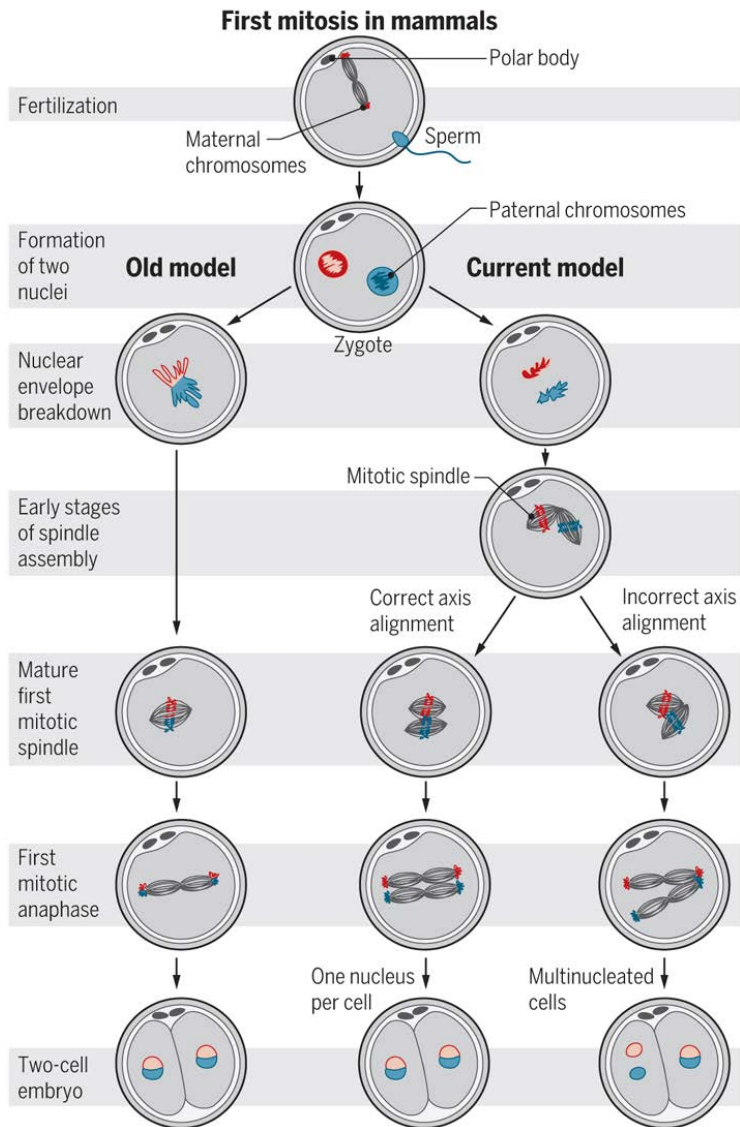


Hsu et al., Cell 2014

Ma et al., Nature Biotech 2018



Parental genomes are kept apart at the first cell division(s) by dual spindle formation



