

Stockholm: 7 December, 2012.

The egg and the nucleus: a battle for supremacy

J. B. Gurdon

Cambridge, England.

Content

Background

Attack by the egg

Defense by the nucleus

Prospects

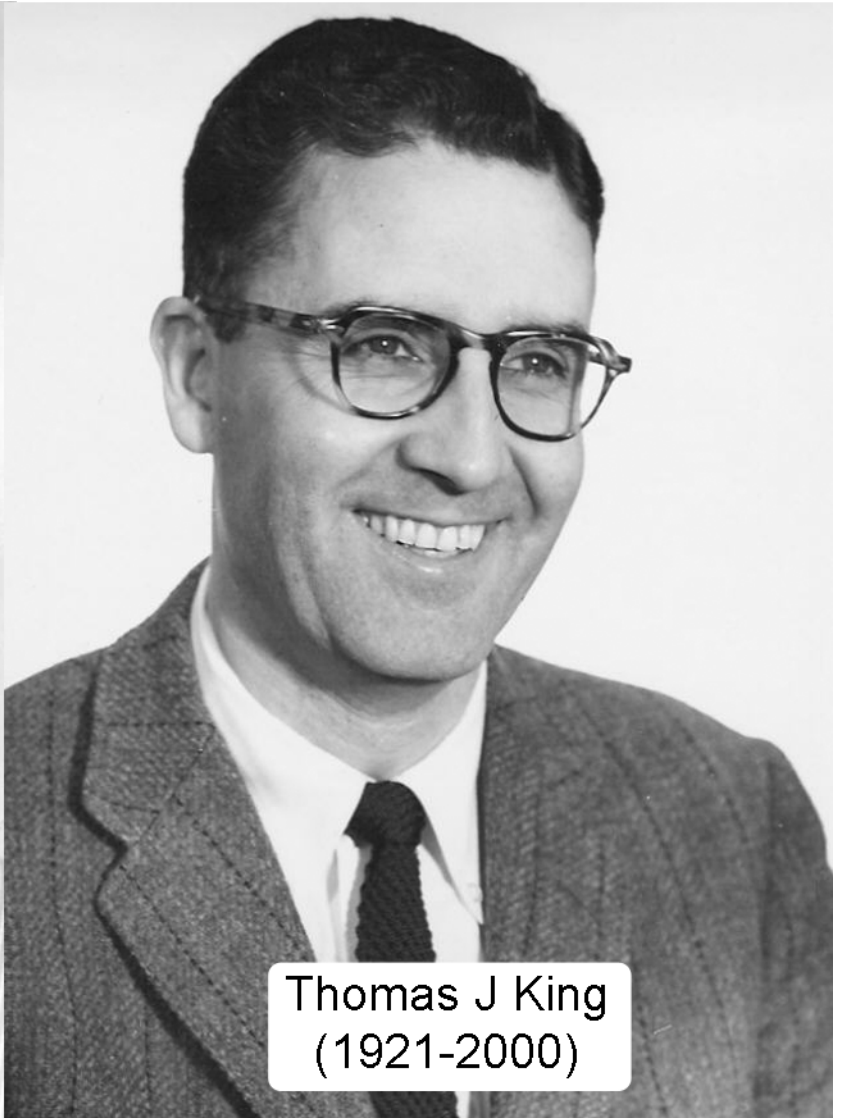
Background

The original question

**Do all cells in the body
have the same sets of genes?**

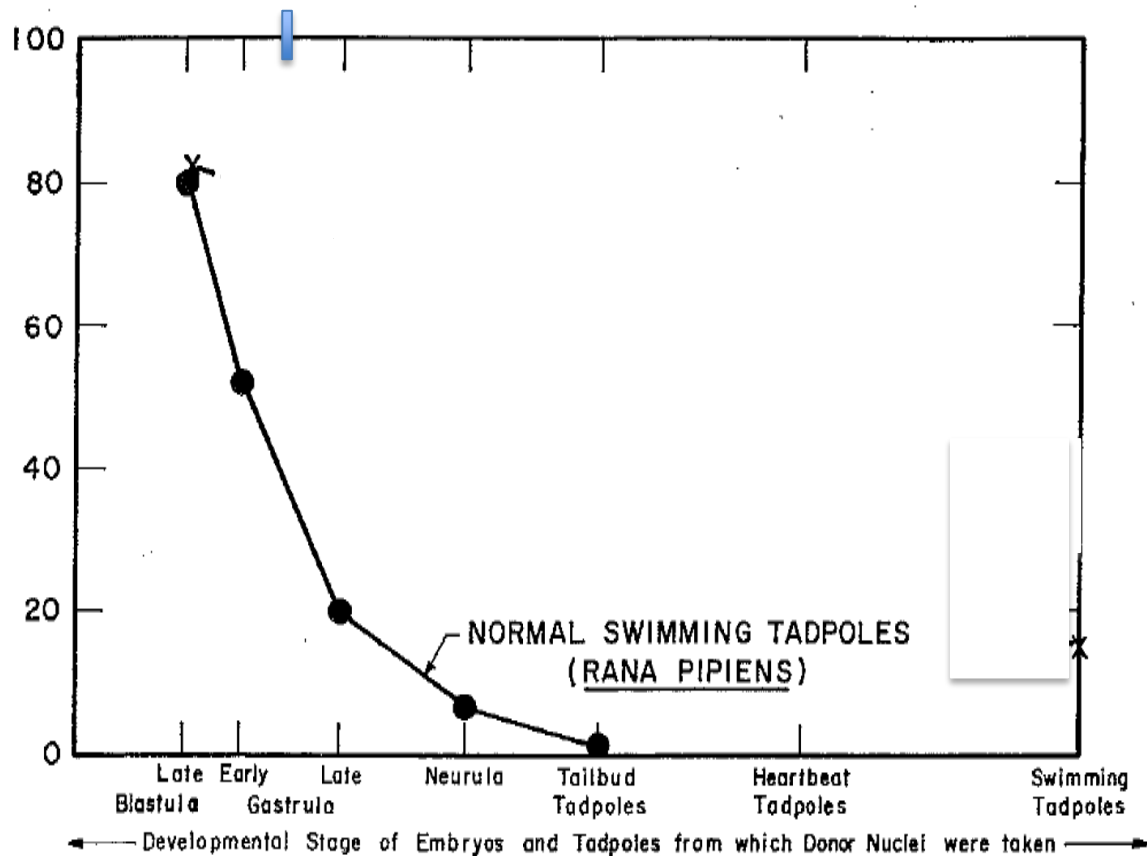


Robert Briggs
(1911-1983)



Thomas J King
(1921-2000)

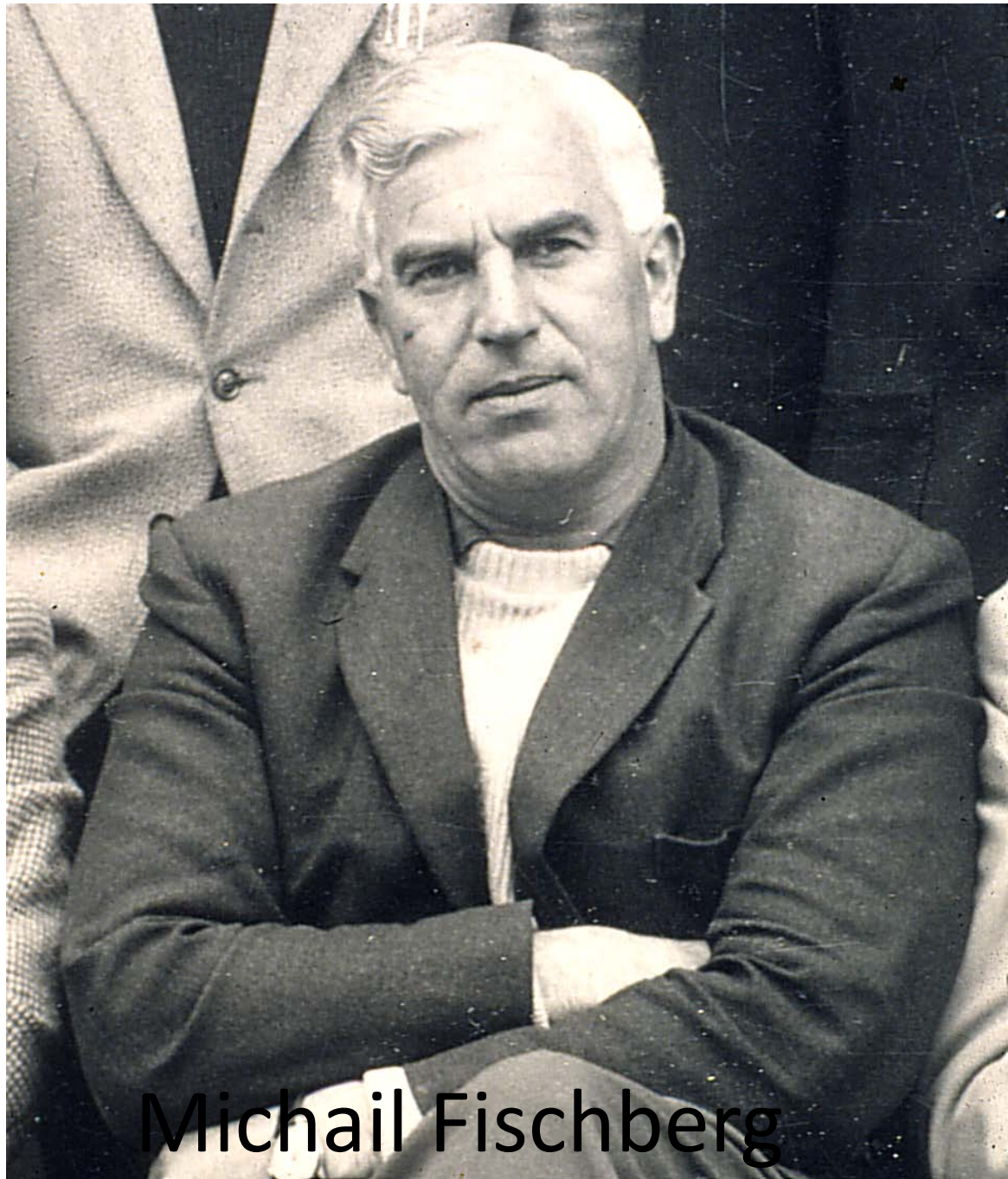
PERCENTAGE OF COMPLETE BLASTULA TRANSPLANT-
EMBRYOS WHICH DEVELOP NORMALLY



Briggs and King, 1952. Proc. Nat. Acad. Sci., USA

Cloned adult vertebrate (1958)





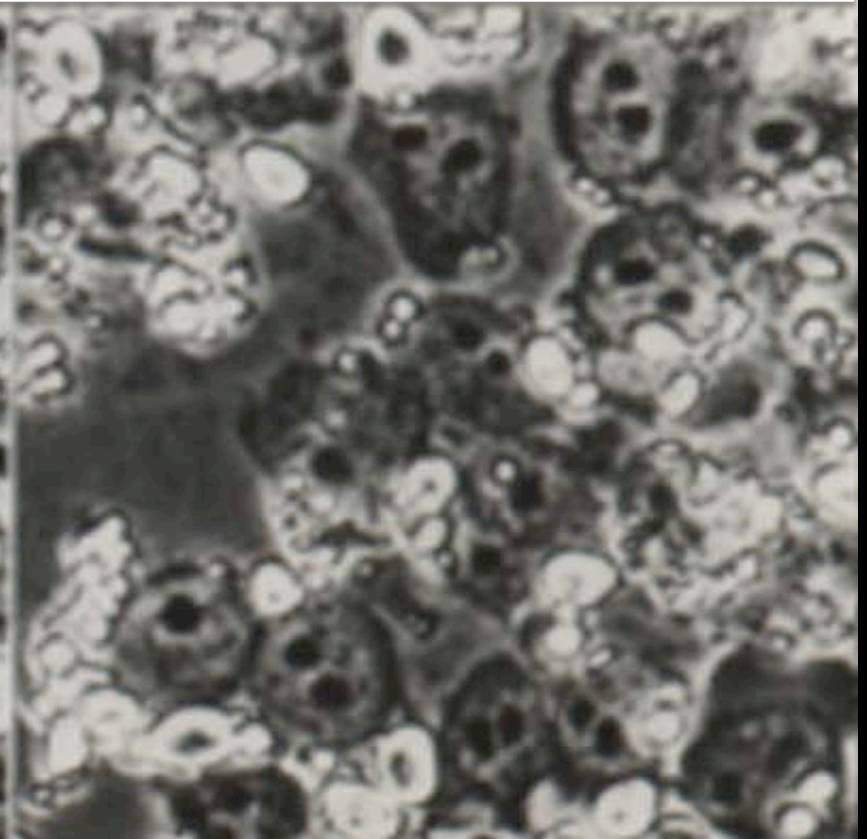
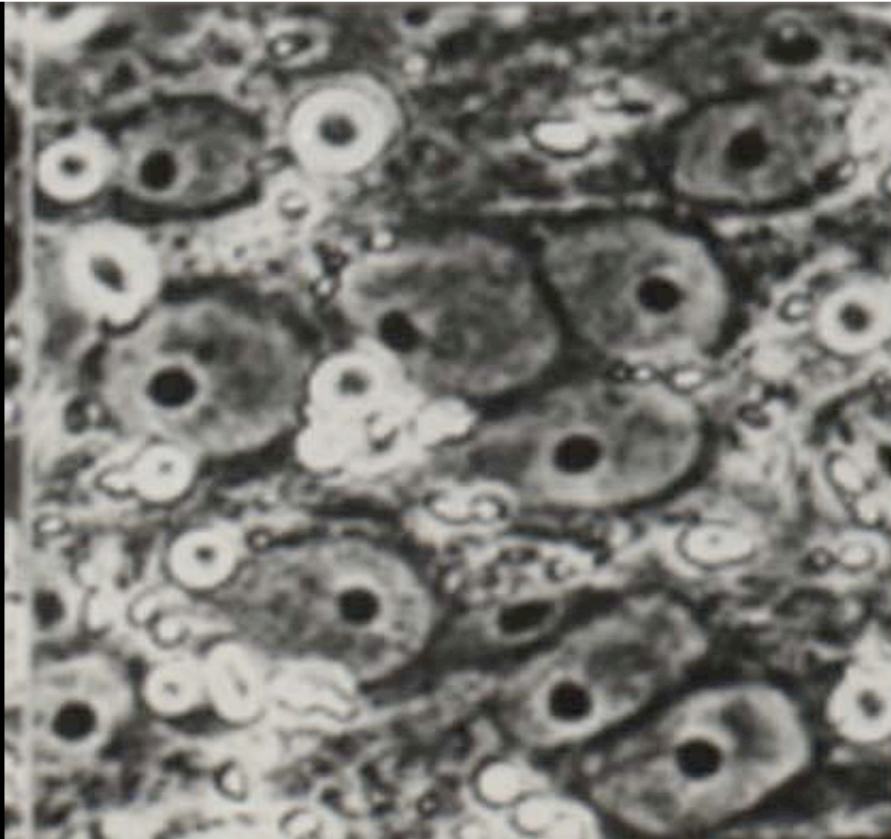
Michail Fischberg



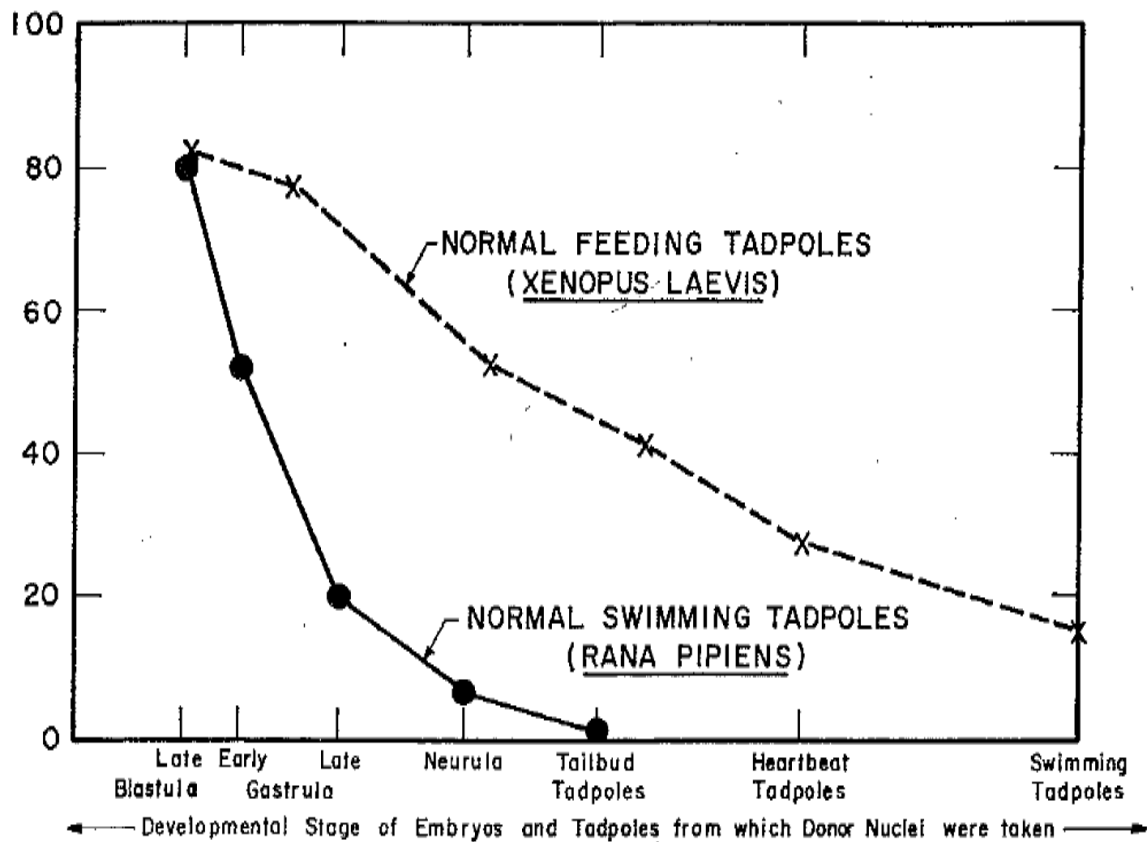
1-nucleolus



2-nucleolus



PERCENTAGE OF COMPLETE BLASTULA TRANSPLANT-
EMBRYOS WHICH DEVELOP NORMALLY

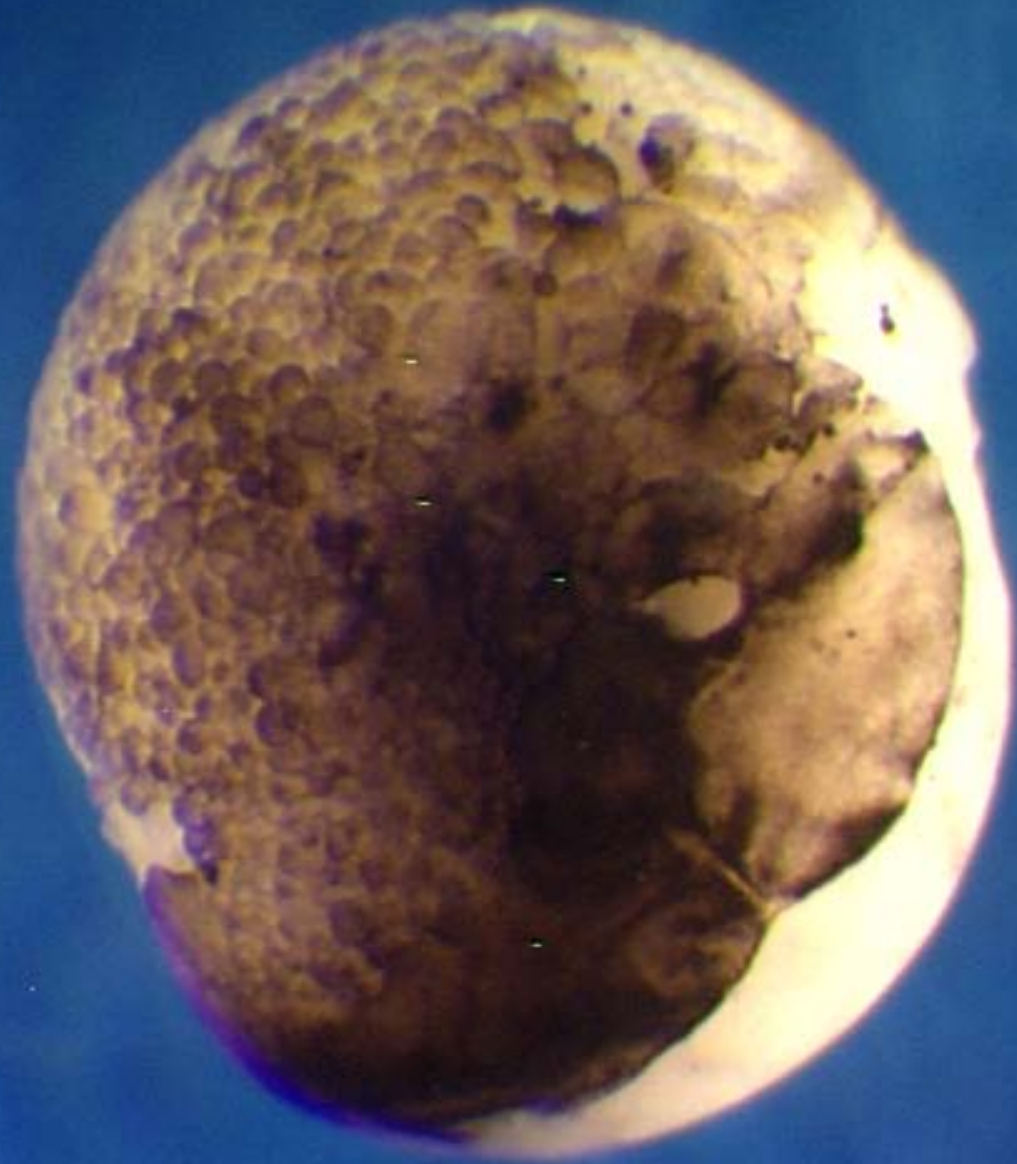


Intestinal Tract of Feeding Tadpole

(GFP-marked)

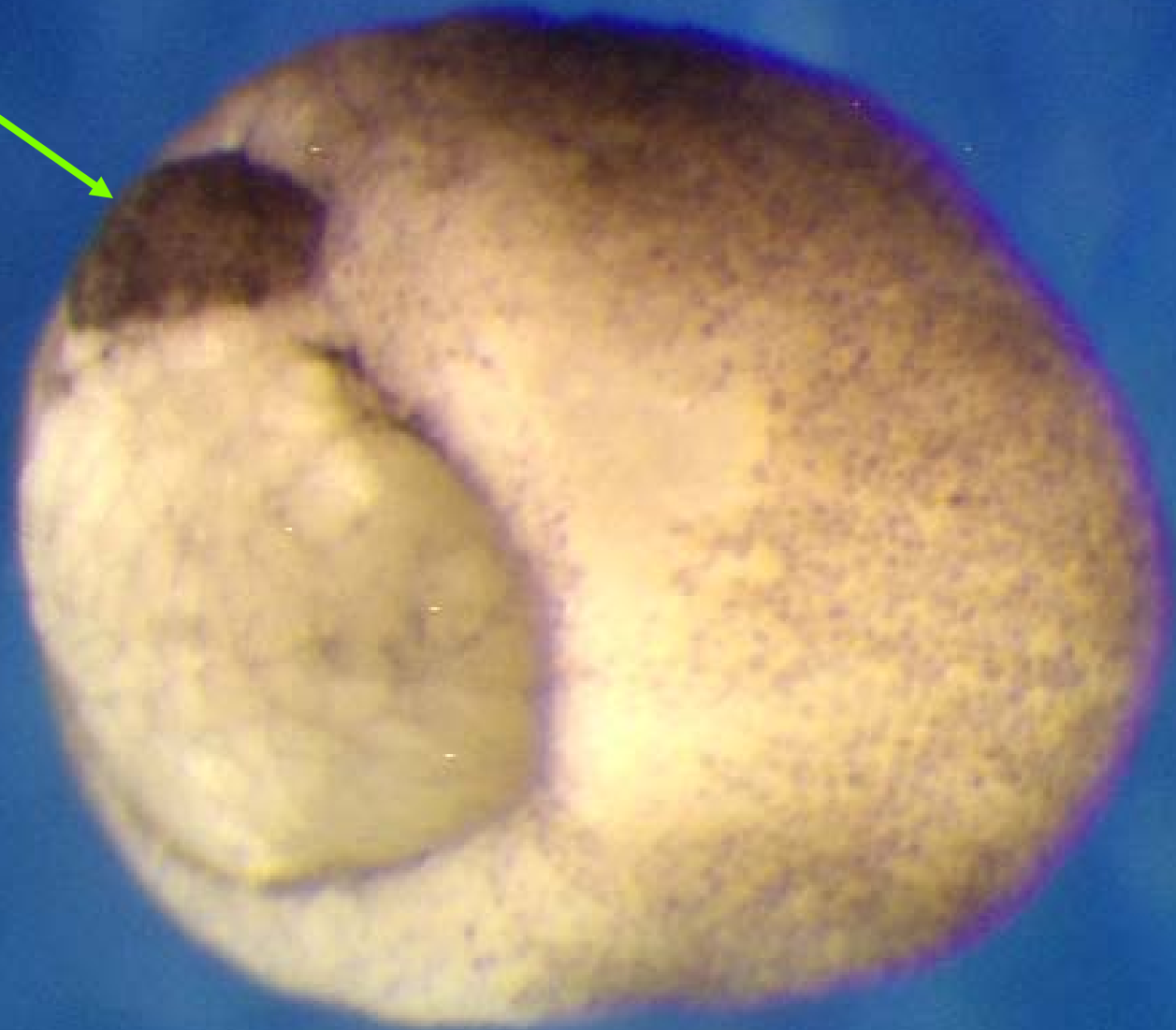


Partial blastula

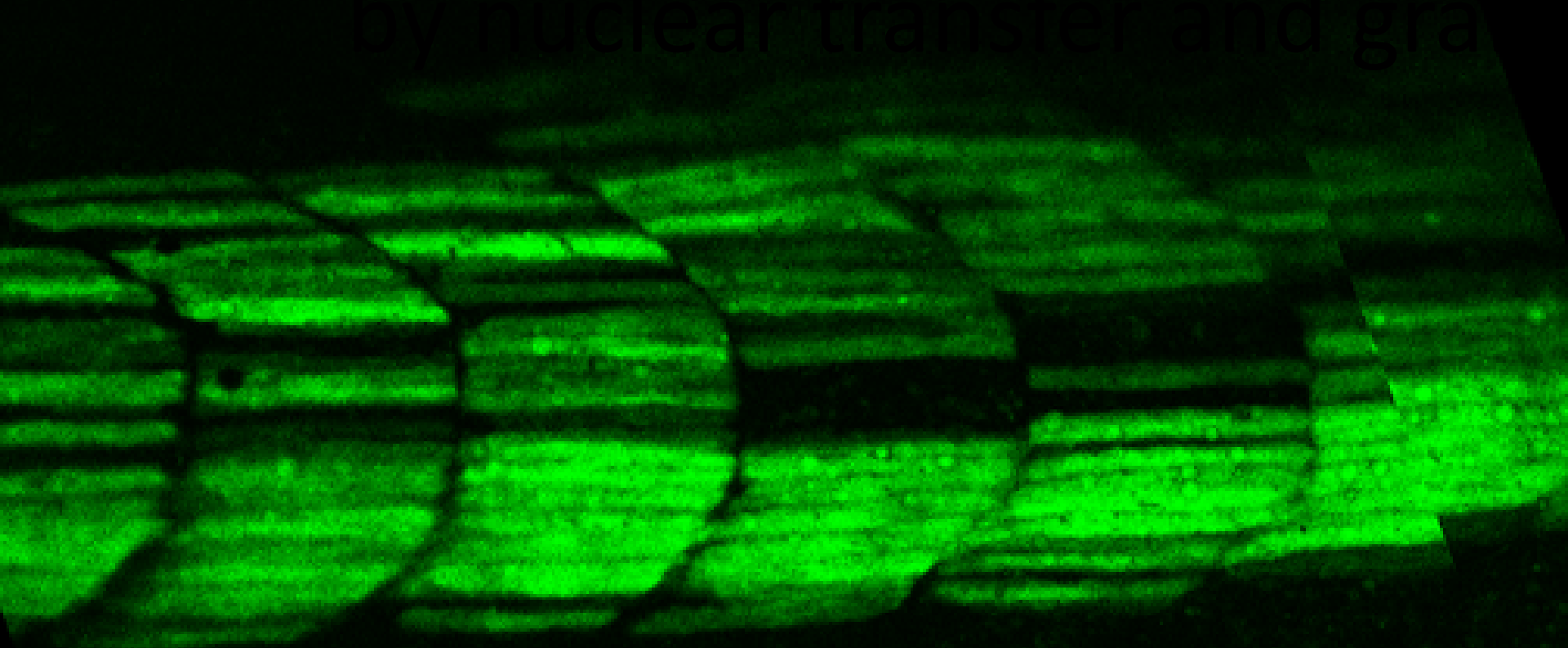


Nuclear Transplant Embryo Graft

GFP



GFP-muscle derived from intestine nuclei



Efficiency of nuclear reprogramming by nuclear transfer to eggs

Switch between cell-types:
e.g. intestinal epithelium to muscle and nerve.

First nuclear transfers.....15%
Serial nuclear transfers.....7%
Grafts from nuclear transfer embryos.....8%

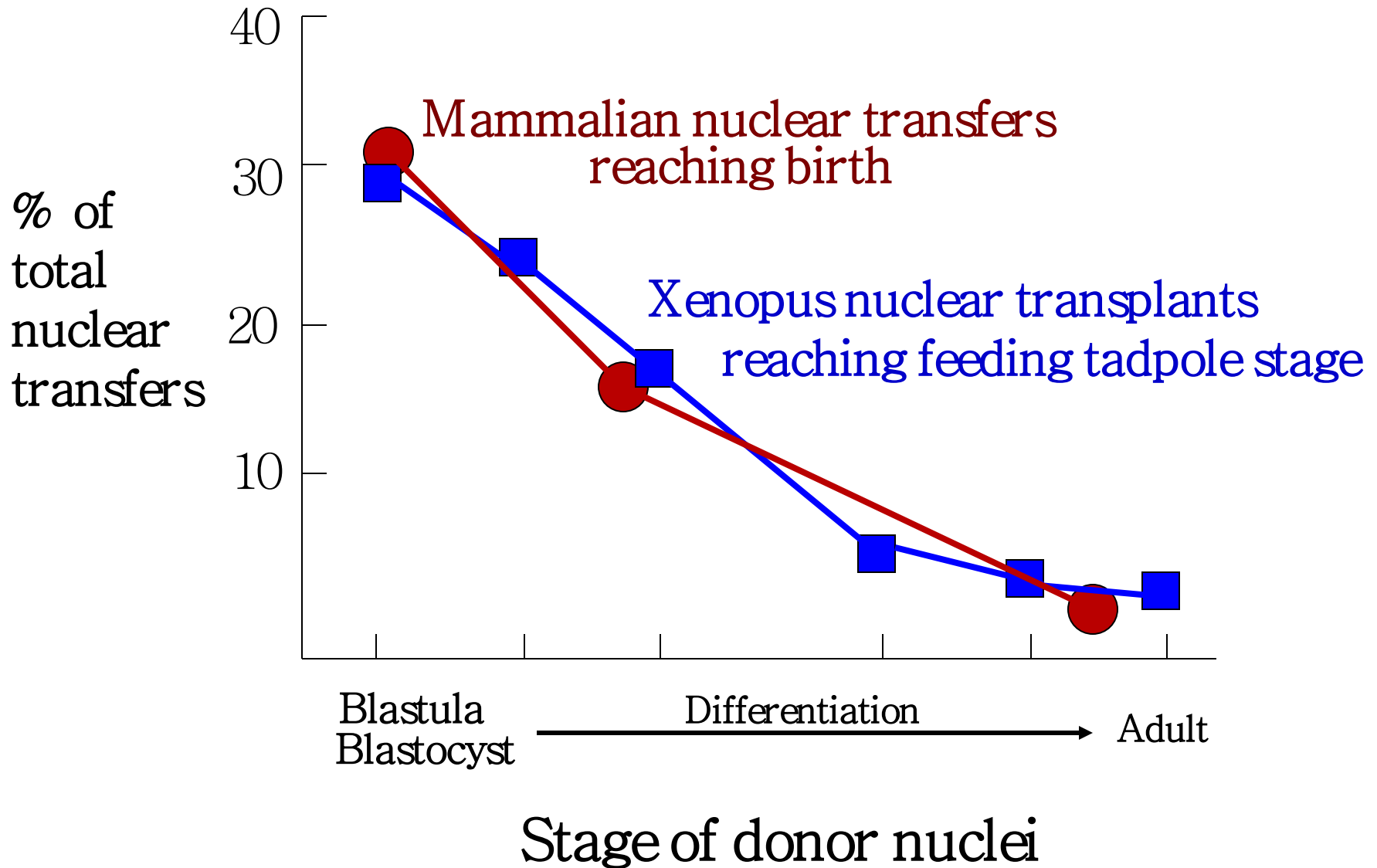
Total: 30%

The cloning of sheep

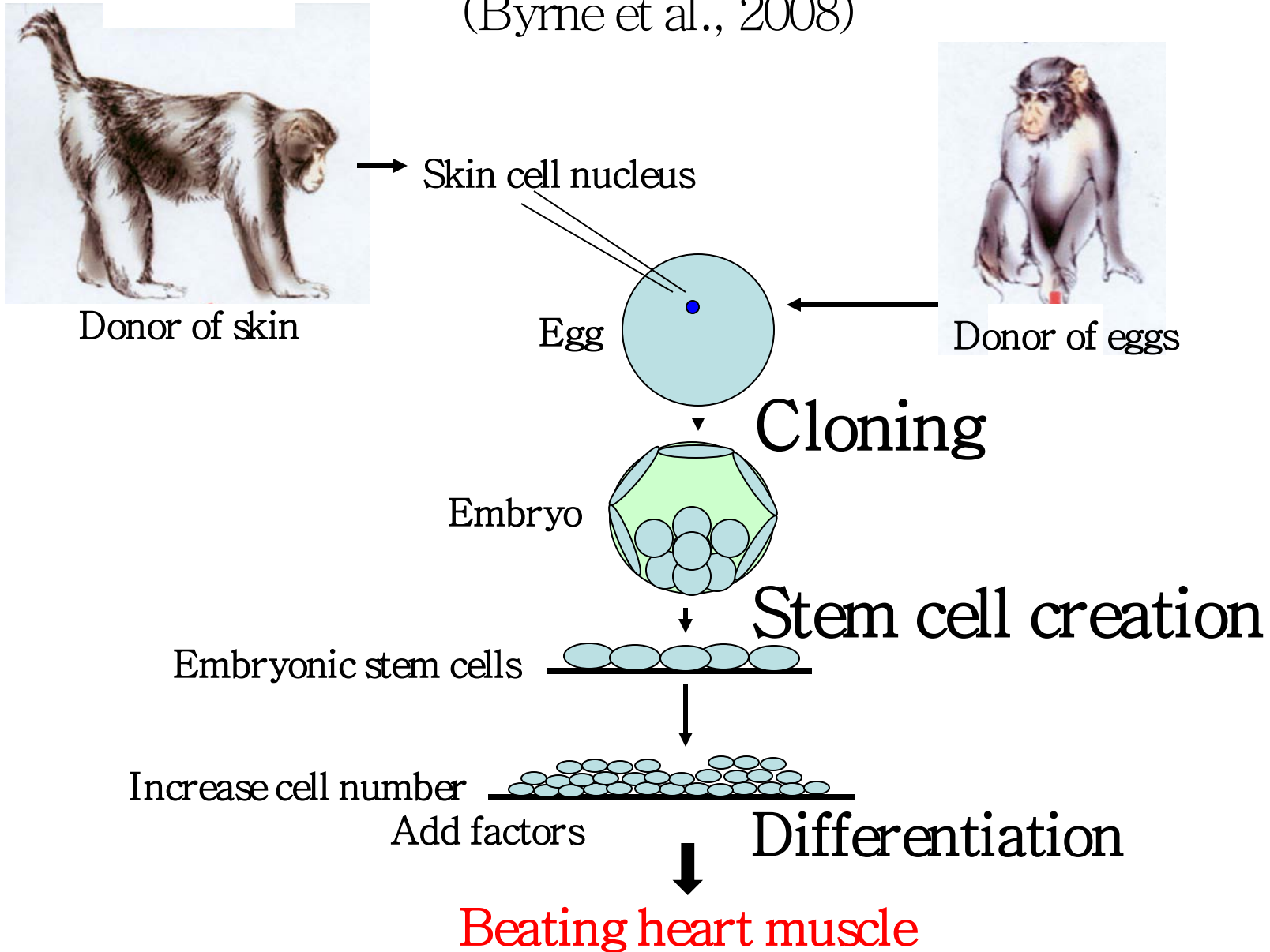


Wilmut, Campbell et al 1996 and 1997.

Nuclear transfer success decreases as donor cells differentiate



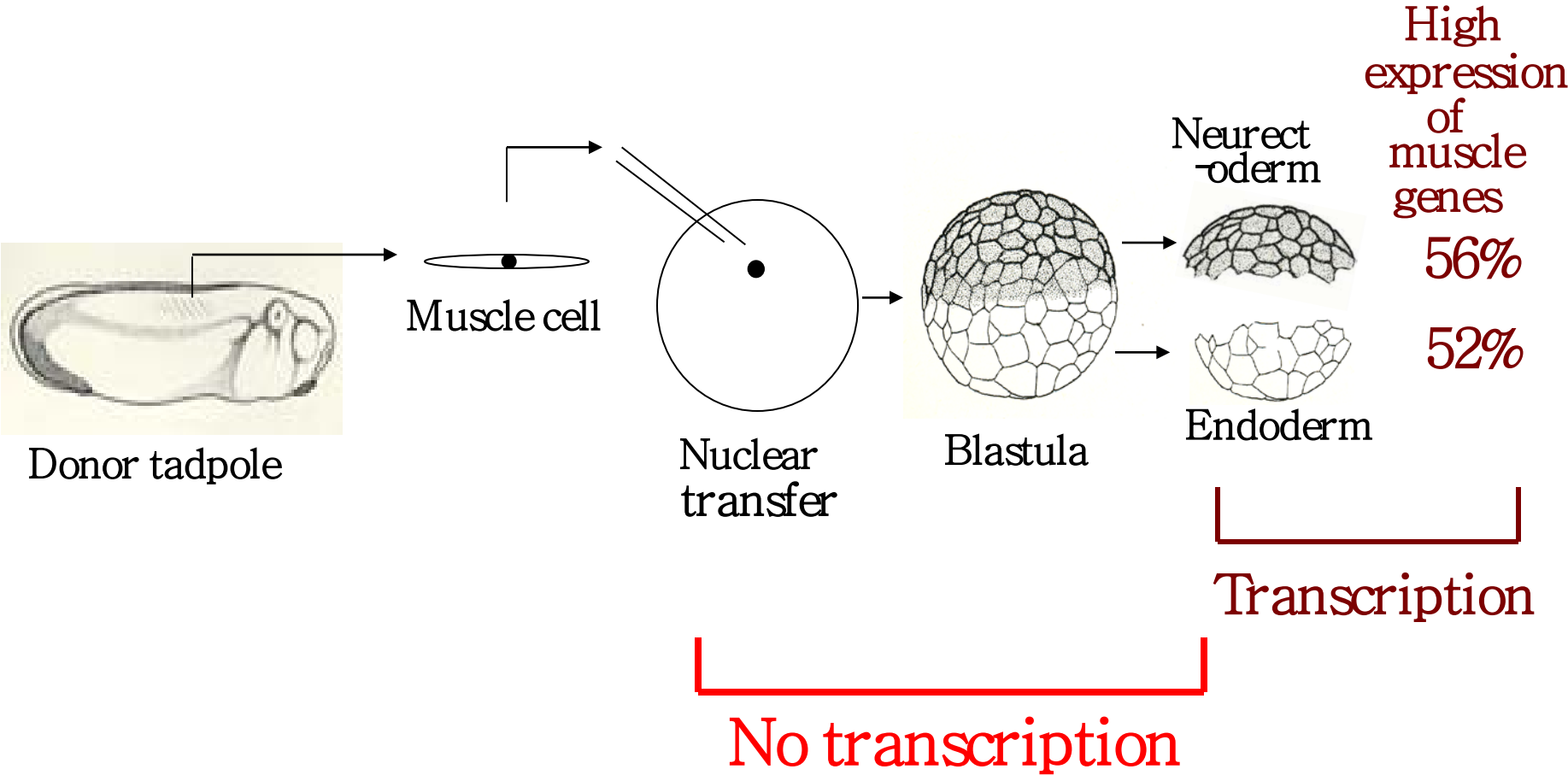
Derivation of functional heart from adult monkey skin (Byrne et al., 2008)



Differentiation of CRES-2 cells into cardiac tissues

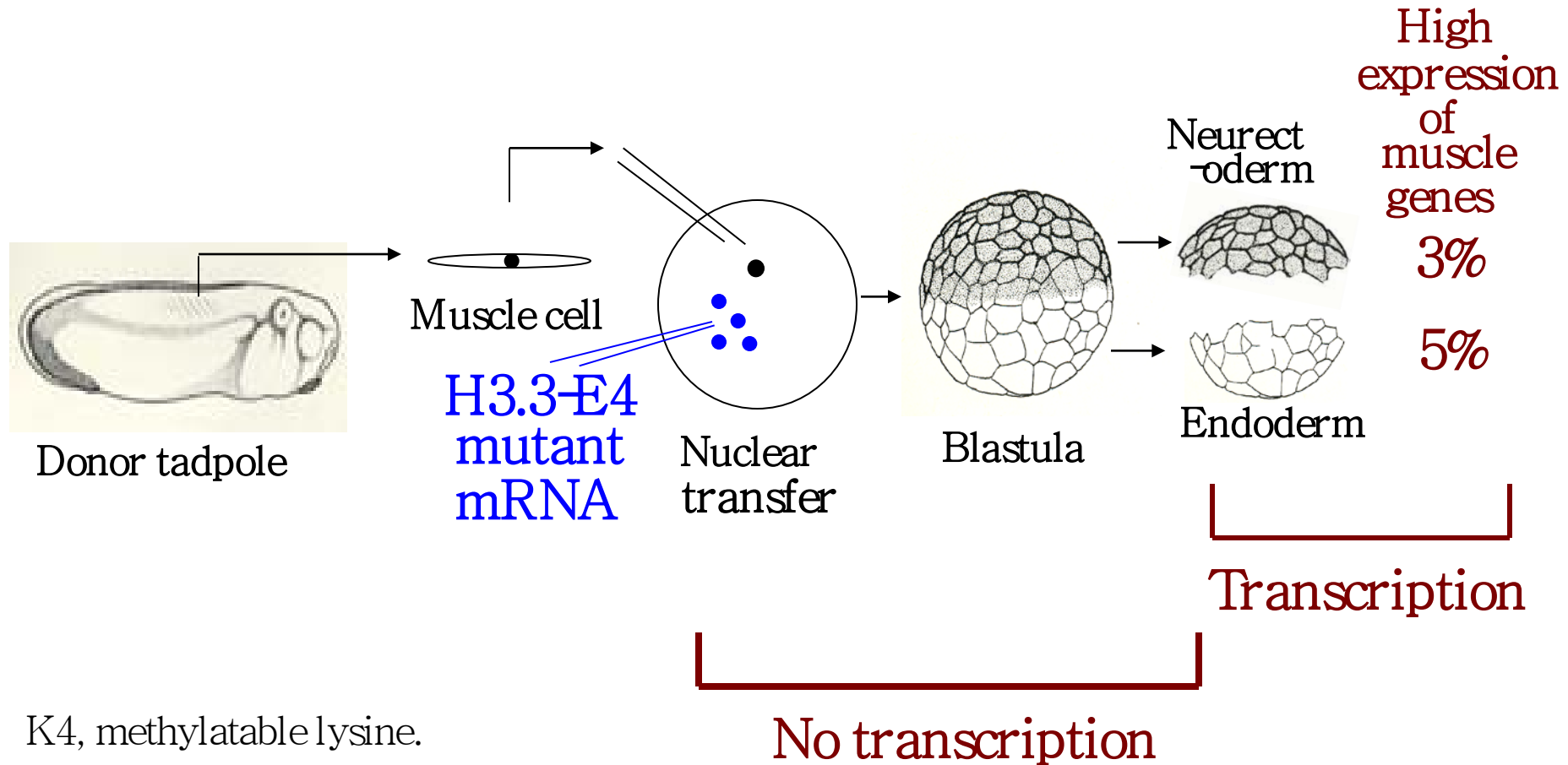
Epigenetic memory

Embryos derived from muscle nuclei remember their origin even in their nerve and endoderm cells



H3.3 is required for epigenetic memory.