Reichmann et al., *Science* 2018
Zielinska and Schuh, *Science* 2018

Spindle abnormalities in early human embryos

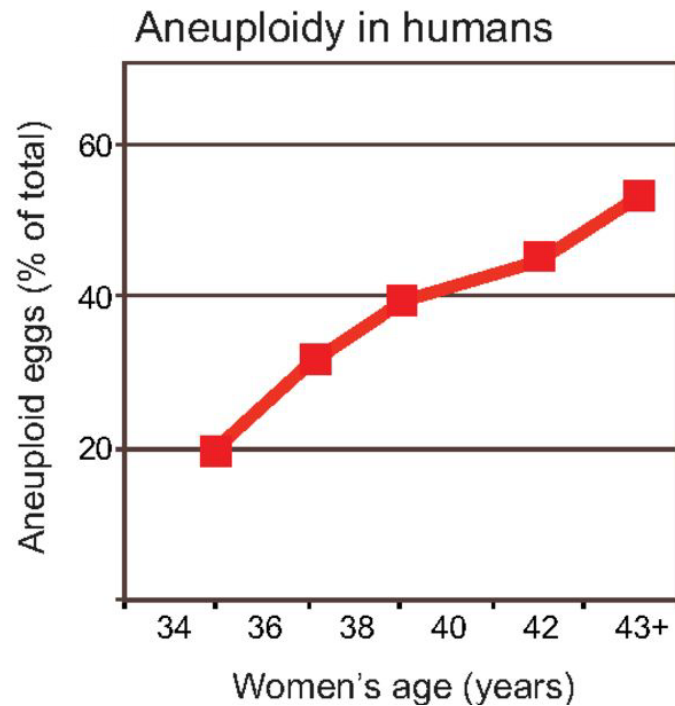
Bipolar spindle formation



Multipolar spindle formation



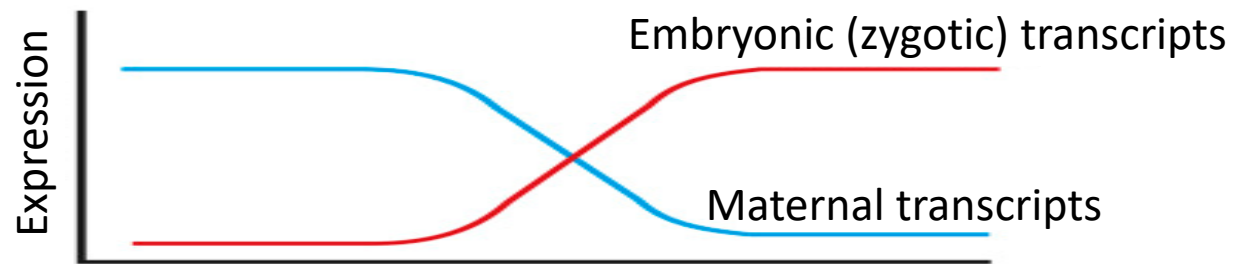
Human embryos exhibit high rates of aneuploidy