Elimination by H3.3 mutated from K4 to E4.



K4, methylatable lysine.

E4, glutamine

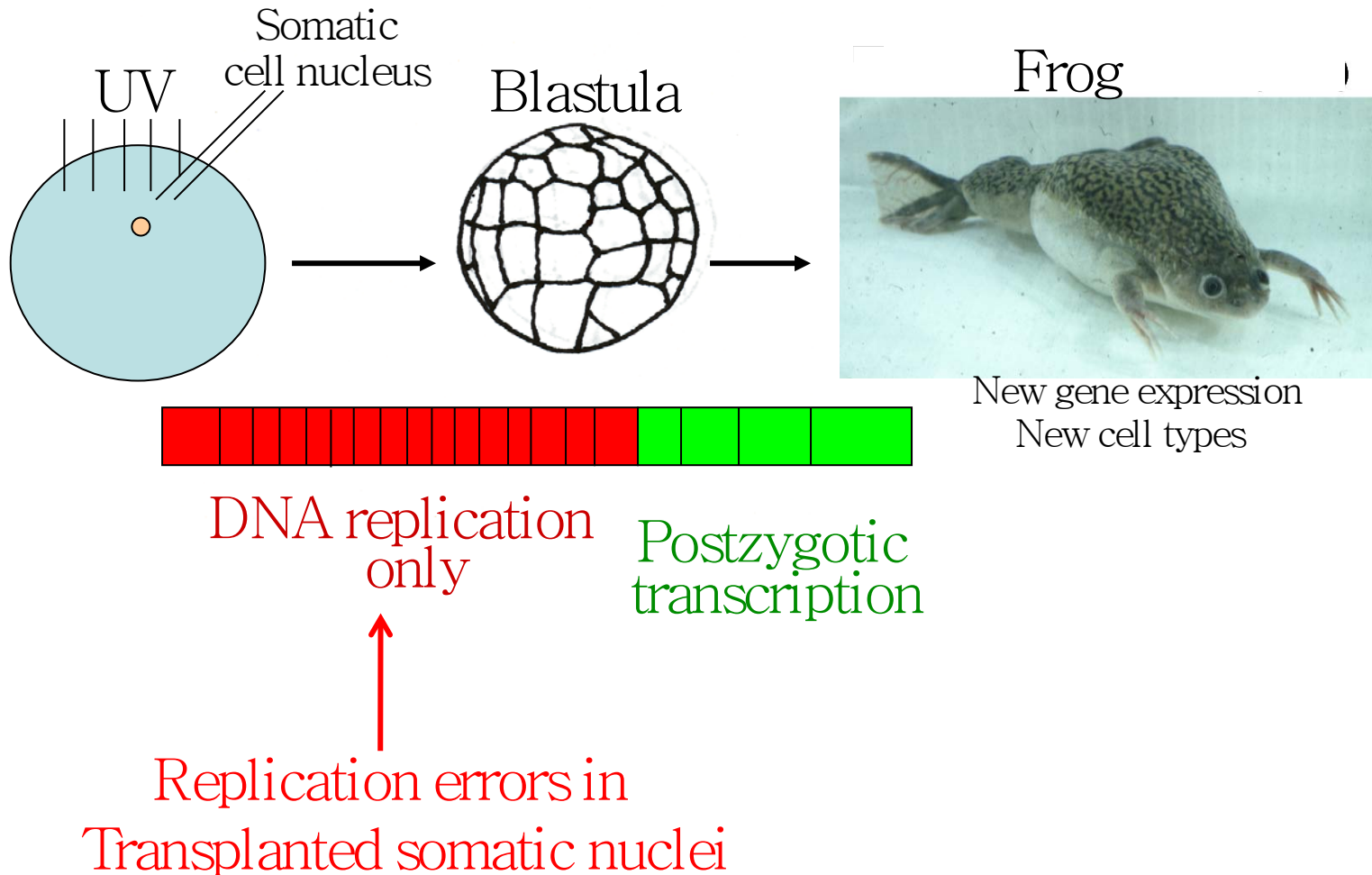
EPIGENETIC MEMORY

Can be explained if:

1. H3.3 promotes continuing transcription of active genes, and if
2. Egg cytoplasm reverses gene transcription with a 50% efficiency.

First meiotic prophase oocytes
to analyse the mechanism of nuclear reprogramming

Single nuclear transfer to eggs in second meiotic metaphase

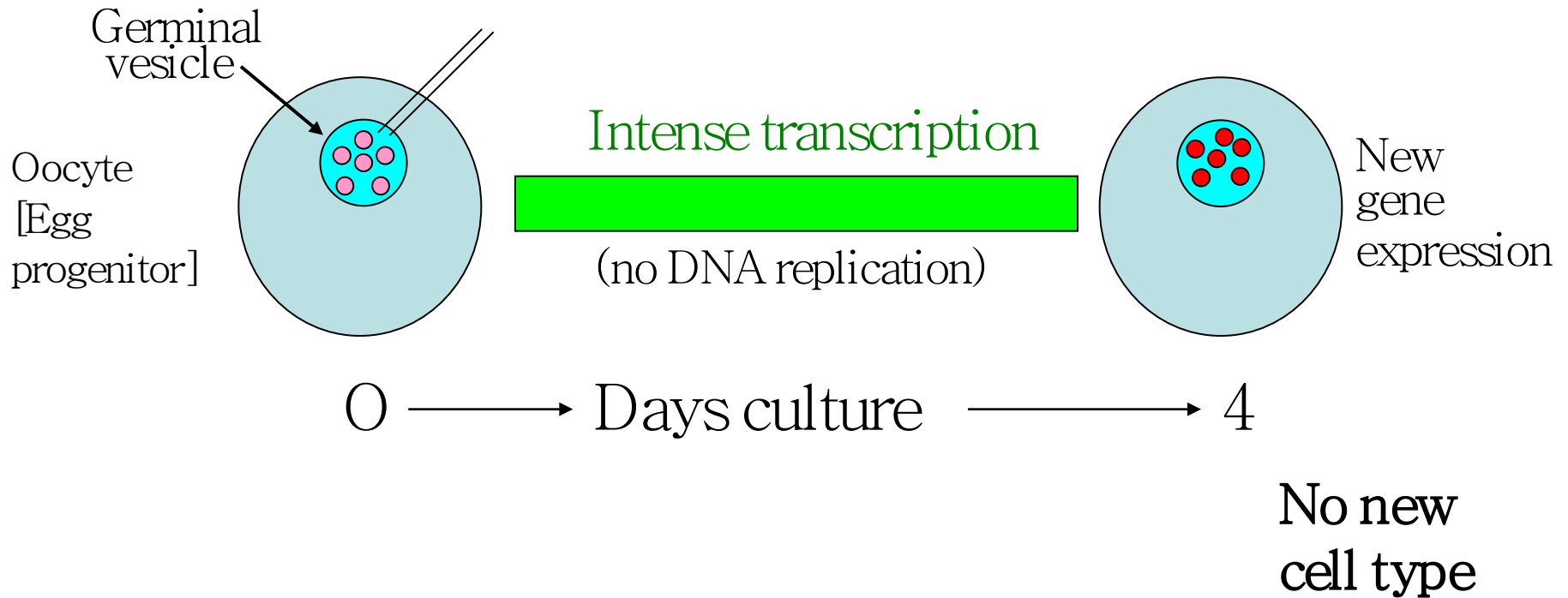


Incomplete DNA replication

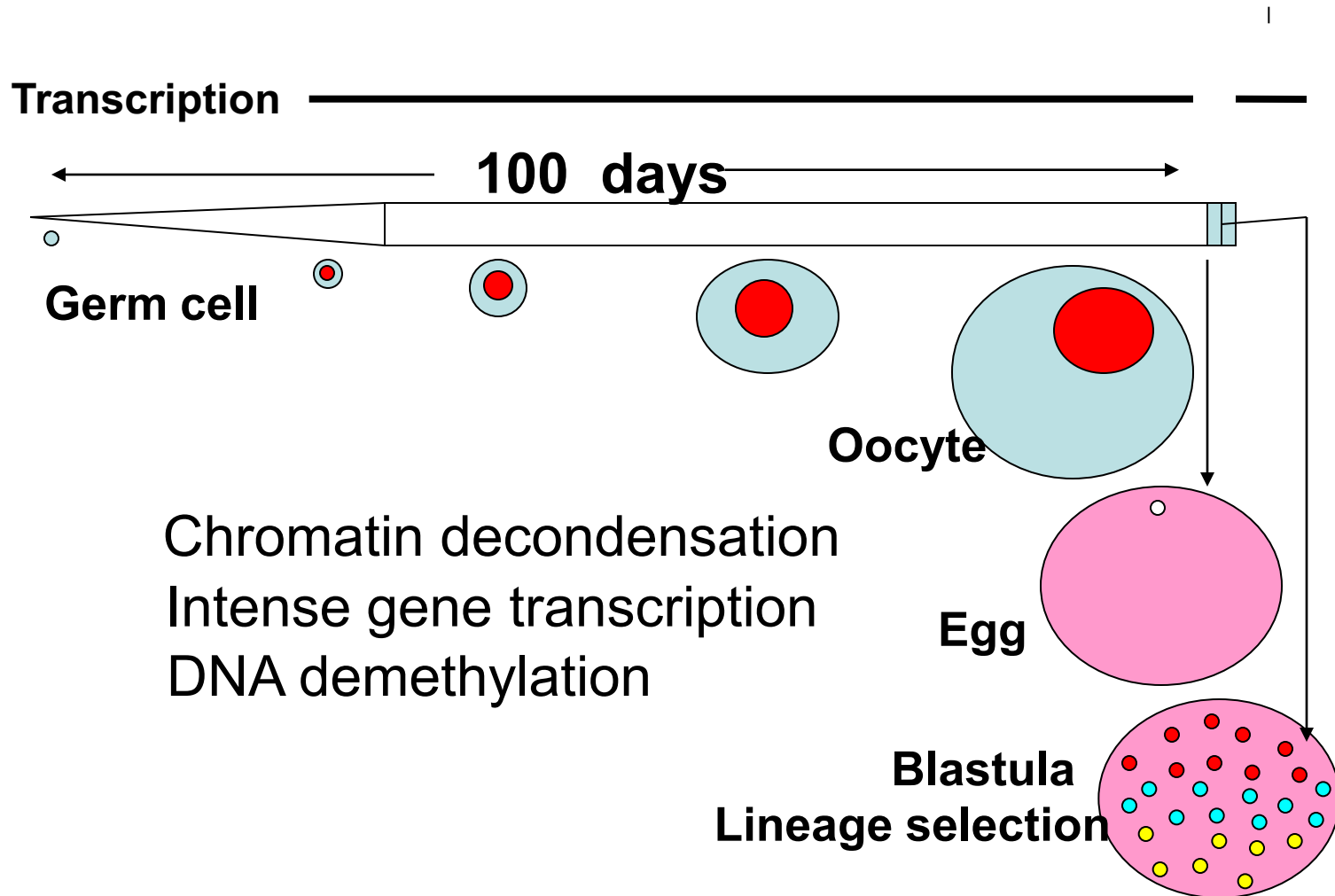
damages somatic nuclei

transplanted to eggs

Multiple nuclei transferred to growing oocytes in first meiotic prophase



Oocyte formation prepares the egg for development



Xenopus oocyte and germinal vesicle

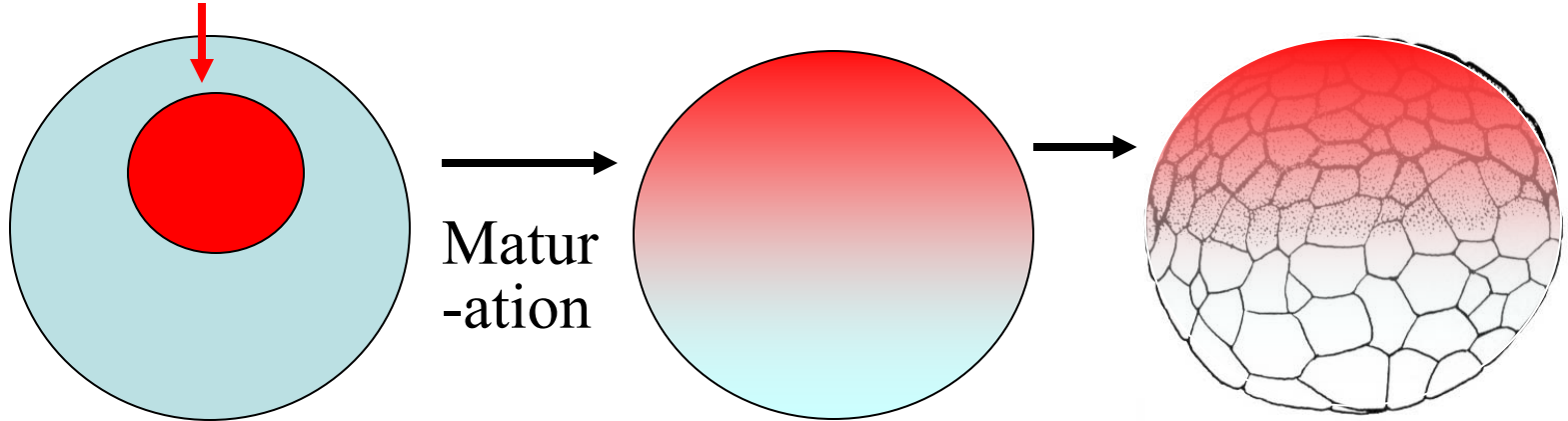
GV

1 mm



The oocyte germinal vesicle contents contribute to post-fertilization development

Germinal vesicle



Oocyte

First
meiotic
prophase

Egg

Second
meiotic
metaphase

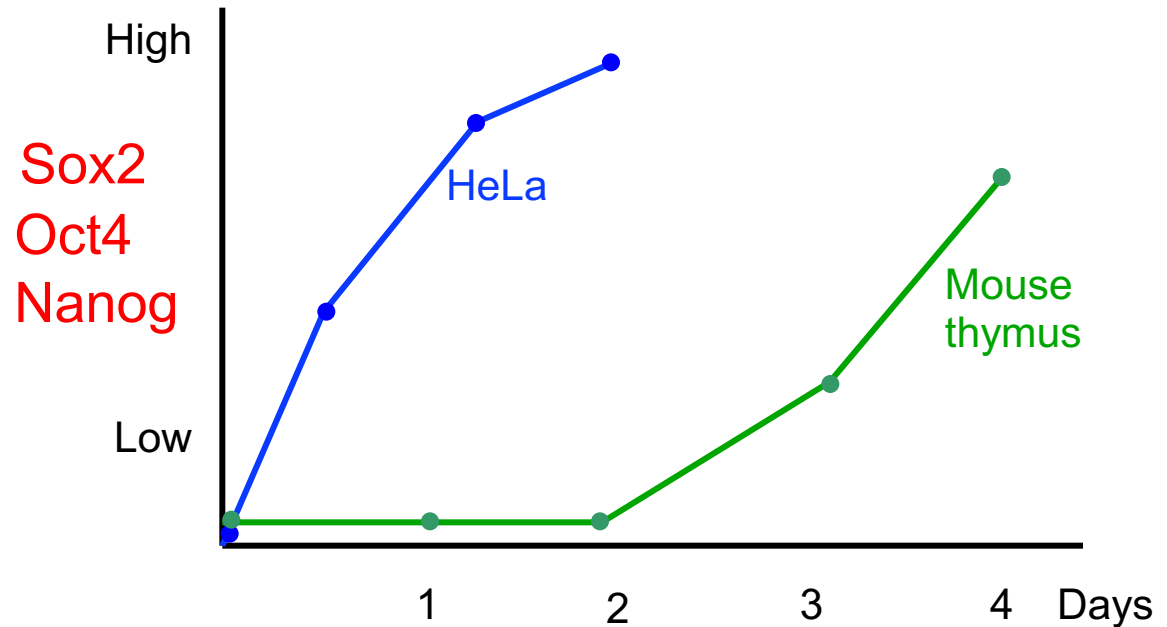
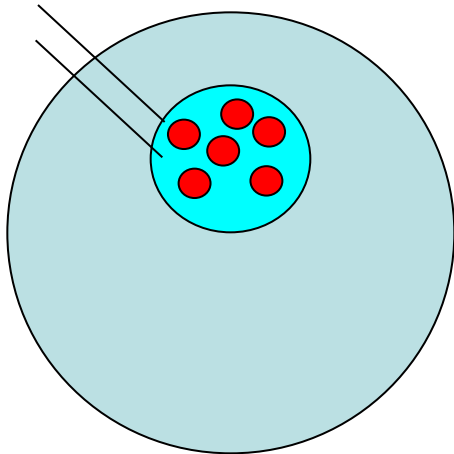
Embryo

Mid blastula

Mammalian stem cell genes are rapidly activated in mammalian nuclei transplanted to *Xenopus* oocytes

Nuclei of differentiated cells are reprogrammed slowly.