Jones and Lane, *Development* 2013

Consequence of the timing of embryo genome activation

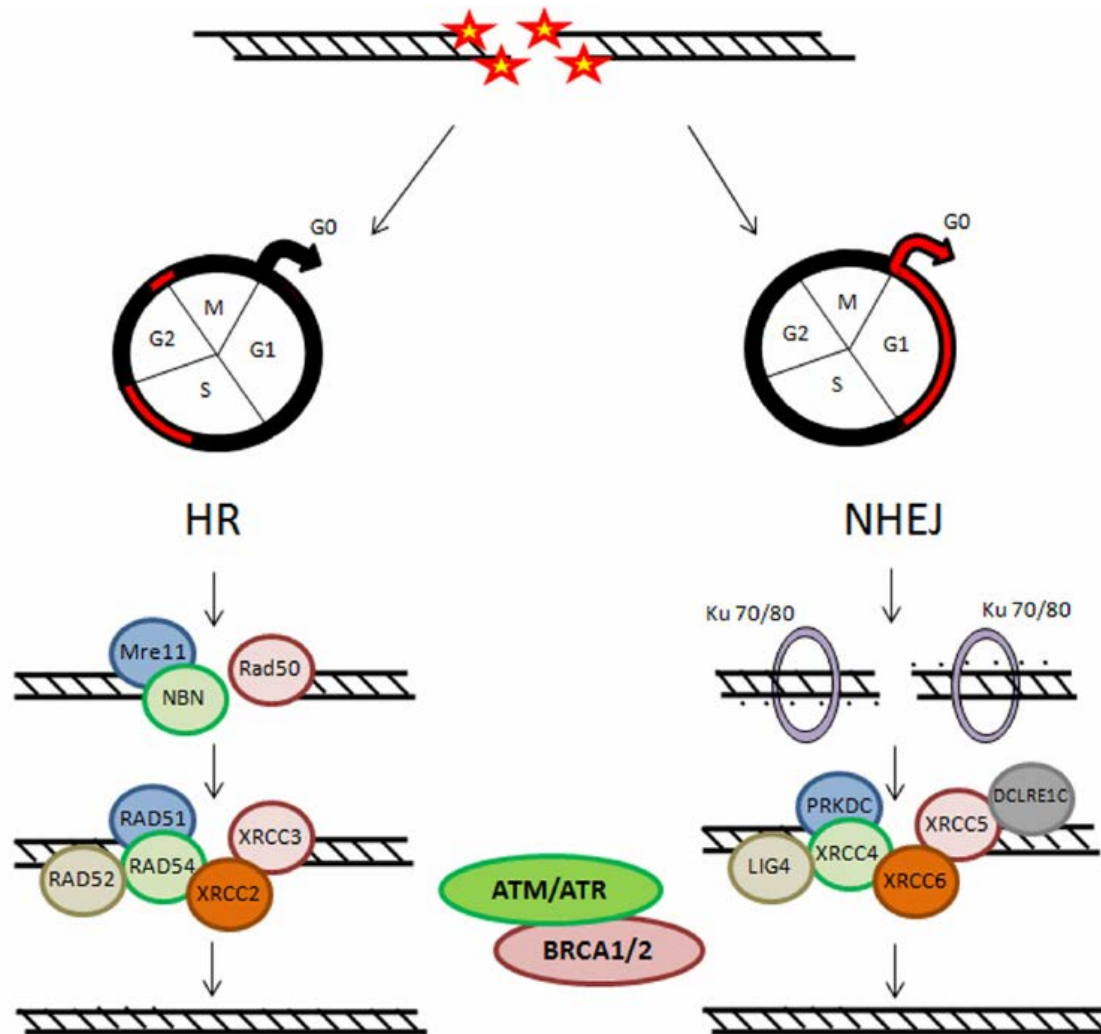
Embryo genome