Mouse/human
somatic cell
nuclei



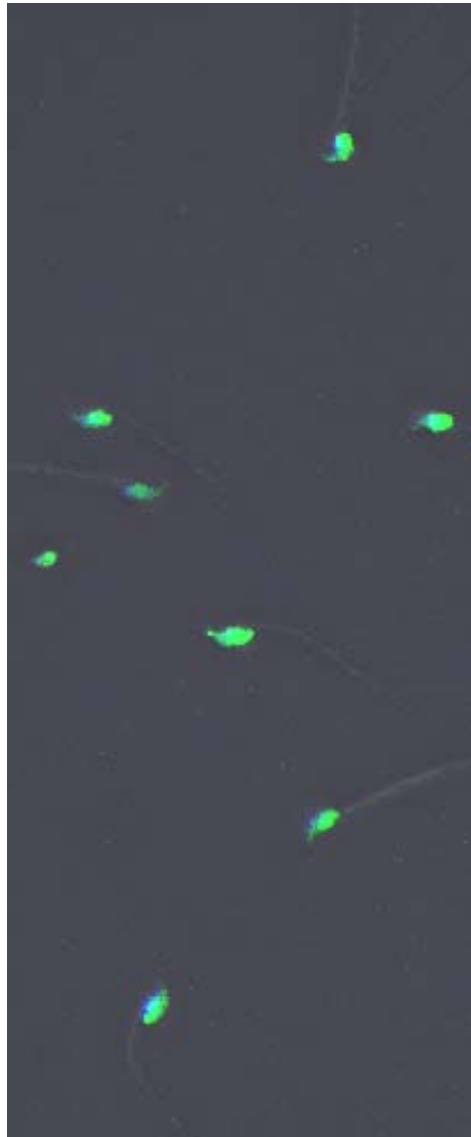
Oocyte transcription assay

Living oocyte transcription assay

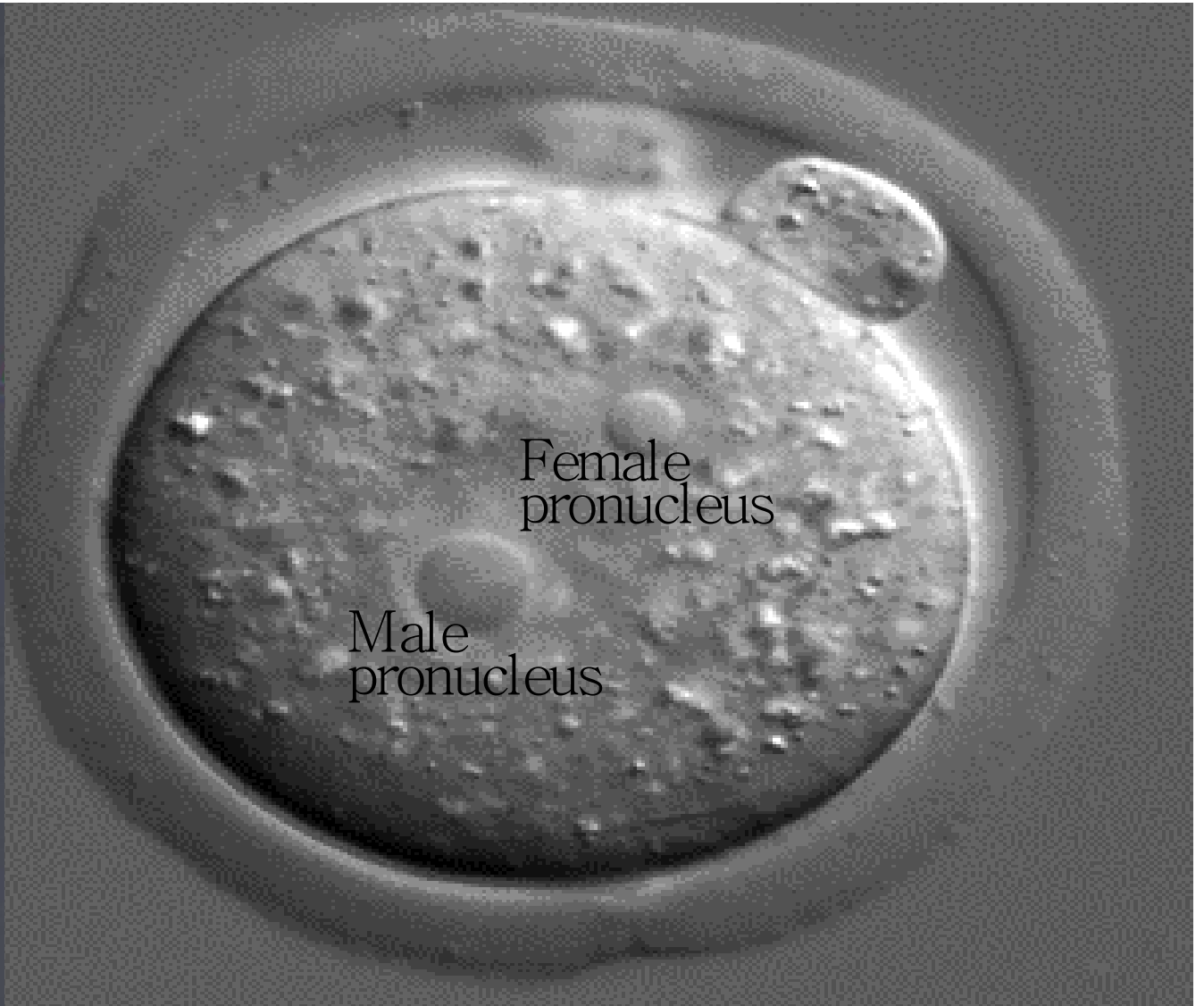
1. Multiple somatic nuclei in one oocyte.
2. Linear accumulation of new transcripts.
3. Multiple initiations of transcription per gene per day.
4. Oocyte injections show resistance

Transcriptional activation:
attack by egg cytoplasm

Mouse sperm

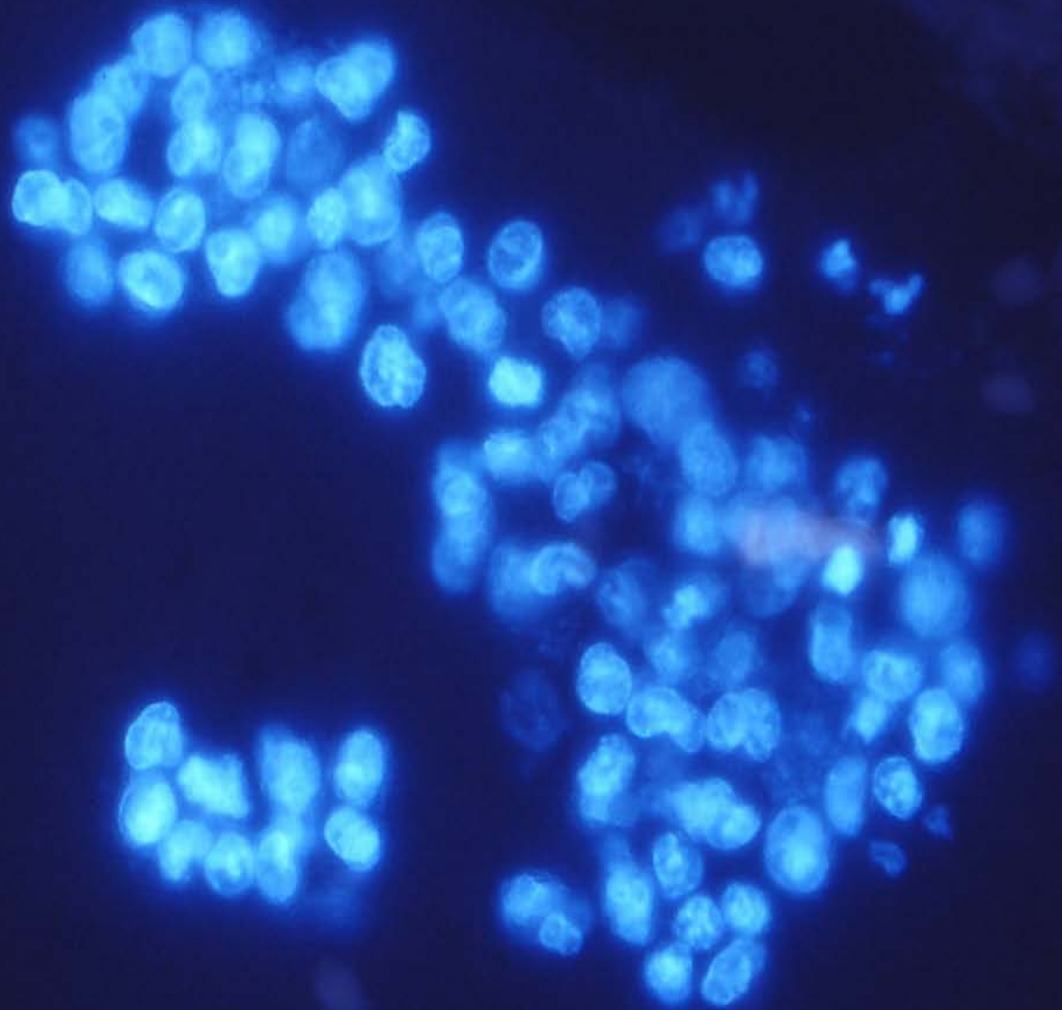


Fertilized mouse egg

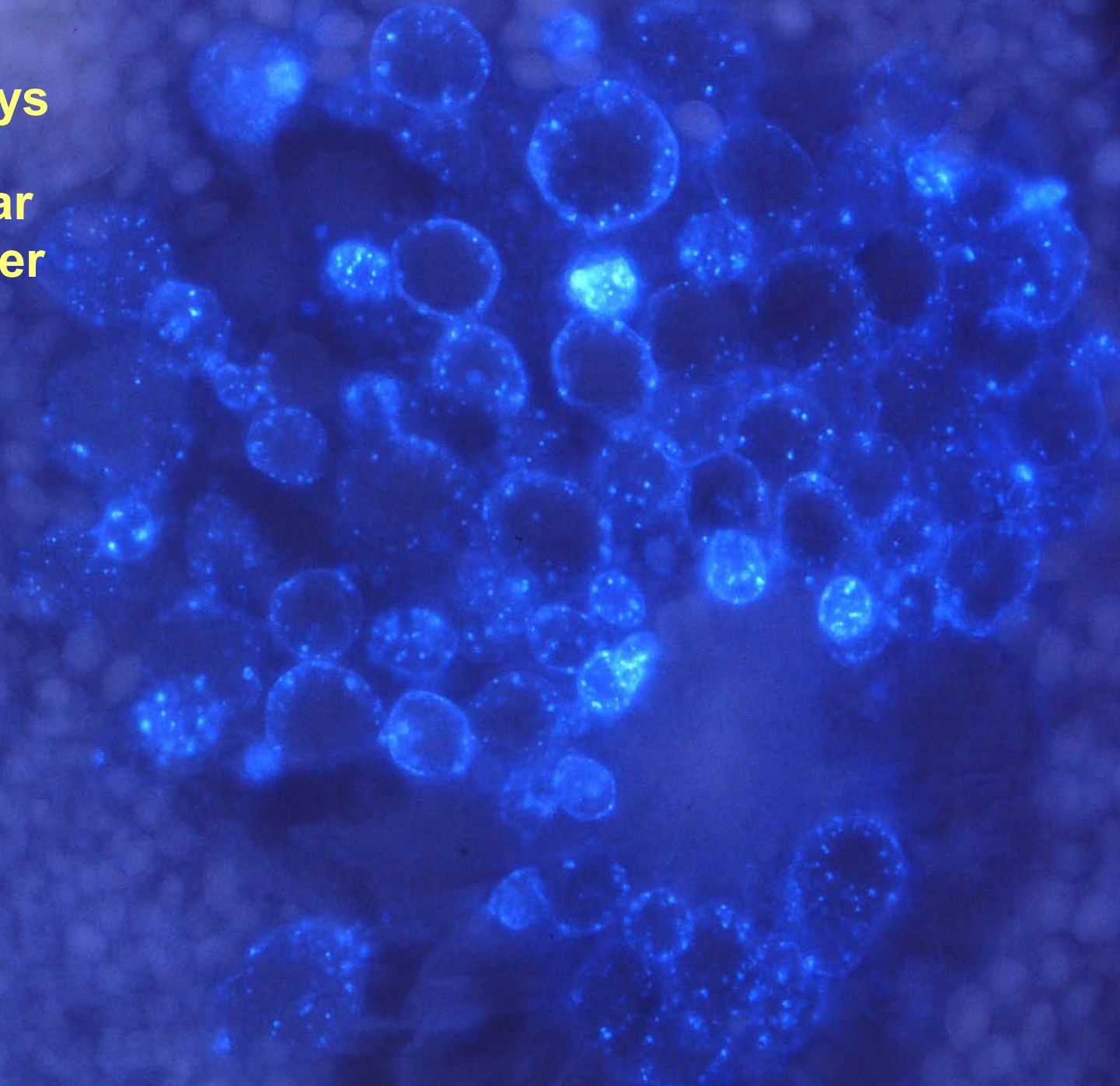


Mammalian
cultured cell
nuclei:

Just after
injection



**1-2 days
after
nuclear
transfer**

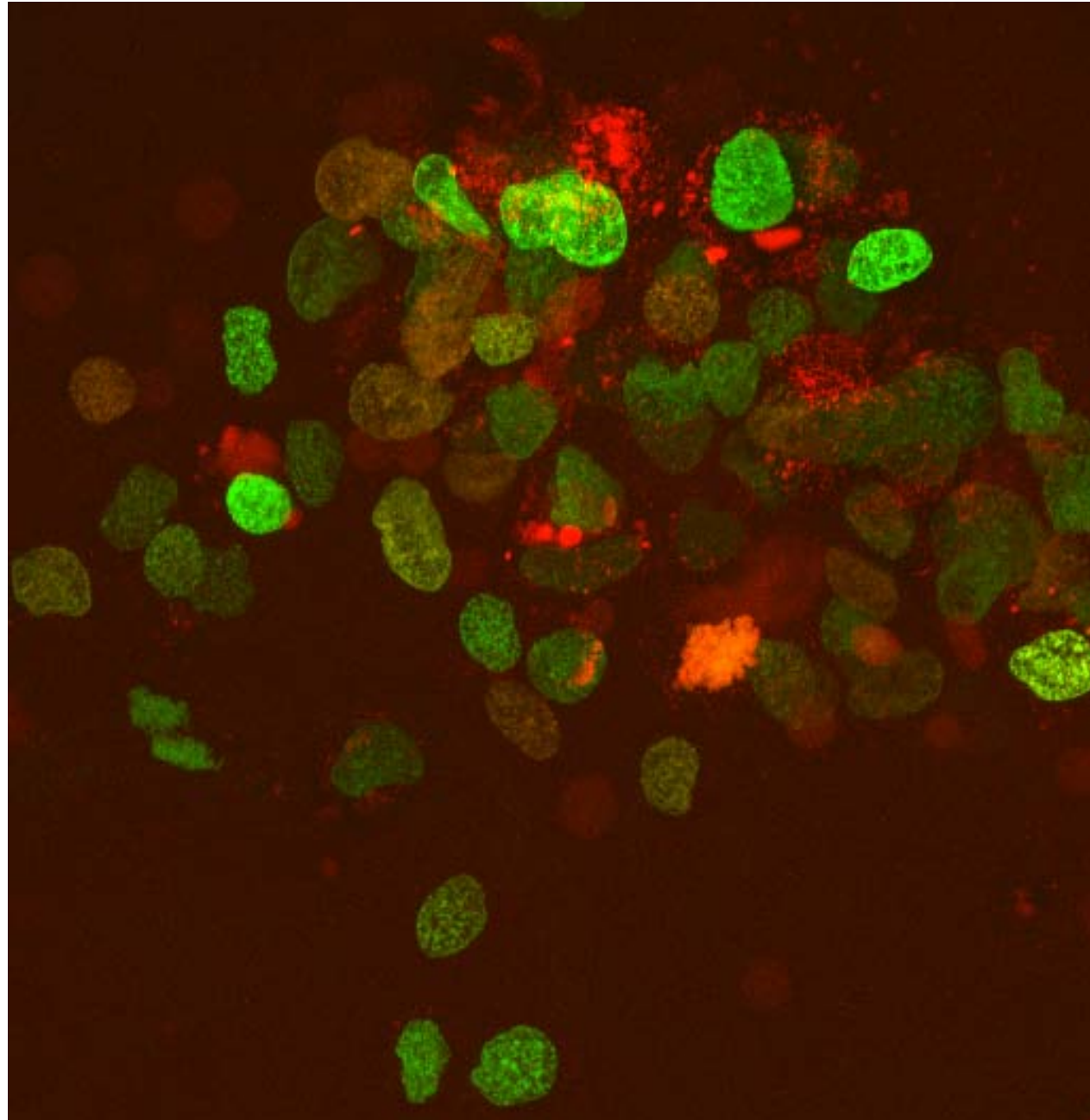


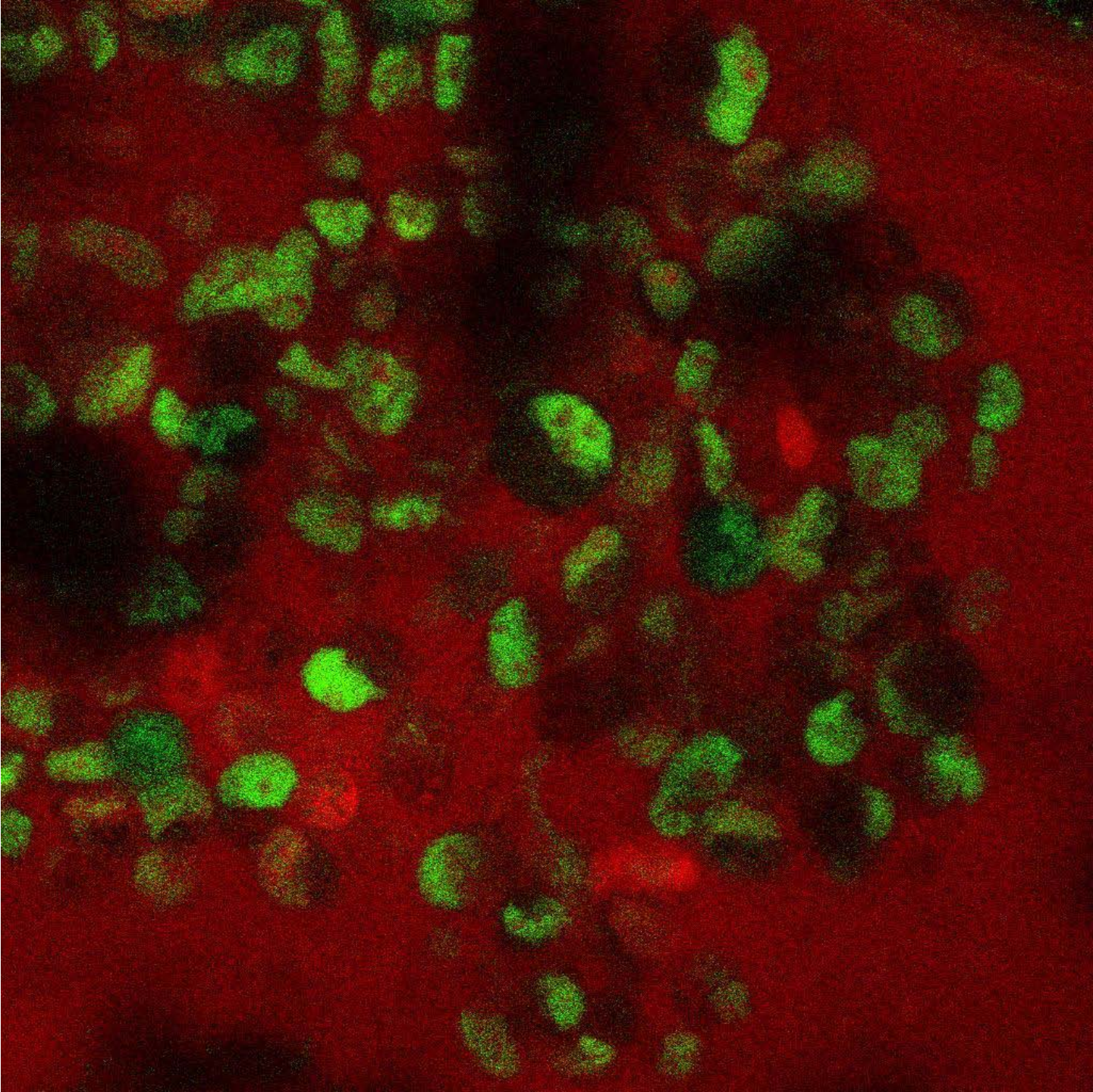
Histone replacement in transplanted nuclei.