activation



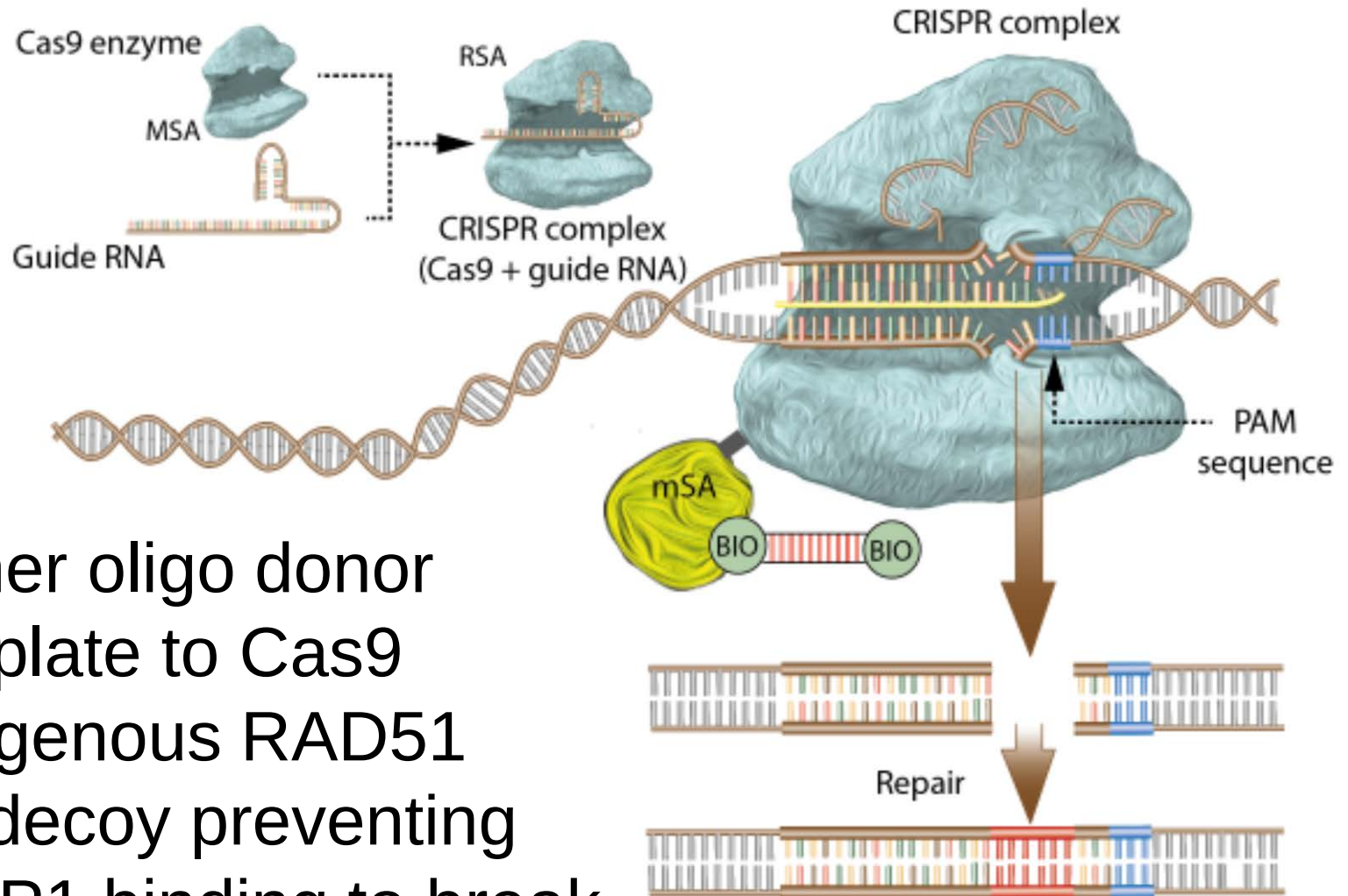
Treen *et al.*, *Current Biology* 2018

Niakan *et al.*, *Development* 2012

Unclear which DDR factors are present in early human embryos

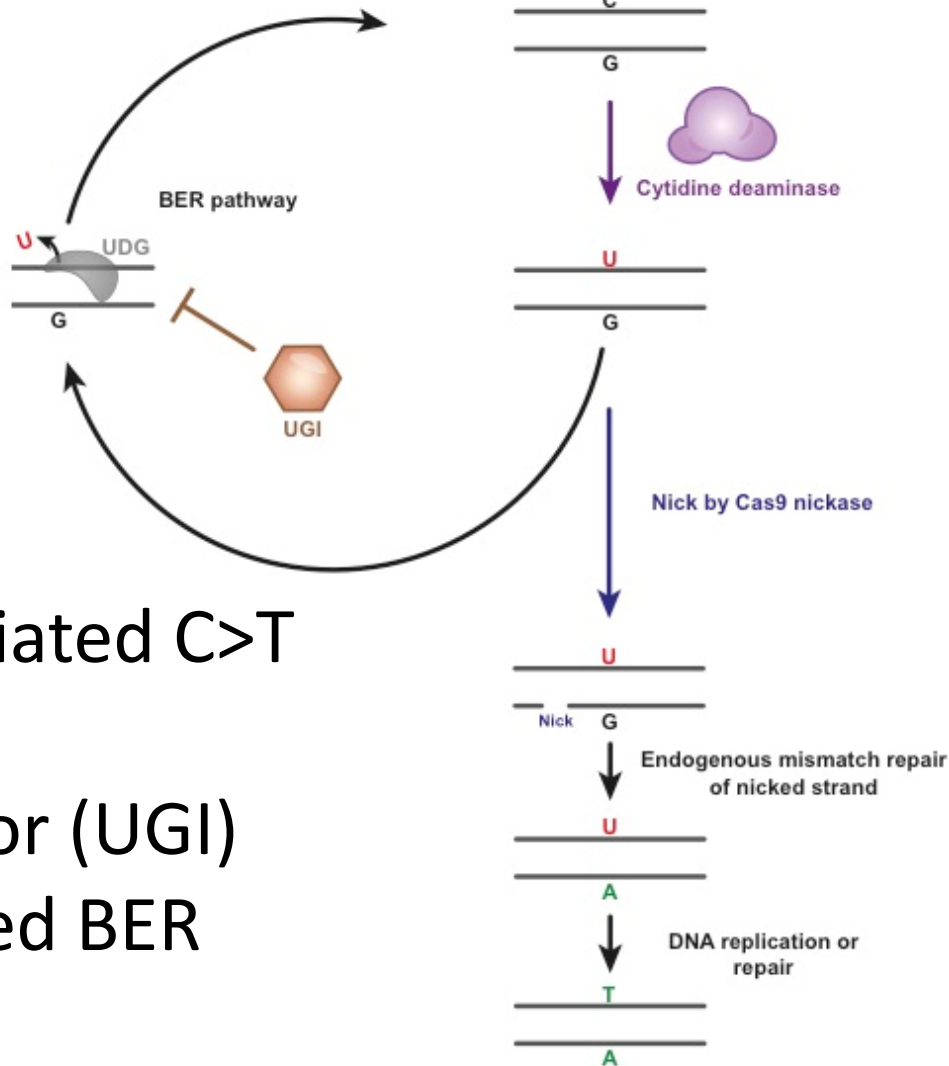
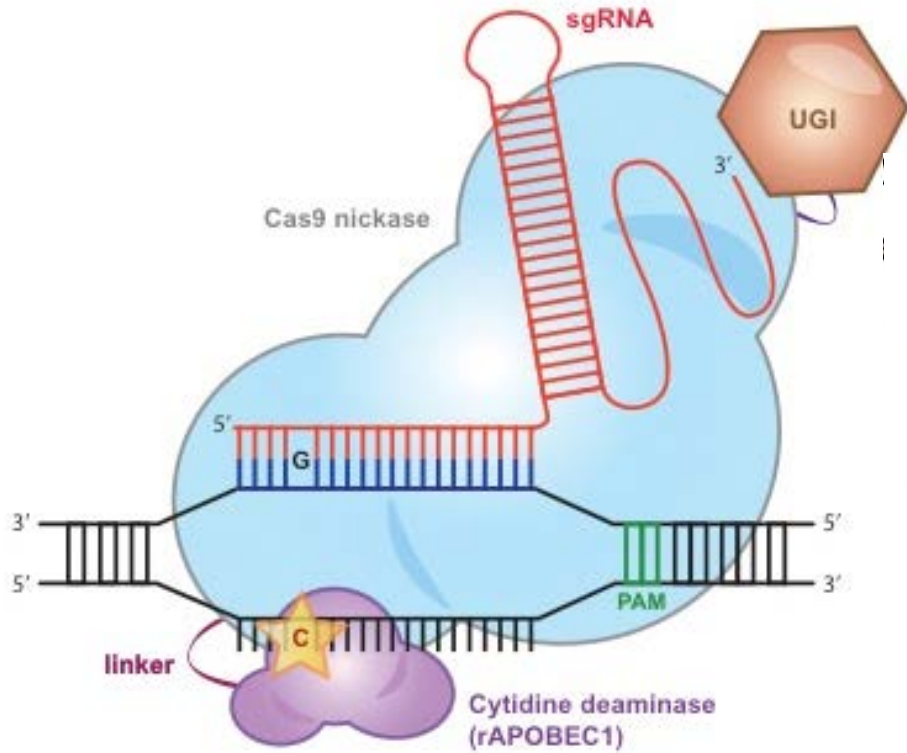