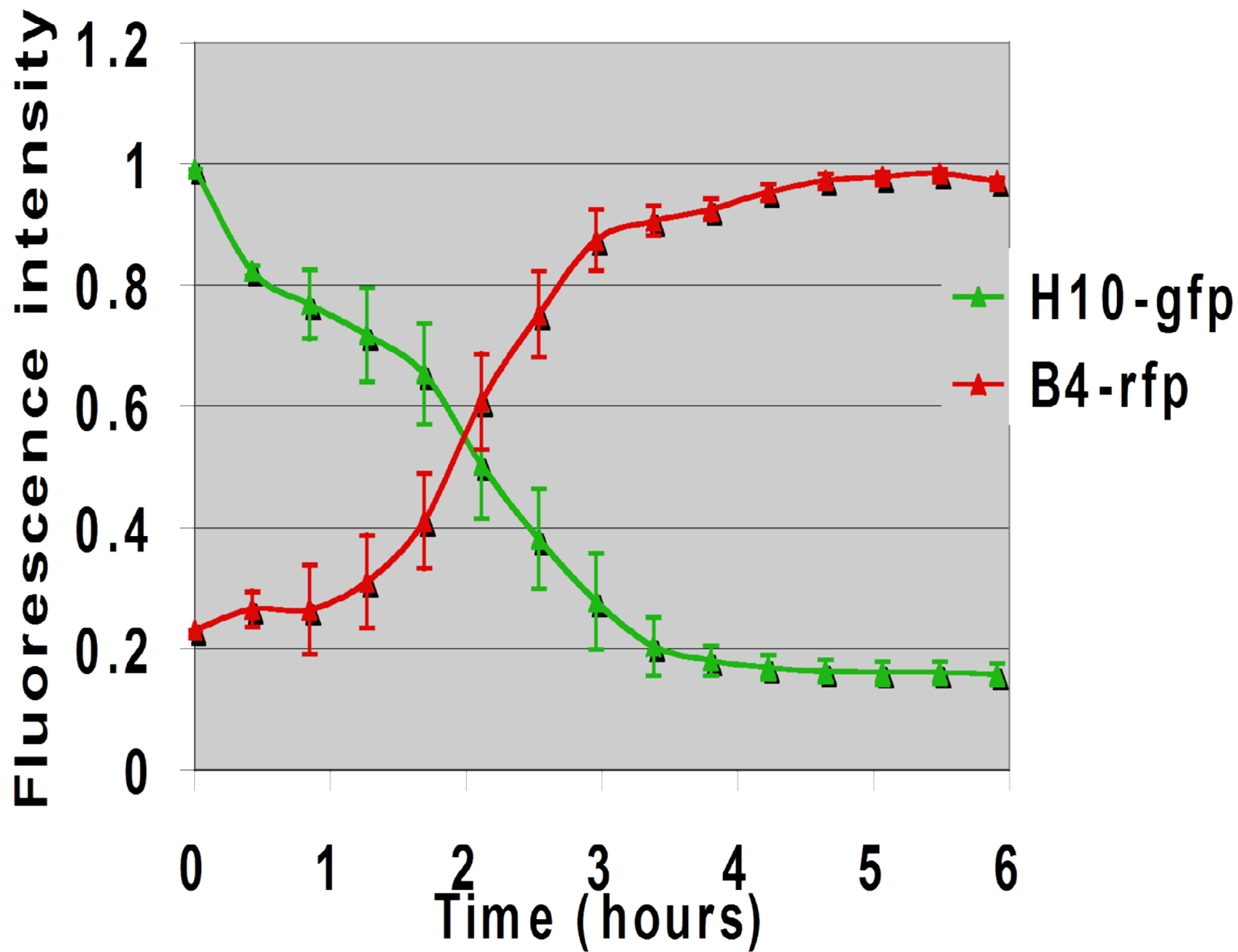
Cultured cell
nuclei in
oocyte GV

Somatic
H1o histone-GFP
replaced by
oocyte
B4 histone-RFP

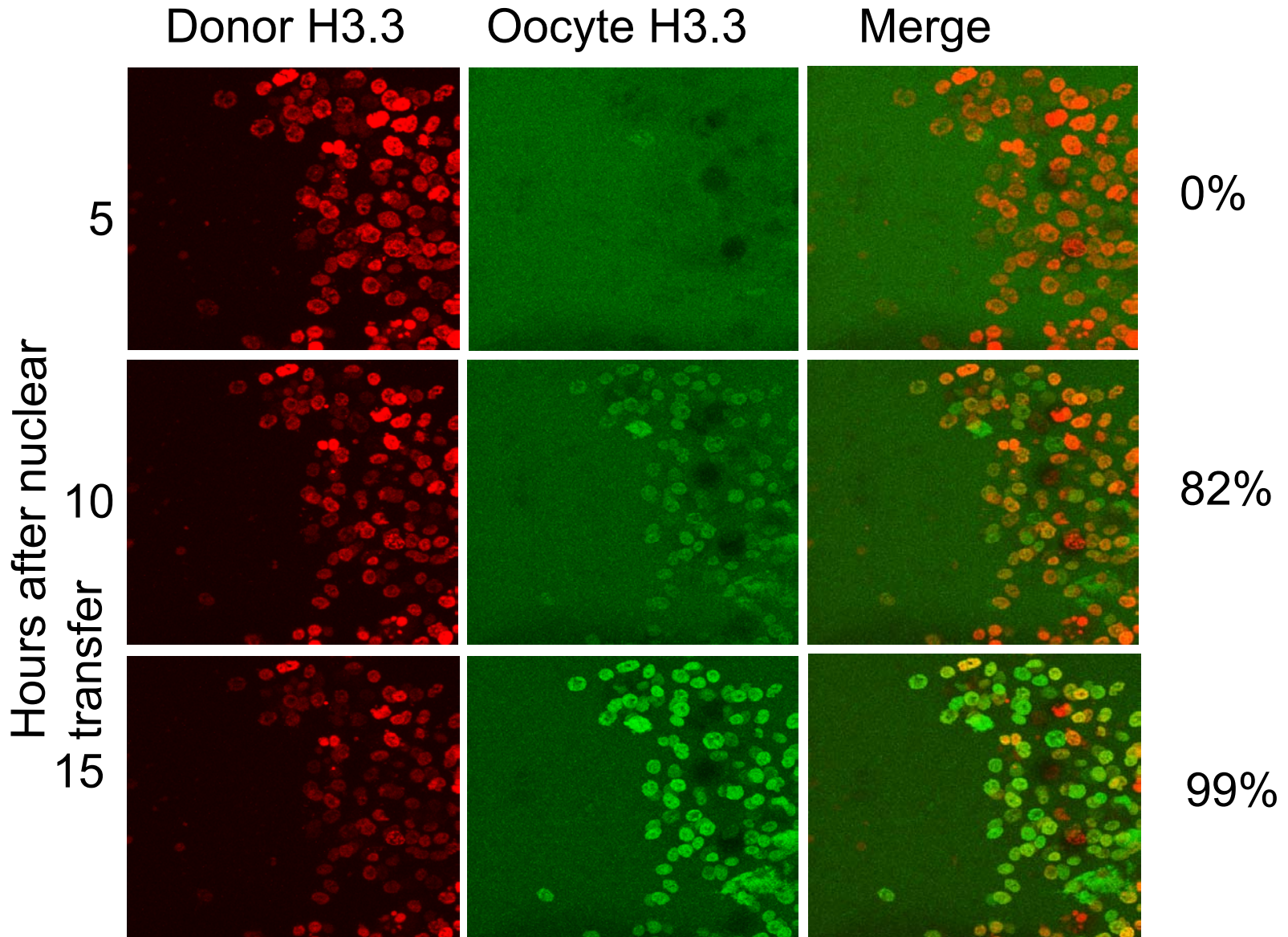
Jerome Jullien



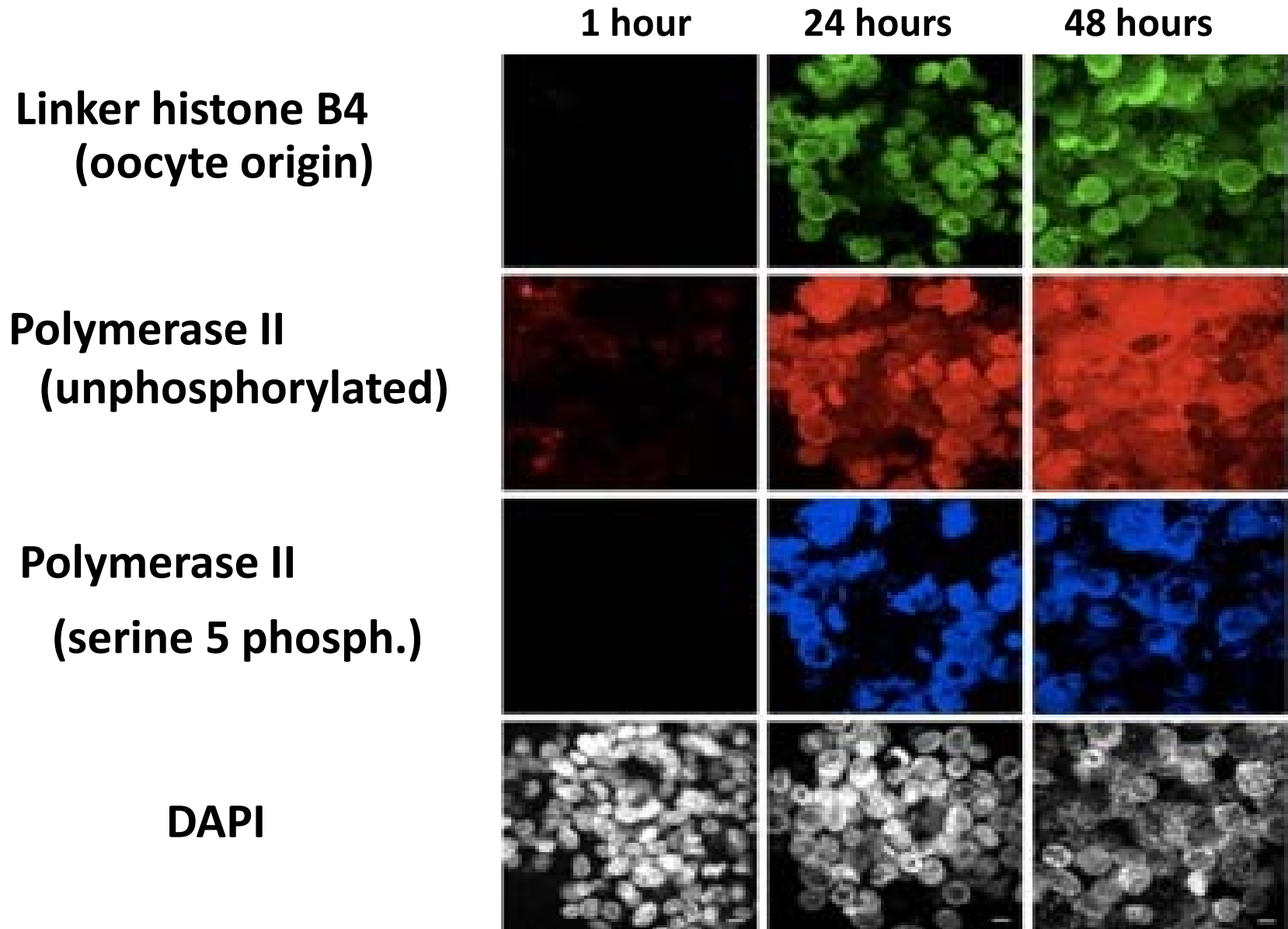




Histone H3.3 is incorporated into transplanted nuclei

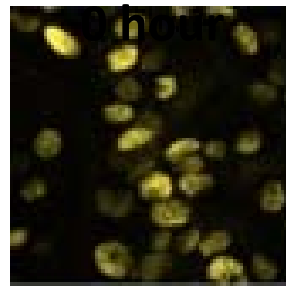


Uptake of linker histone and pol II correlates with reprogramming



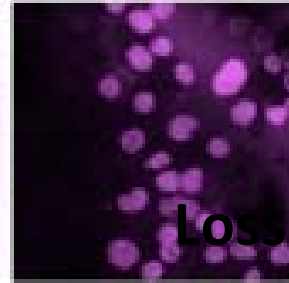
Uptake and loss of chromosomal proteins

YFP-RPB1
(RNA polymerase)



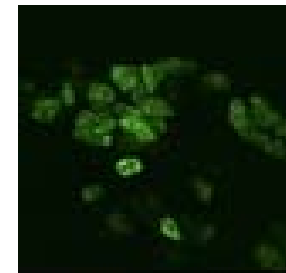
Loss

H2B-cherry
(core histone)



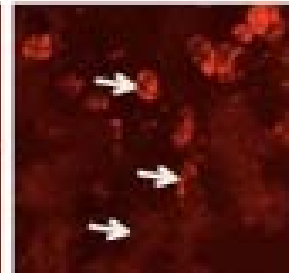
Gain

GFP-TBP
(TATA-binding protein of somatic cells)



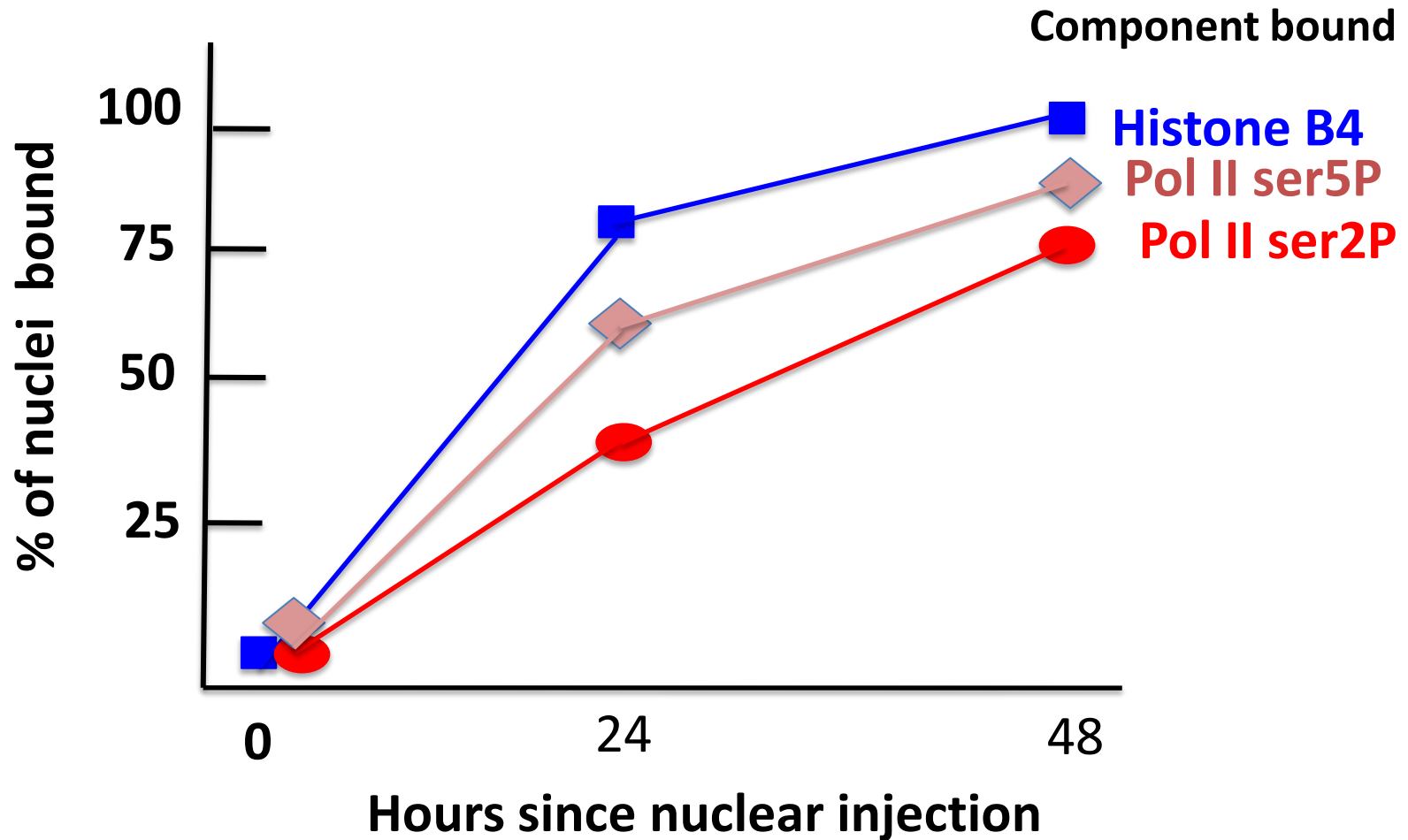
Loss

TBP2 -cherry
(TATA-binding protein of oocytes)

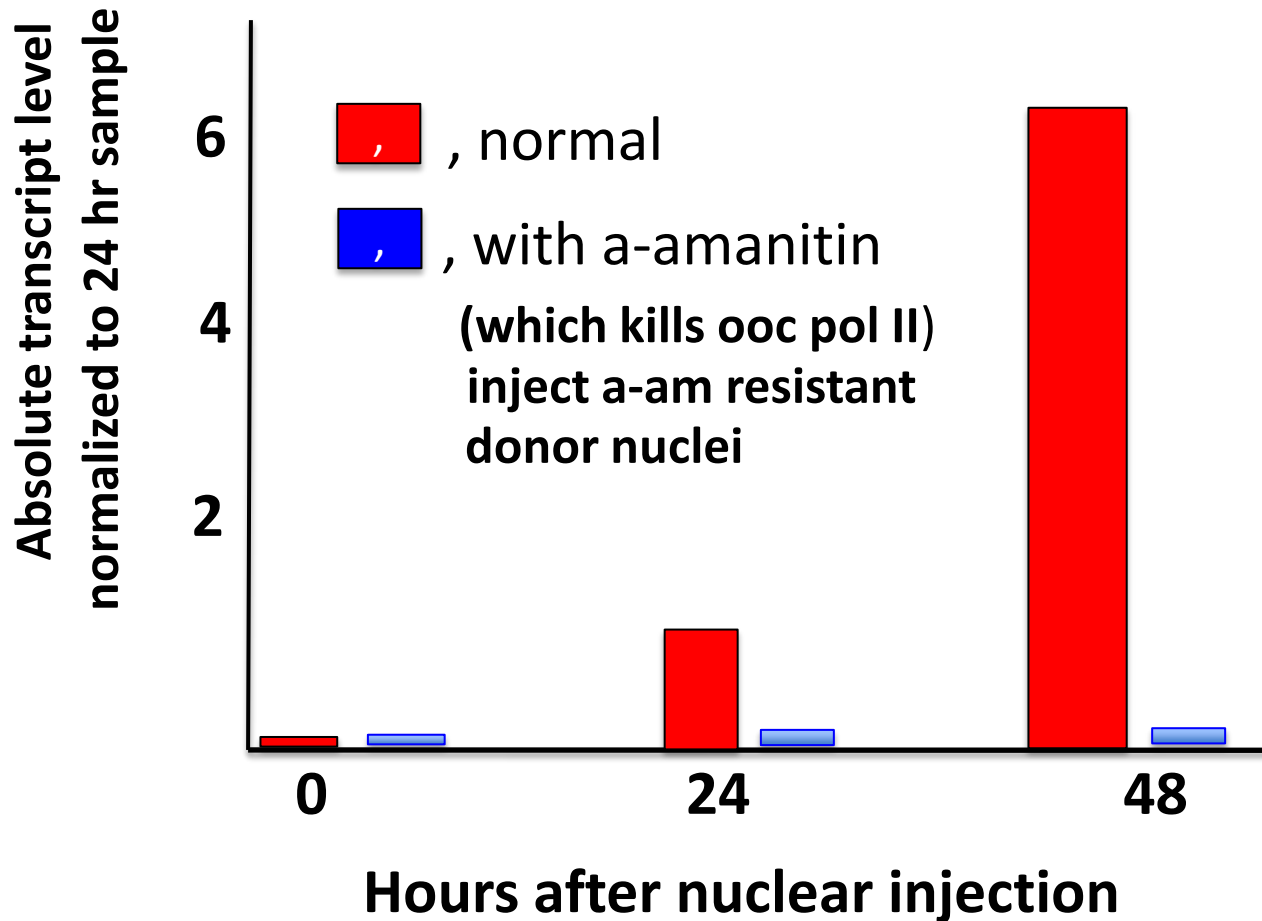


Gain

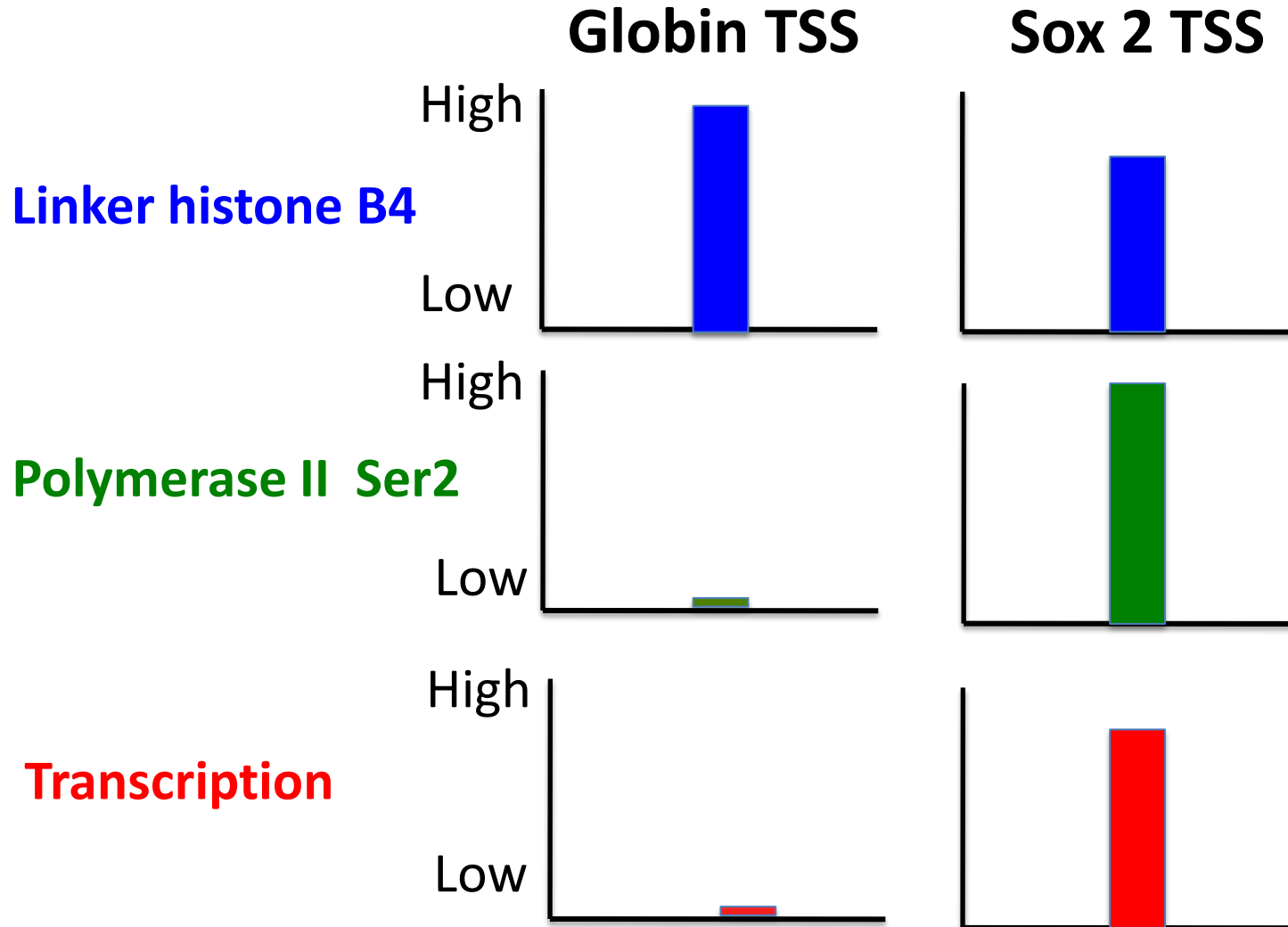
Time sequence of polymerase II components binding to genes