Promoting homologous repair



- Tether oligo donor template to Cas9
- Exogenous RAD51
- i53 decoy preventing 53BP1 binding to break sites

Base editing



- Cytidine deaminase mediated C>T (G>A)
- Uracil glycosylase inhibitor (UGI) inhibits the UDG-mediated BER pathway
- Cas9 nickase promotes mismatch repair

HFEA – Scientific and Clinical Advances Advisory Committee, Horizon Scanning Panel and the Ethics and Law Advisory Group

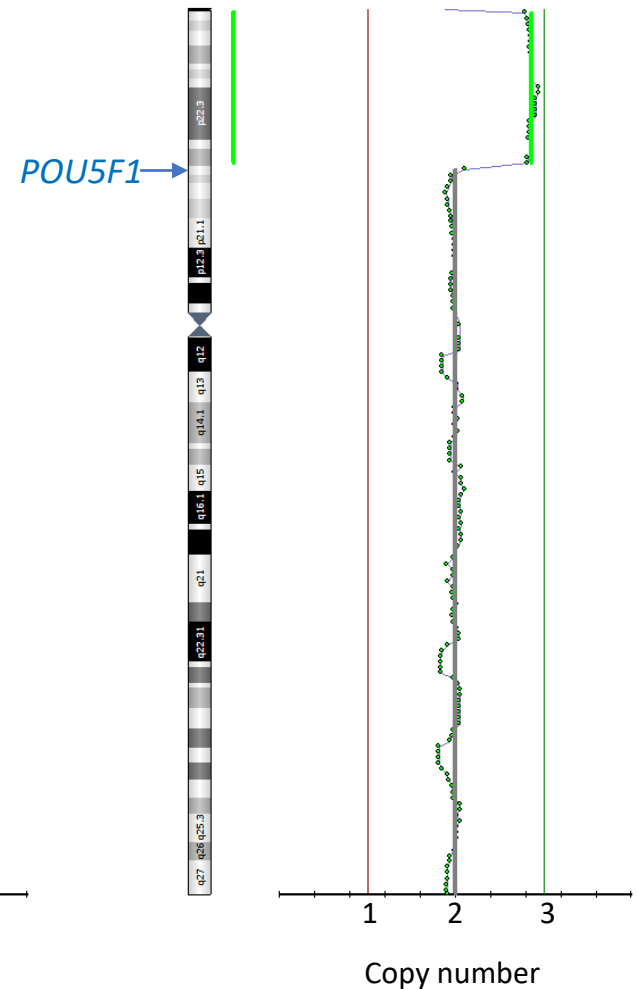
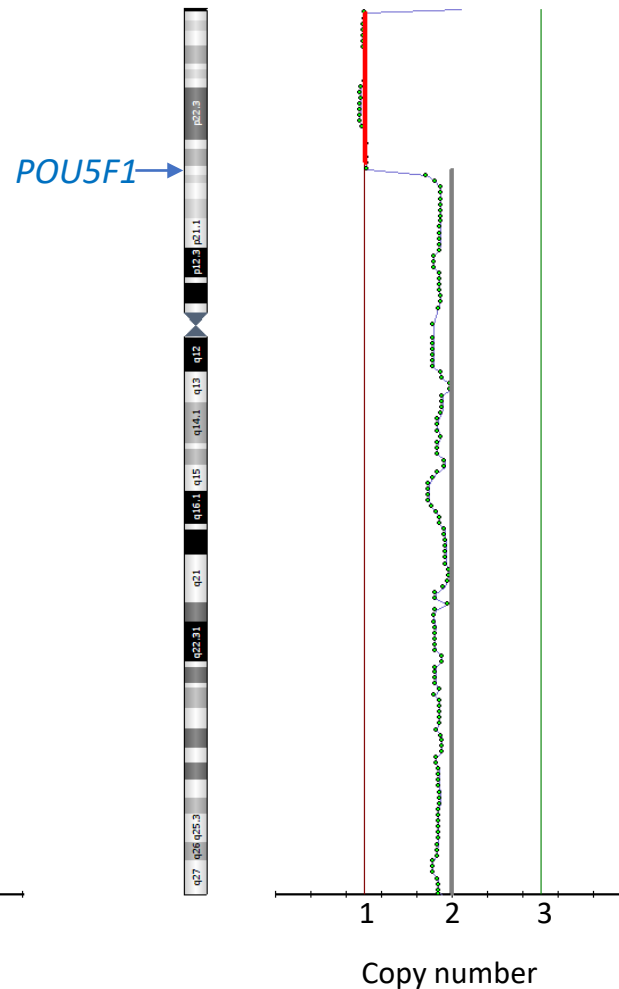
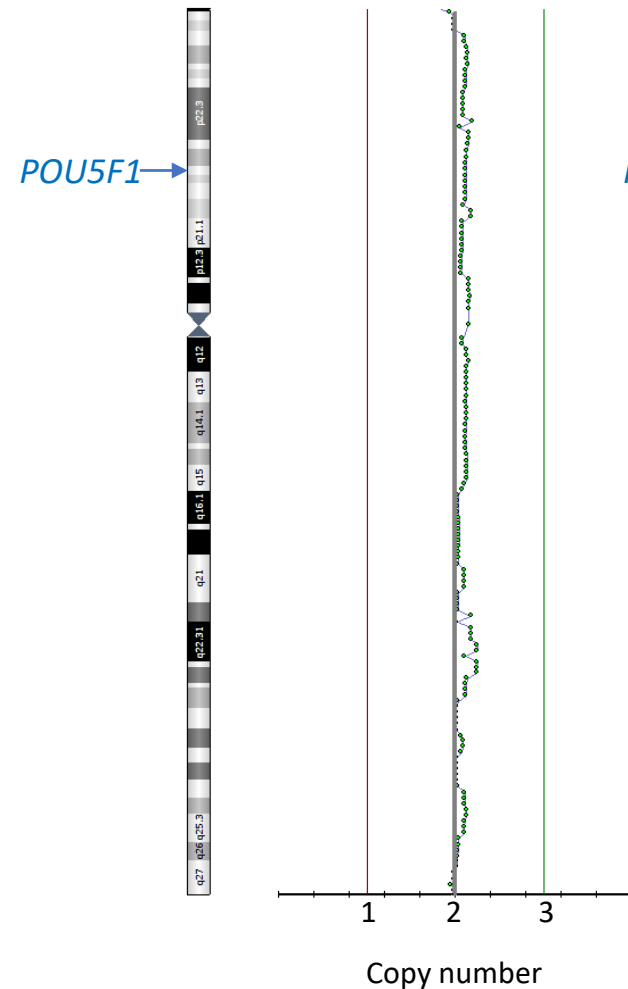
1. Research into human embryo development – specifically the roles of genes and growth factors involved in early development
2. The development of more objective criteria for embryo selection – by investigating gene function in early embryogenesis
3. Research into the genetic background of adverse medical conditions – genetically modified ES cells could be create to model medical conditions
4. Research into the fate of cells during embryo development
5. Introducing a gene to increase the efficiency of stem cell derivation

Segmental loss or gain of Chr6 in human embryos following CRISPR/Cas9 genome editing

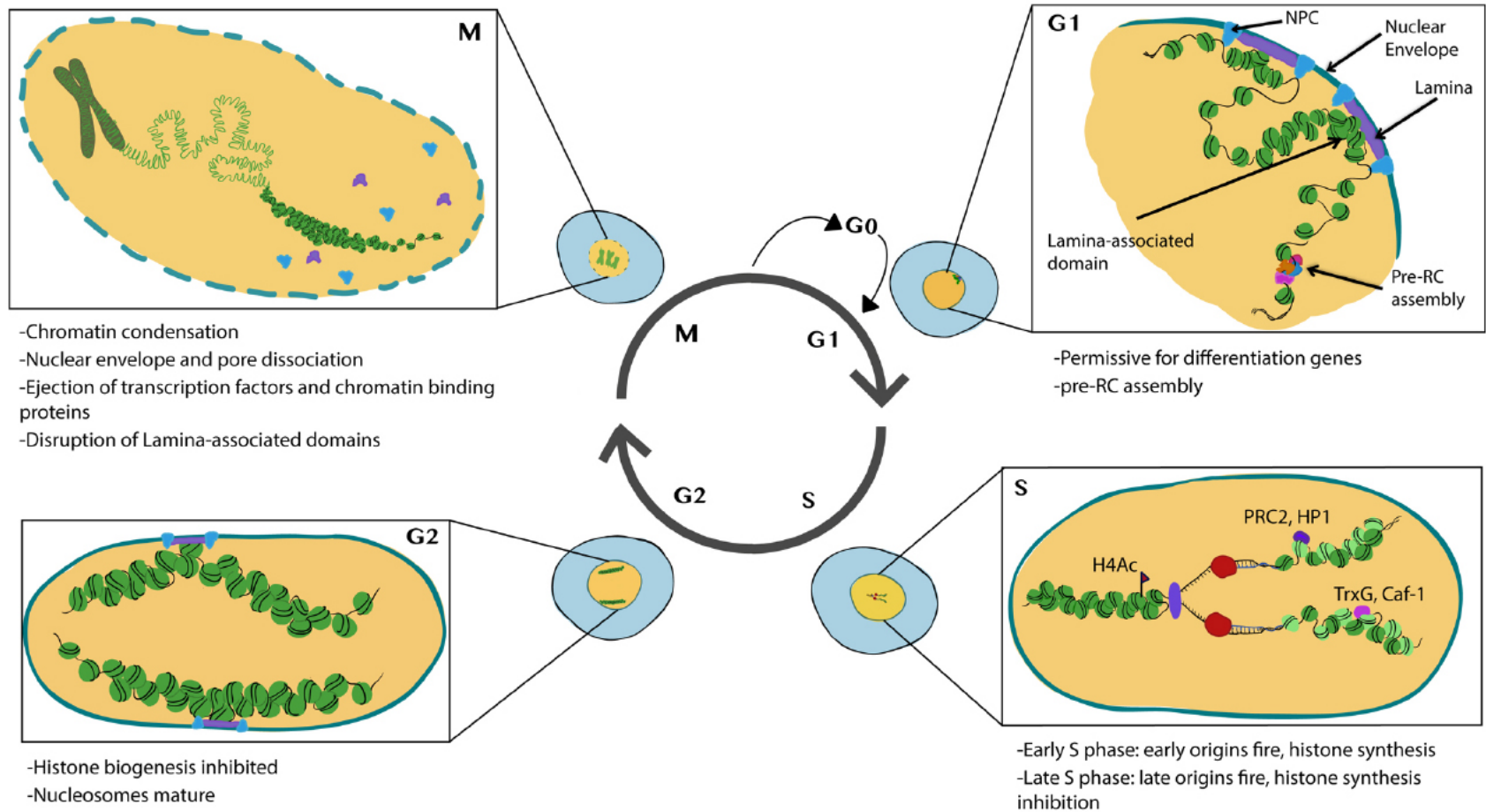
Segmental loss of Chr6
-6p22.3 (29.1 Mb)

Segmental gain of Chr6
+6p22.3 (29.1 Mb)