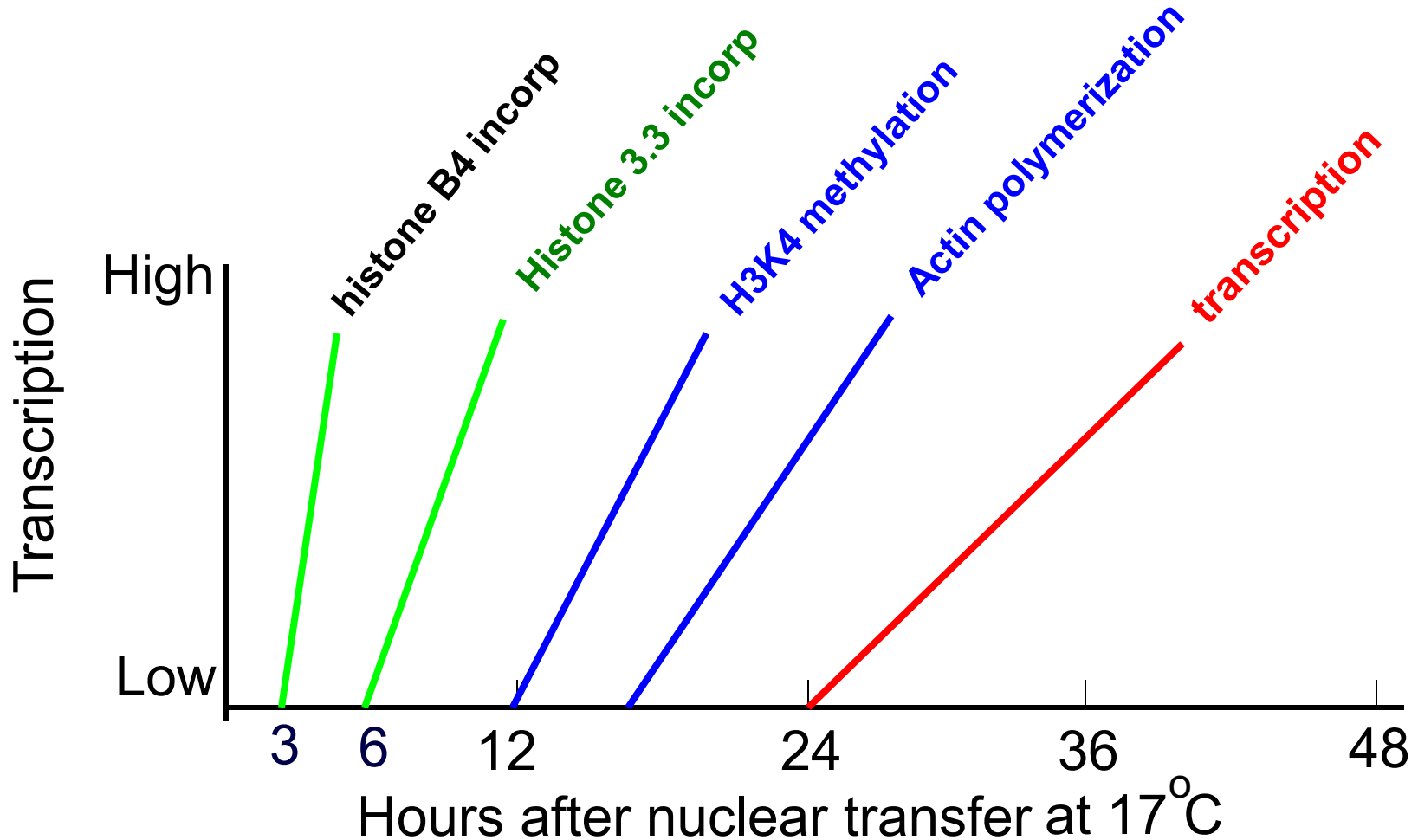
Transcriptional reprogramming depends on polymerase II of oocyte origin



Reprogramming is selective at the level of polymerase II



Time course of transcriptional activation of somatic cell nuclei by oocytes

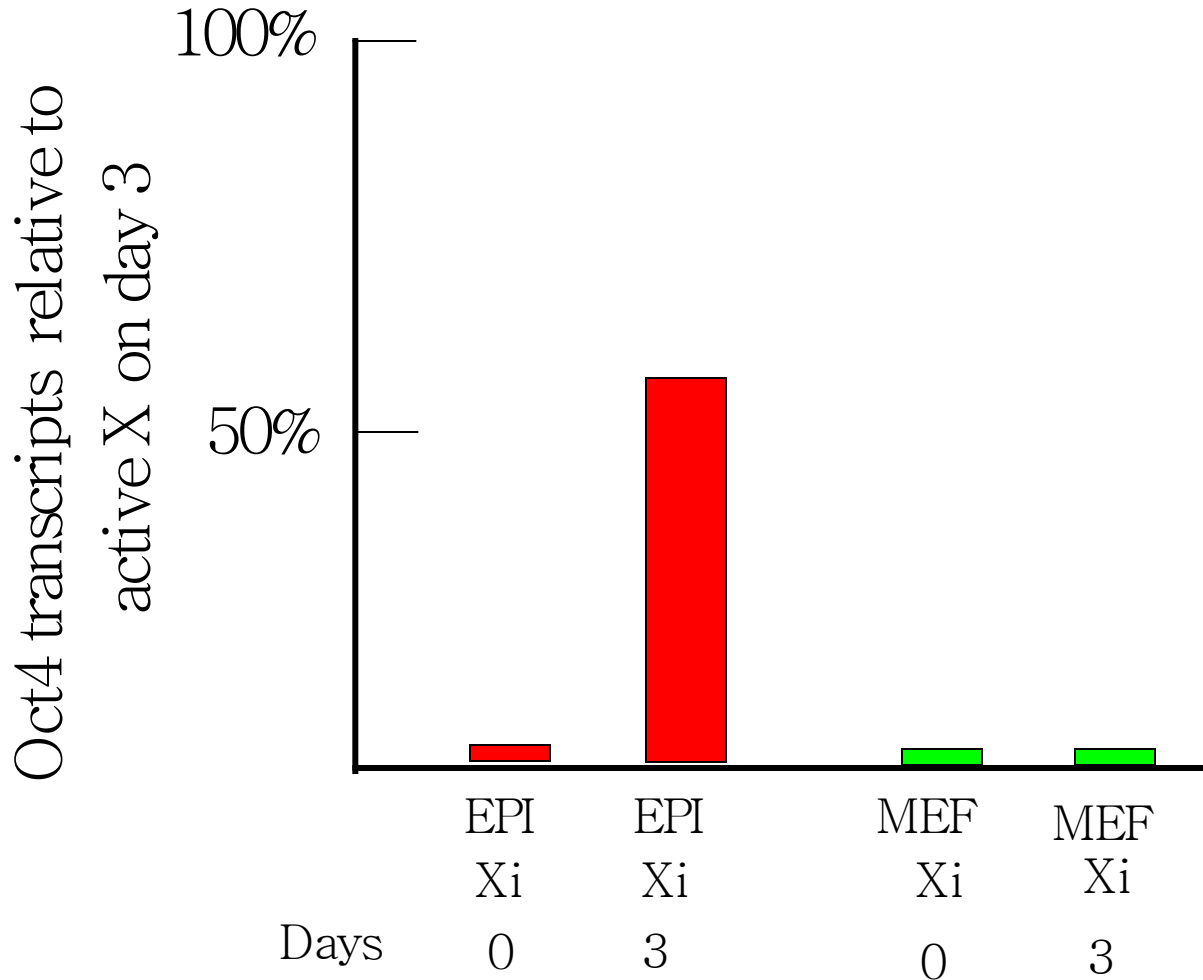


Resistance to reprogramming:

defence by the nucleus

Repressed Xi in female mammals

Epiblast-Xi, but not MEF-Xi, genes are strongly reactivated in injected oocytes



MacroH2A is knocked down by inhibitory RNA,
and induces Oct4 and Sox2 in MEF-Xi cells.

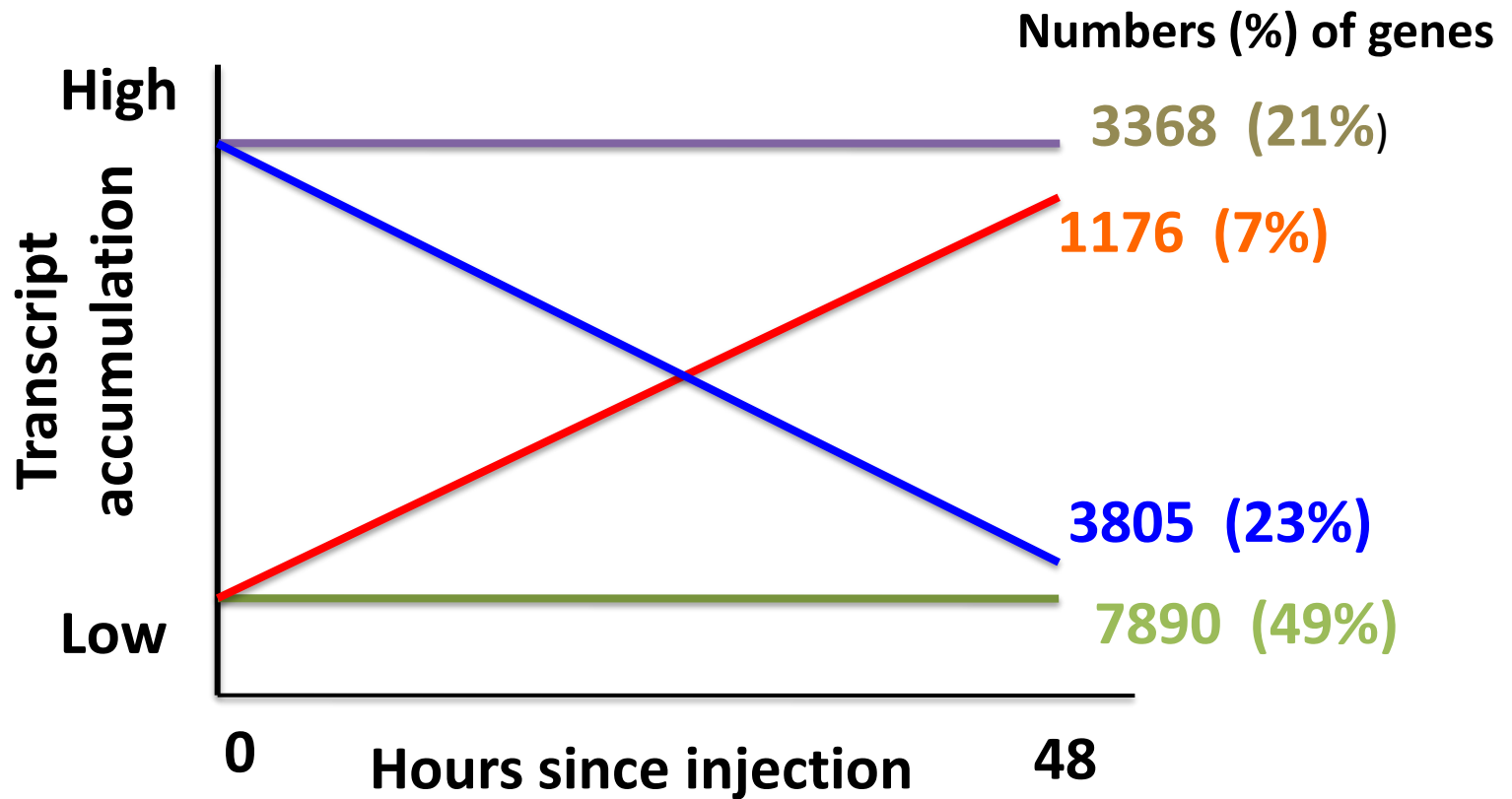
MacroH2A helps to explain
resistance to reprogramming



Conclusion

macroH2A marks embryonic
differentiation and acts as an
epigenetic resistance to nuclear
reprogramming

Selective gene transcription 48 hours after nuclear transfer to *Xenopus* oocytes



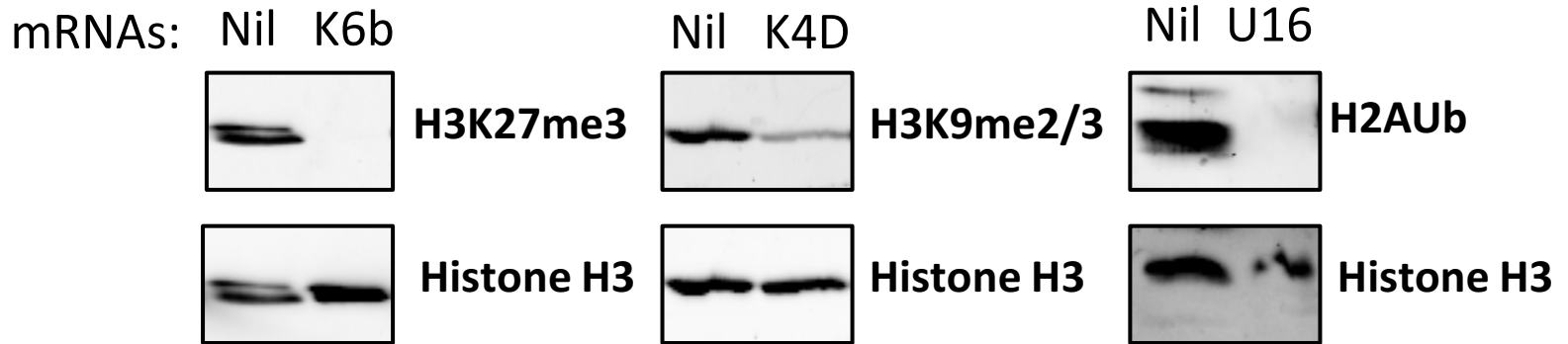
Resistance is gene and cell-type specific

Chromatin modification

Histone modifications in nuclei can be changed after transfer to oocytes

Inject mRNAs on day 1. Transplant nuclei on day 2.

Reisolate transplanted nuclei on day 3 for Western analysis

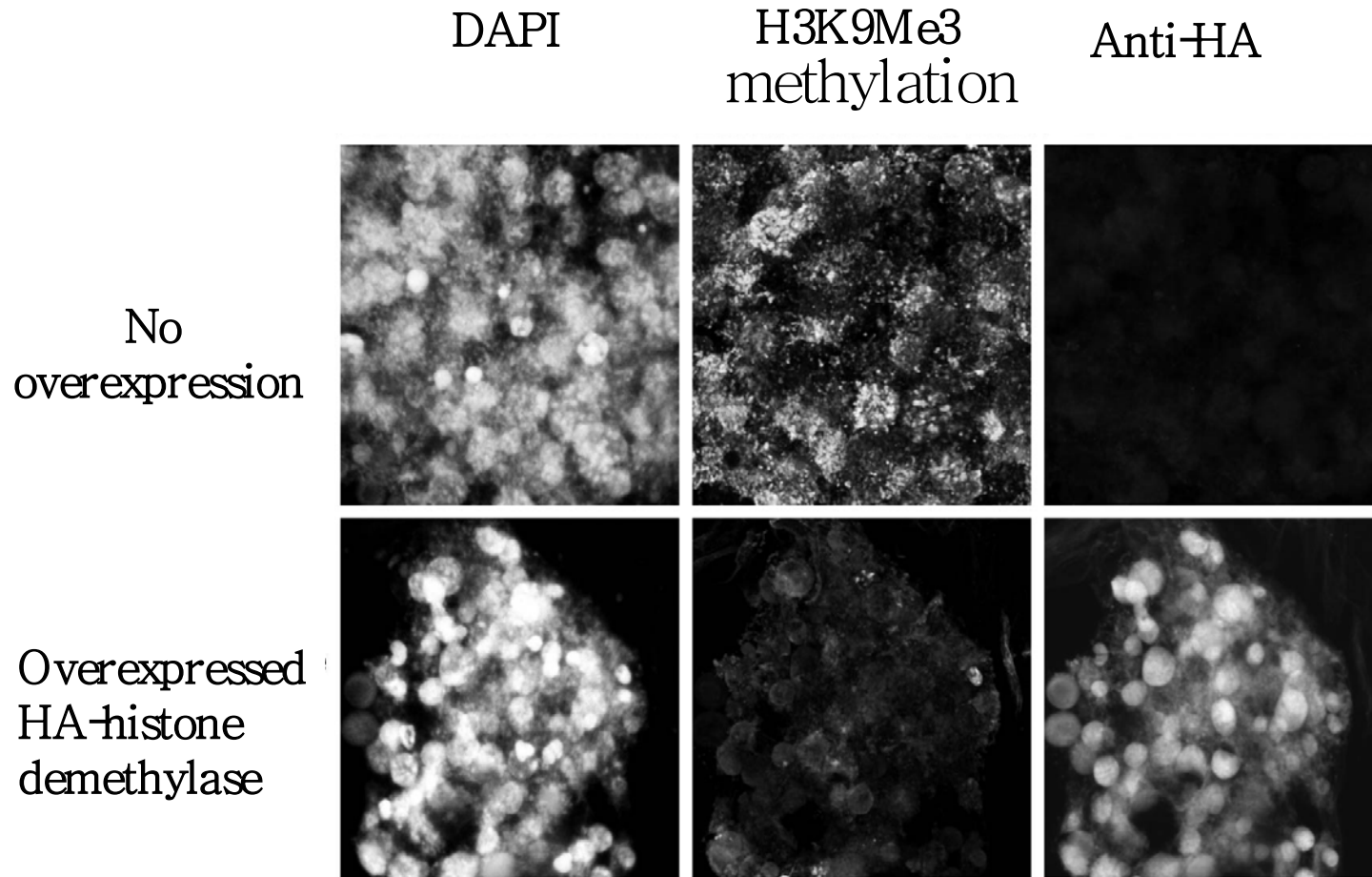


K6b, H3K27me3, H3K27 demethylase.

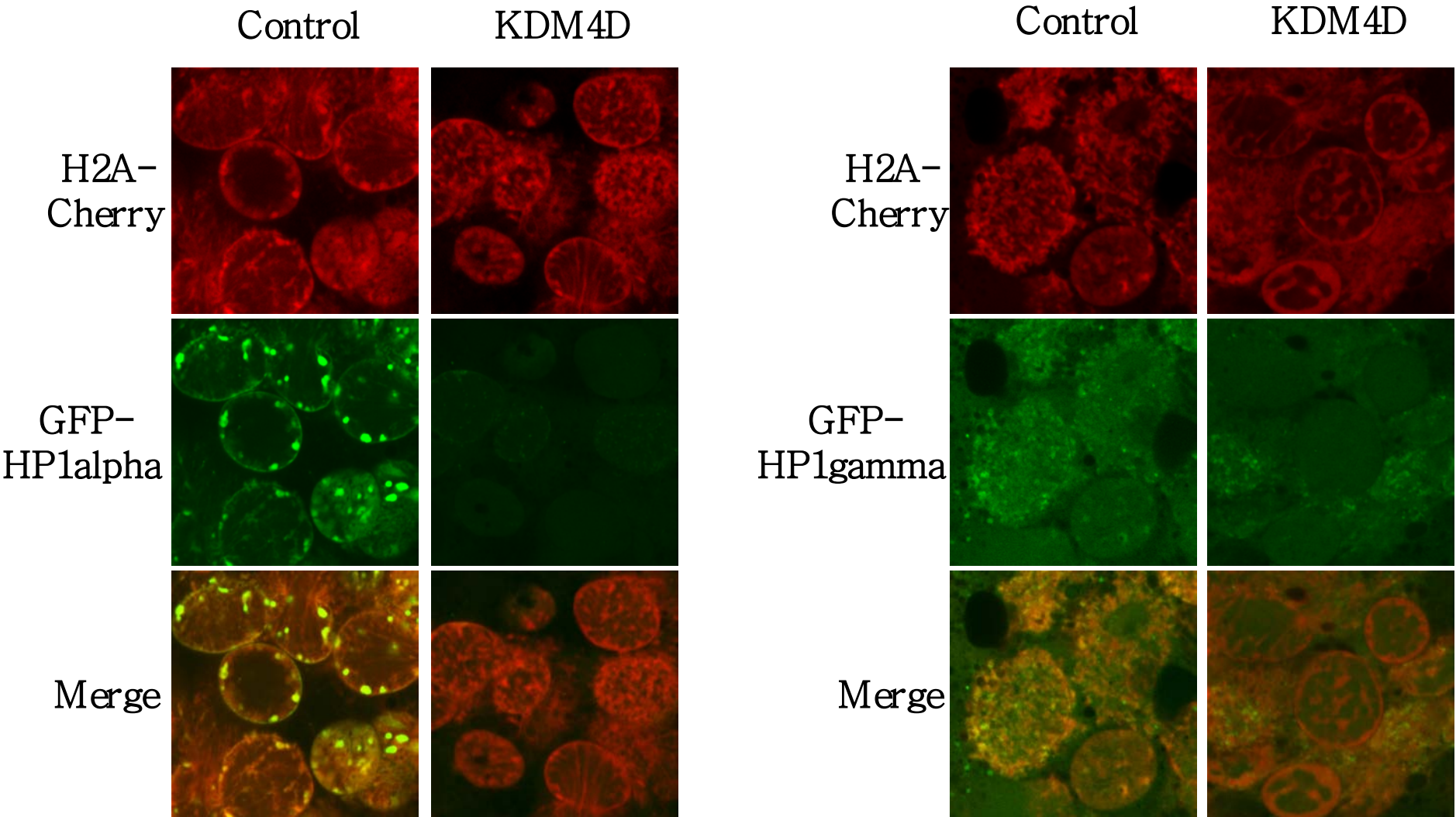
K4D, H3K9 demethylase.

U116, H2A deubiquitinase.

Histone modifiers overexpressed in the oocyte efficiently modify transplanted nuclei chromatin

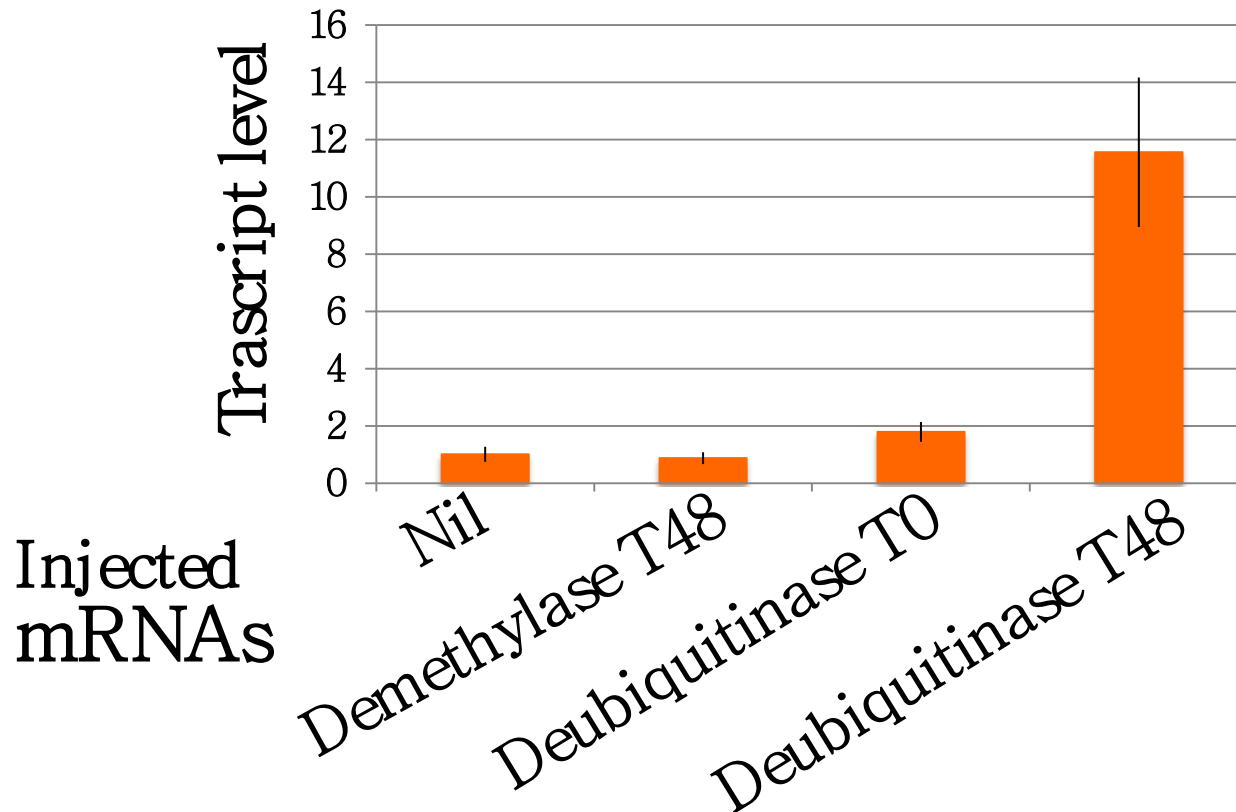


Loss of HP1 alpha binding to transplanted chromatin after lysine demethylase overexpression



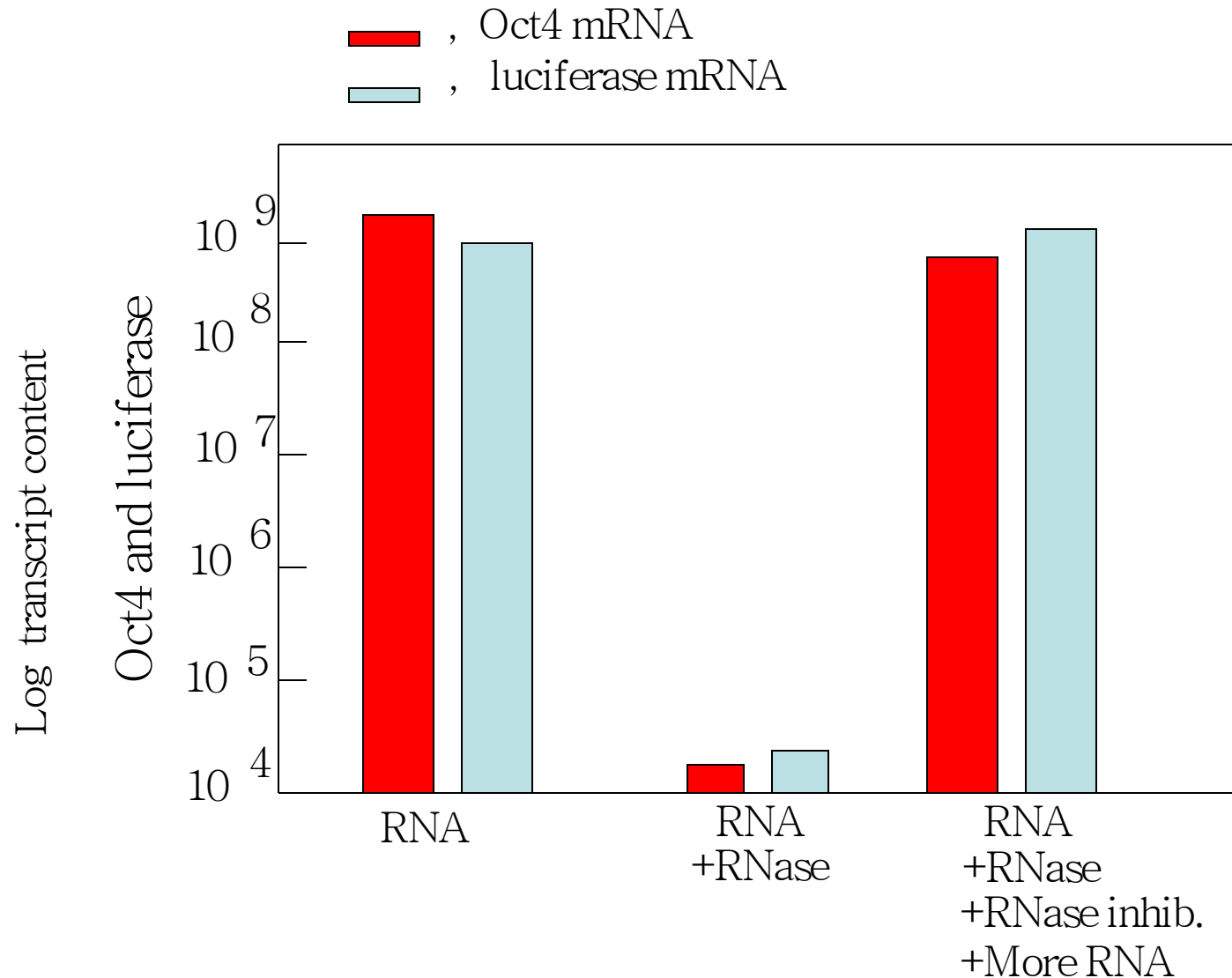
Overexpression of histone 2A deubiquitinase removes resistance

Prok 2



Chromatin depletion

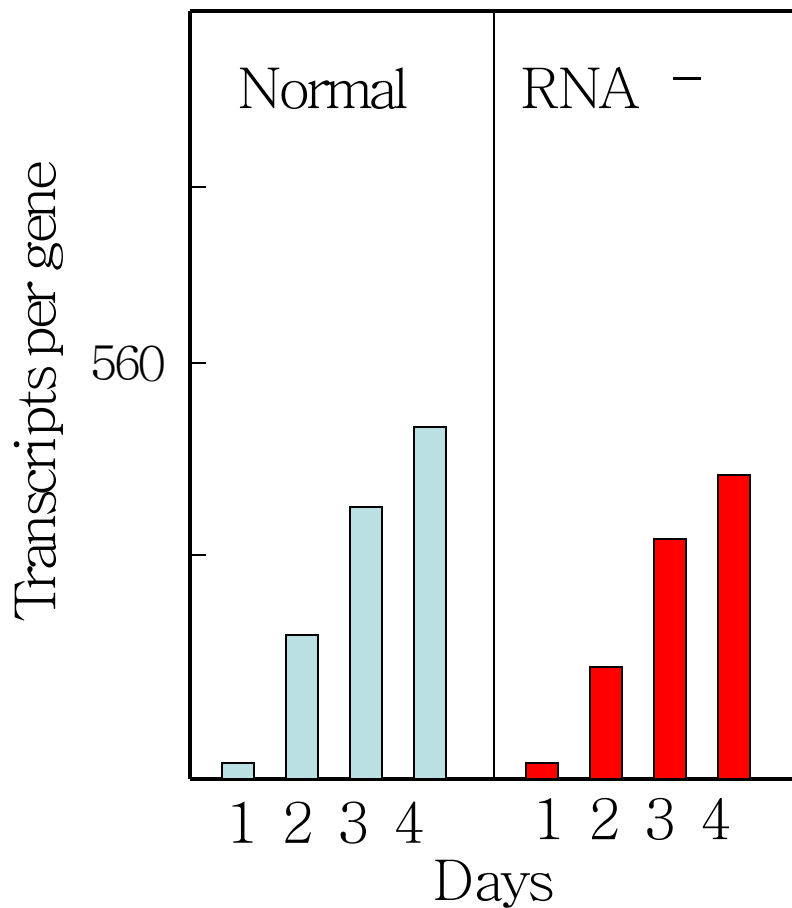
RNA is removed, replaced, and quantitated by RT-PCR



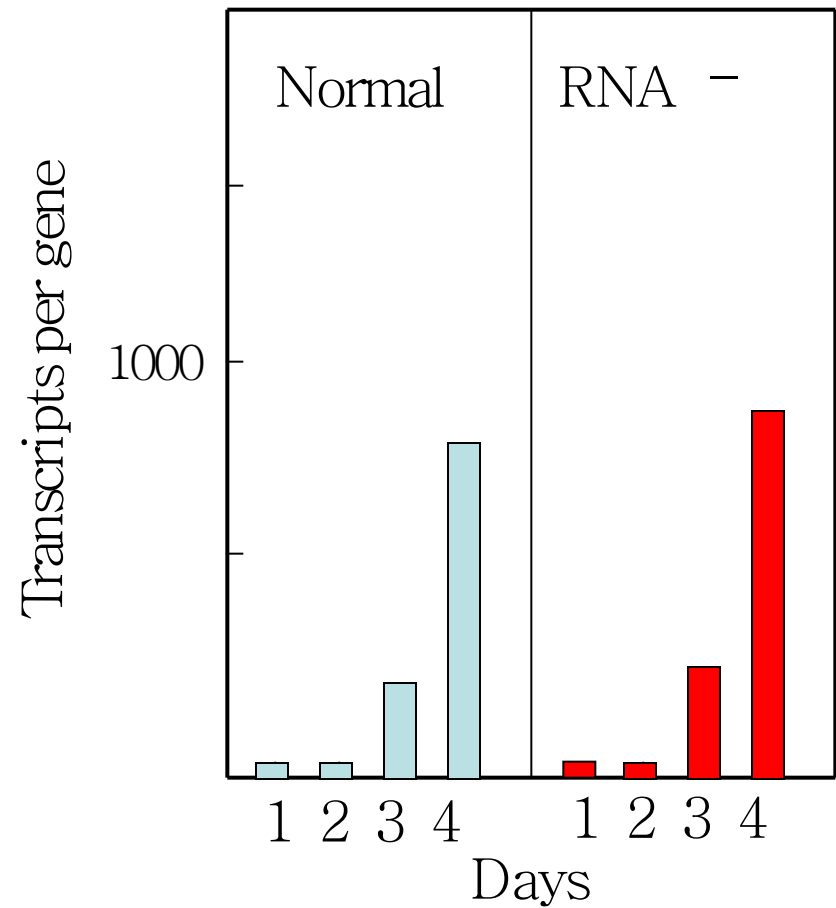
Resistance of nuclei transplanted to oocytes

RNA depletion from donor nuclei does not affect rate or extent of reprogramming

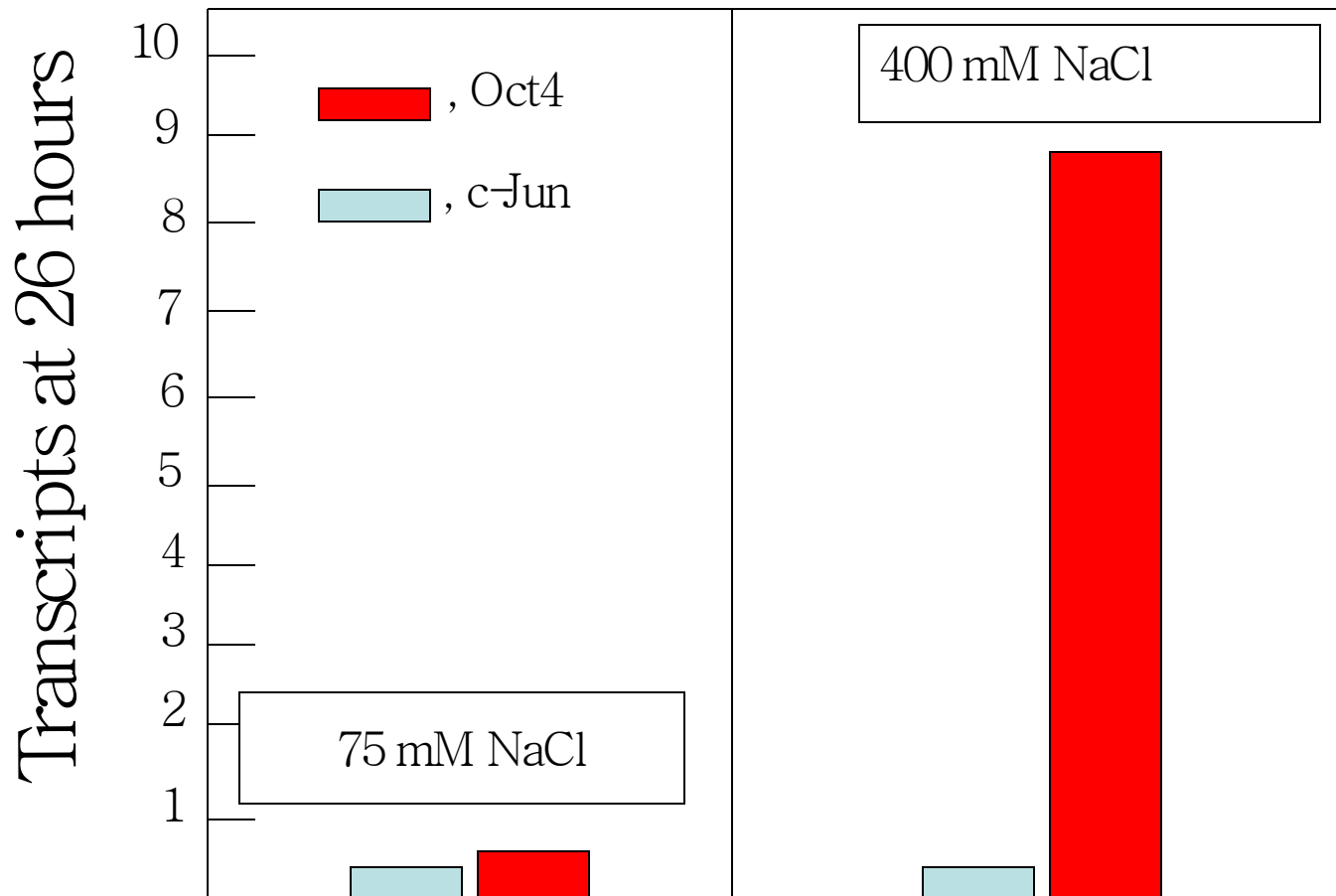
Oct4. Pluripotency.
Thymus nuclei



Sox2. Pluripotency.
Thymus nuclei

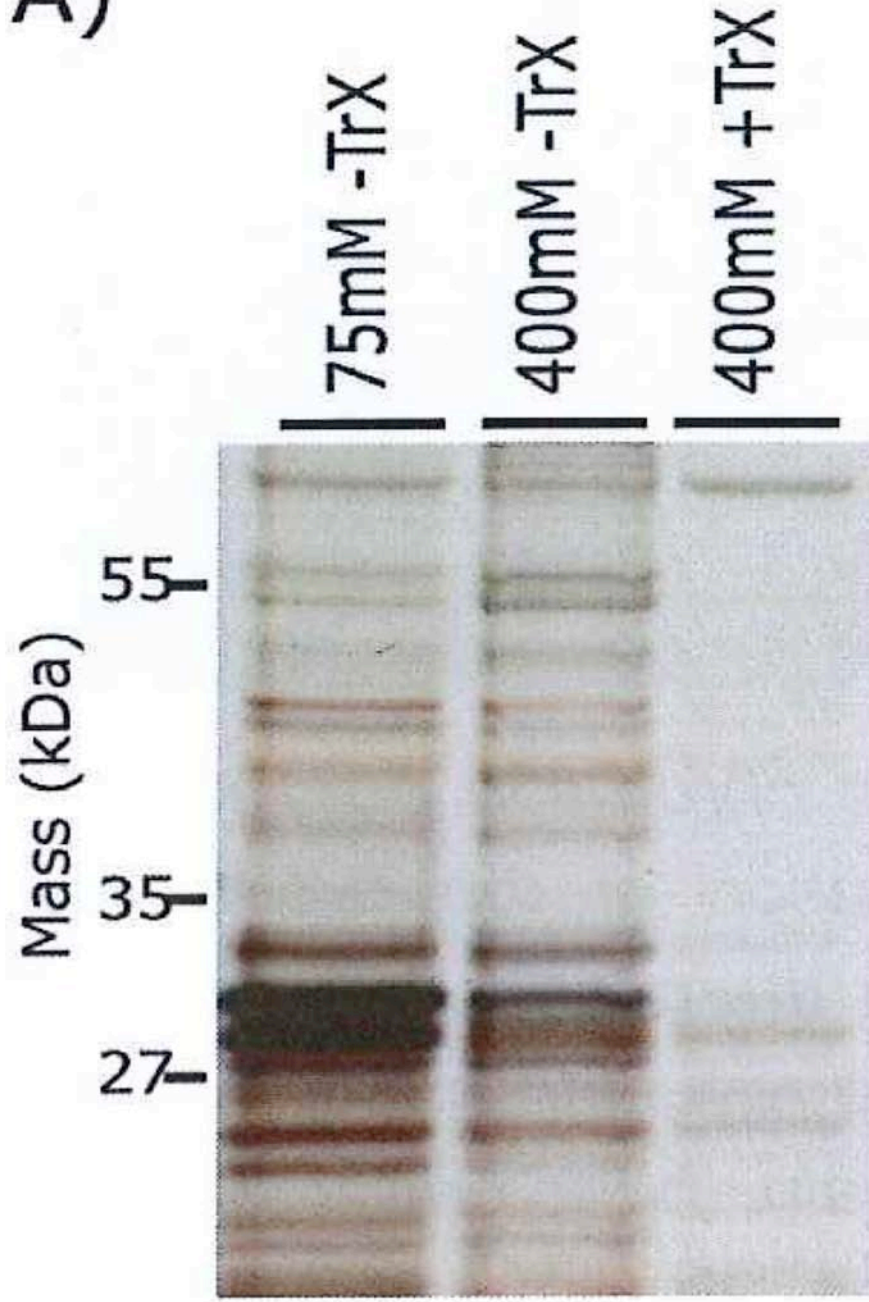


Protein depletion in somatic nuclei removes memory and enhances transcription

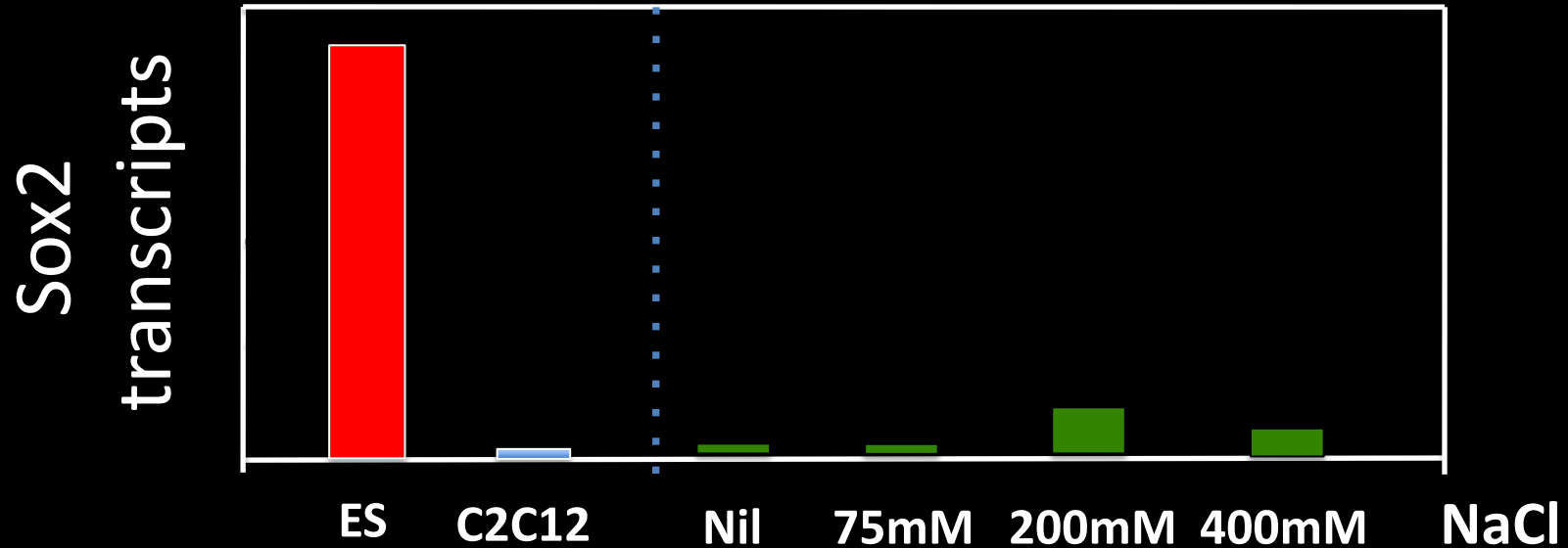


Protein
removal
from
nuclei
by salt
and Triton

A)



Resistance to reprogramming is maintained at high salt concentrations



Levels of chromosomal proteins after salt treatment	HMG	10	10	3	1
	EED	10	10	8	0
	BMi1	10	10	5	0
	HP!	10	10	9	0
	Brg1	10	10	6	1
	Pol II	10	10	9	0
	H2A	10	10	10	10