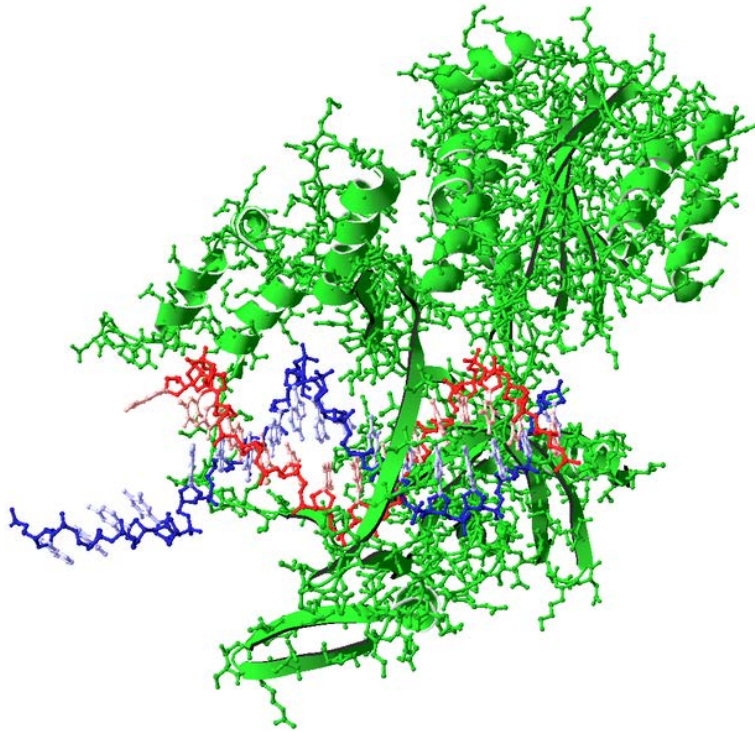
Chr6



Unpublished

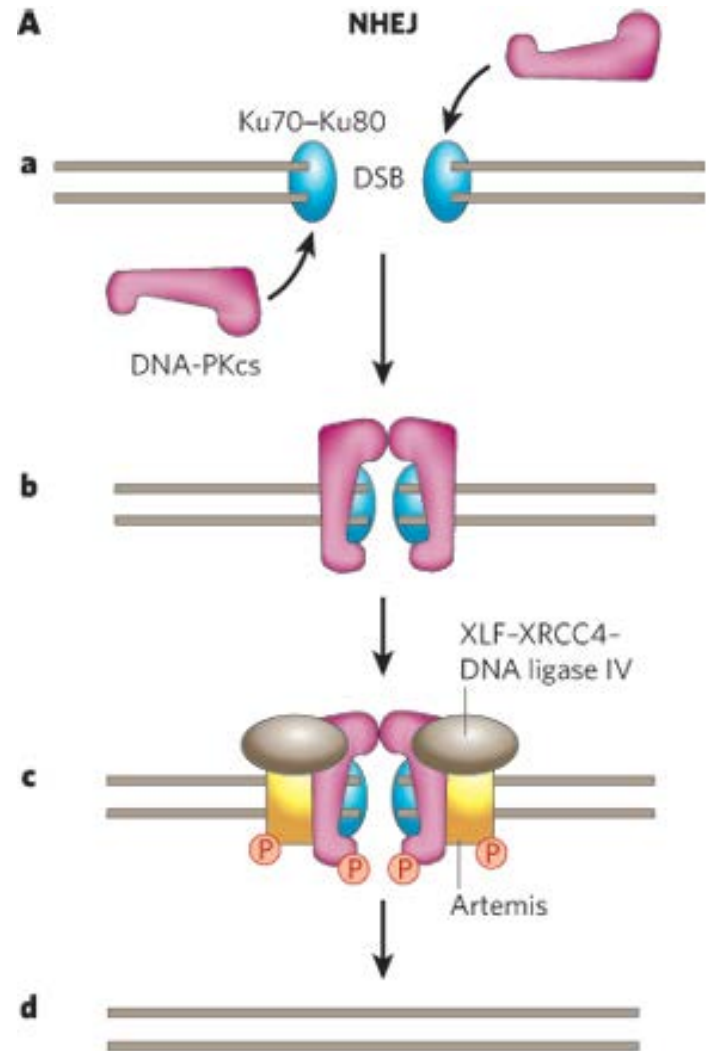


Non-homologous end joining (NHEJ)



Ku70/Ku80 heterodimer

- Homology independent
- RAD51-independent
- Error prone



Individual bipolar spindle formation around each pronucleus