The battle for supremacy

The egg

Designed to transform
sperm to an embryo
active nucleus

Tries to do the same
for somatic nuclei

The nucleus

Designed to maintain
the same pattern of
gene expression

Tries to resist
any change

Prospect_s

To defeat resistance and win
efficient cell replacement

Acknowledgements

Gurdon lab



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i

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James Byrne

Stina Simonsson

Carolina Astrand

Present colleagues

Jerome Jullien

Kei Miyamoto

Rick Halley-Stott

Vincent Pasque

Marta Teperek

Eva Hoermanseder

Stan Wang

Celia Delahay

MEDICAL RESEARCH COUNCIL

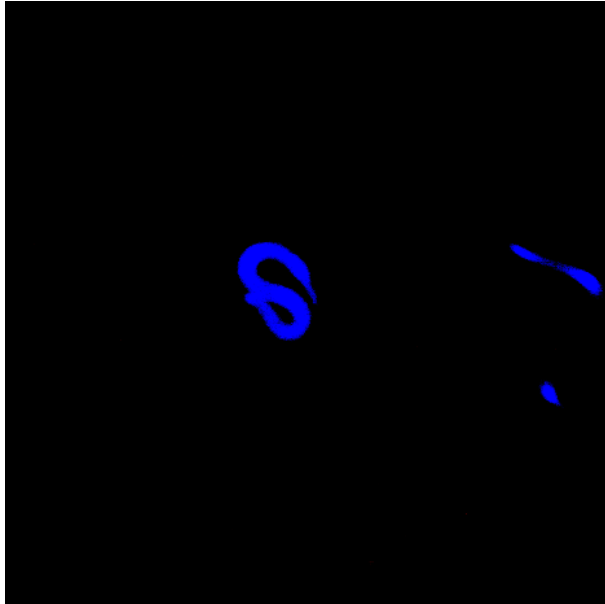
WELLCOME TRUST

CANCER RESEARCH CAMPAIGN

END

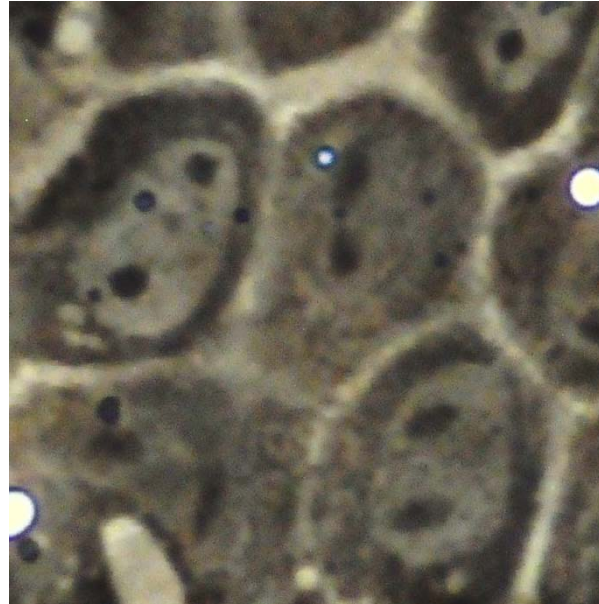
A sperm nucleus is specially designed to yield normal development

Sperm cell



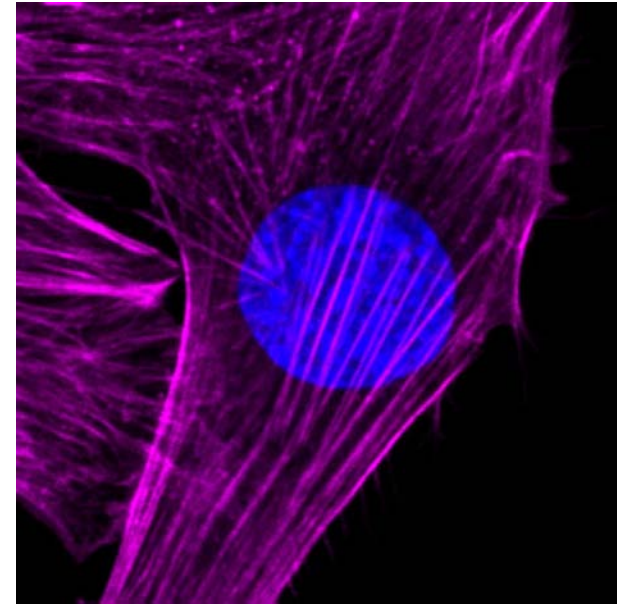
99%

Embryonic cell



35%

Specialised cell



1%

% of normal development after nuclear transfer (to a feeding tadpole)

*Images from
Dr Kei Miyamoto
Marta Teperek*

Conclusions

1. Some cells (endoderm) undergo a very early stable commitment to their lineage pathway.
2. Stable commitment can be reversed by nuclear transfer to eggs.
3. Nuclei from differentiated cells show a strong resistance to reprogramming.
4. Resistance is strongly cell-type and gene specific.
5. Resistance depends on histone modifications and on other stable chromosomal components.

Acknowledgements

Nuclear reprogramming

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Kei Miyamoto Polym. Actin)

Vincent Pasque (Xi)

Richard Halley-Stott (Trn)

Kazutaka Murata (Histone mods)

Marta Teperek (Sperm)

Other Laboratories

Welcome
Trust

G. Crabtree (Stanford)

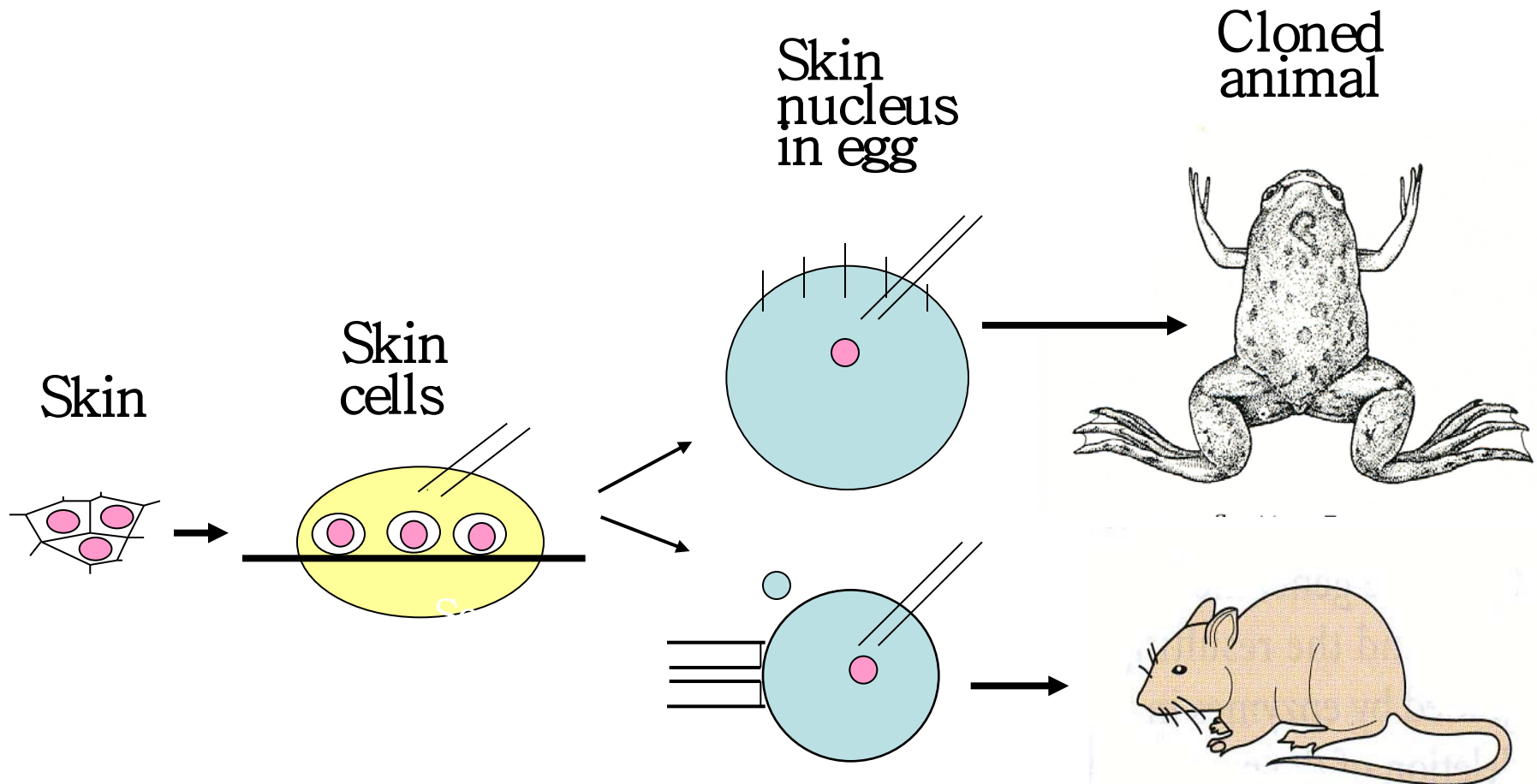
K. Ohsumi (Nagoya)

G. Almouzni (Paris)

K. Shinkai (Kyoto)

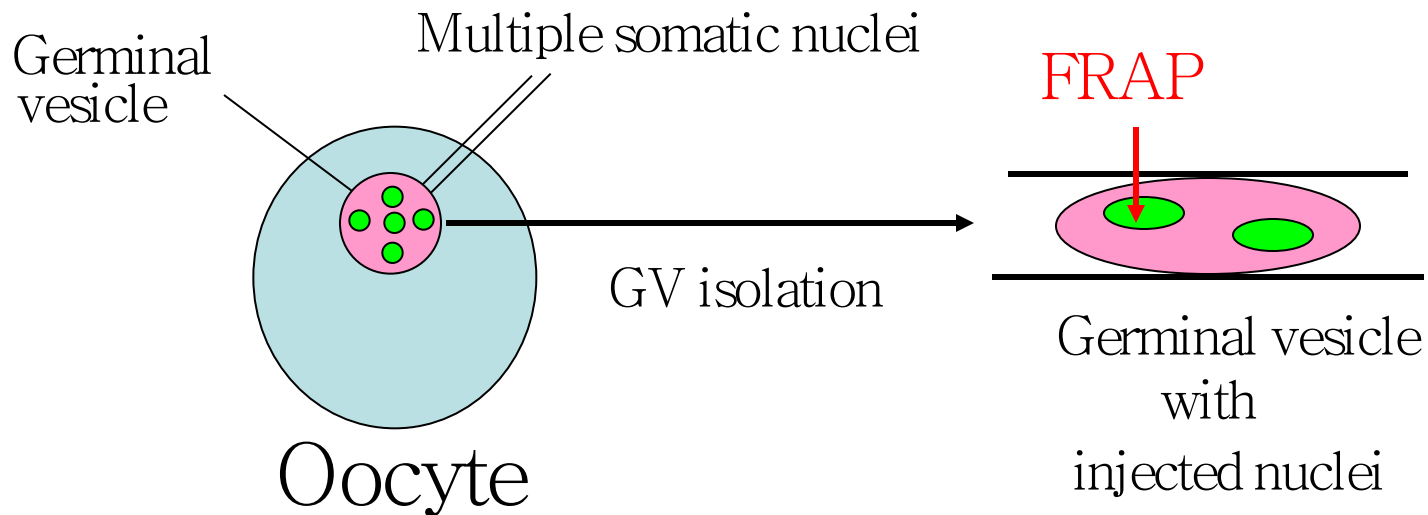
Medical
Research
Council

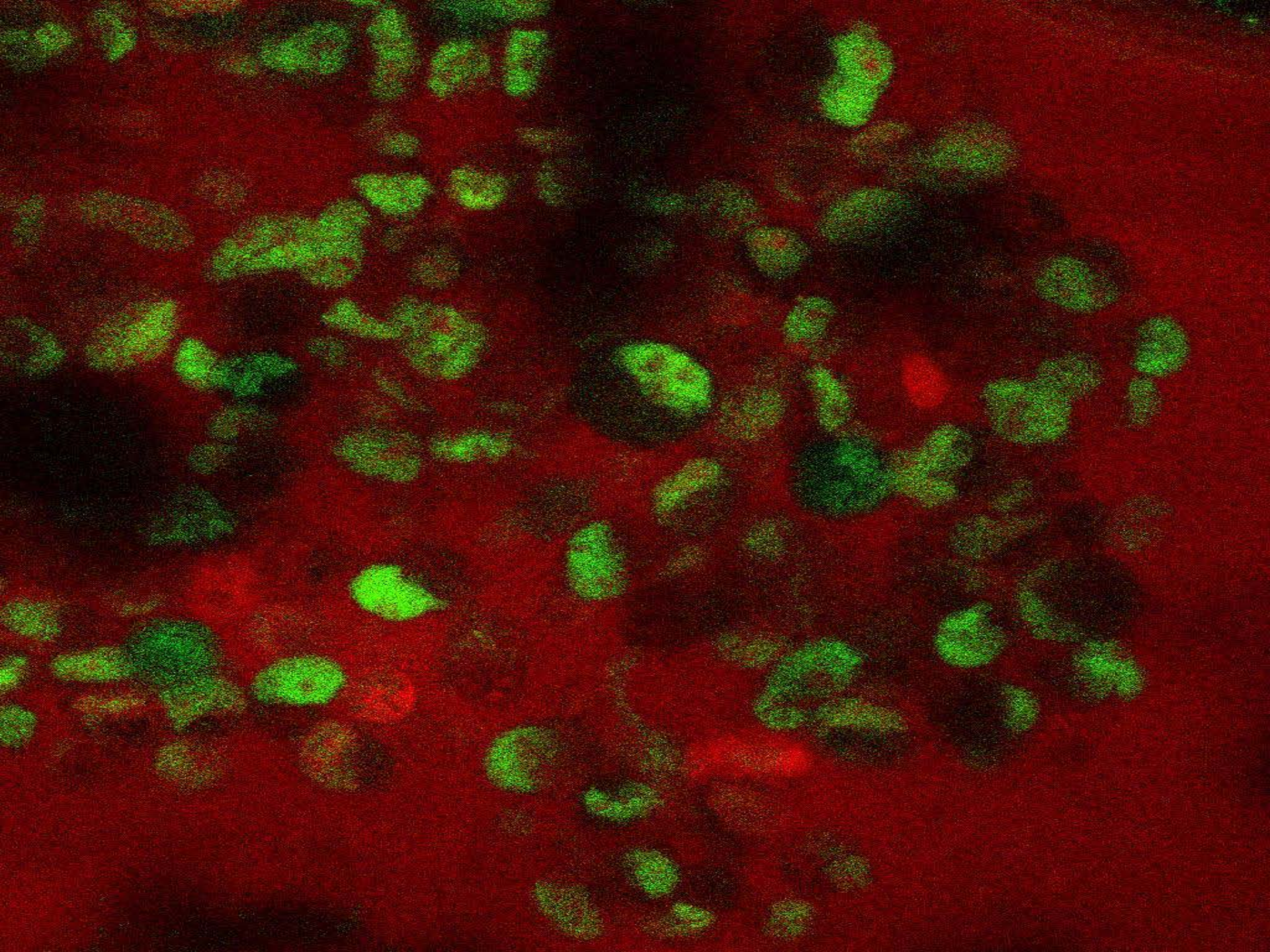
Single nuclear transfer to unfertilized eggs



Fluorescence recovery after photobleaching

To determine the exchange rate of a defined protein
in transplanted nuclei





Increase in polymerase II after nuclear transfer

Time after nuclear transfer

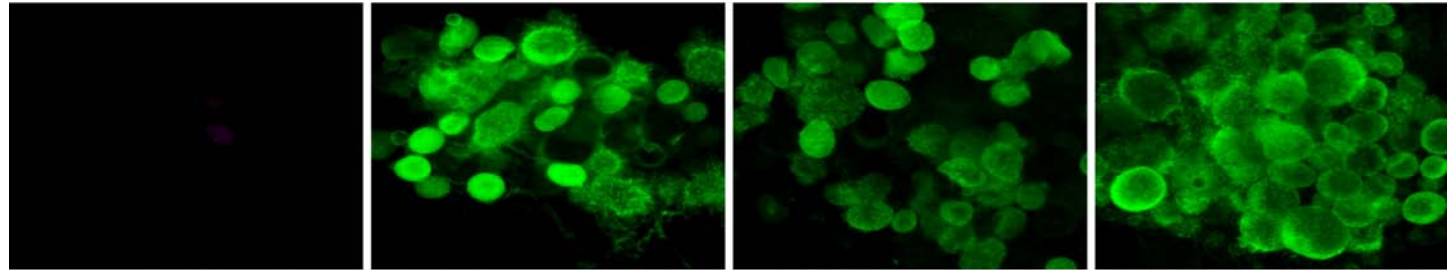
1 hour

6 hours

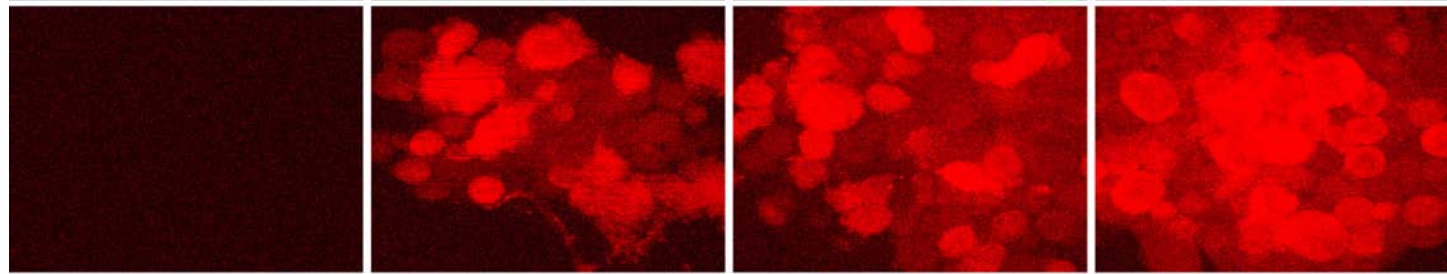
24 hours

48 hours

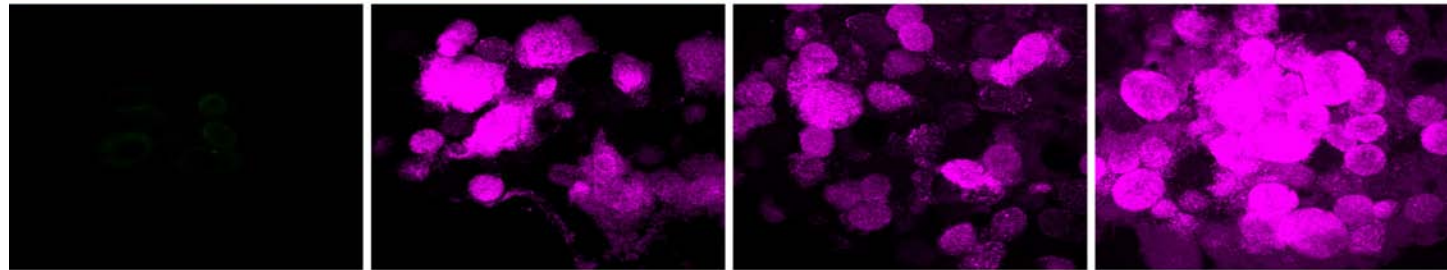
Histone B4



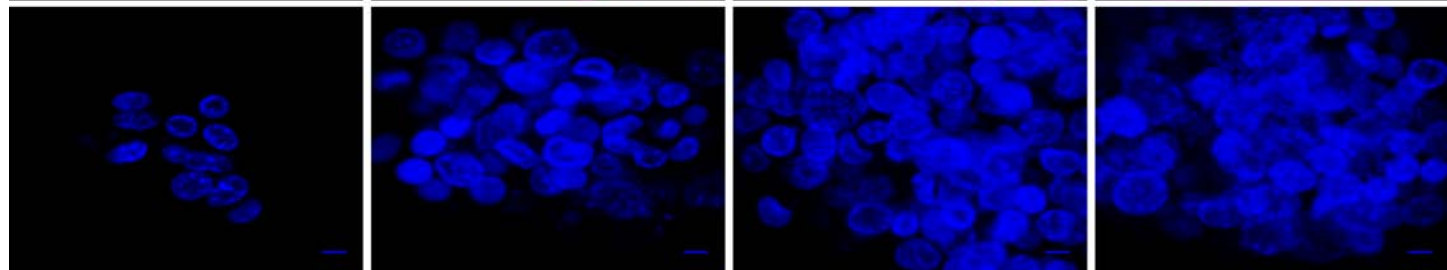
Pol II total



Pol II Ser 2

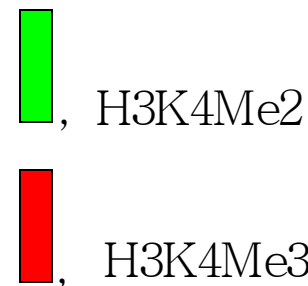
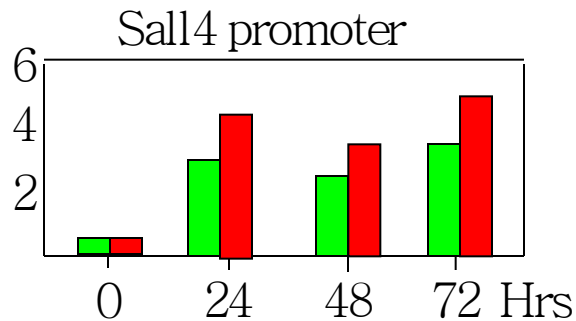
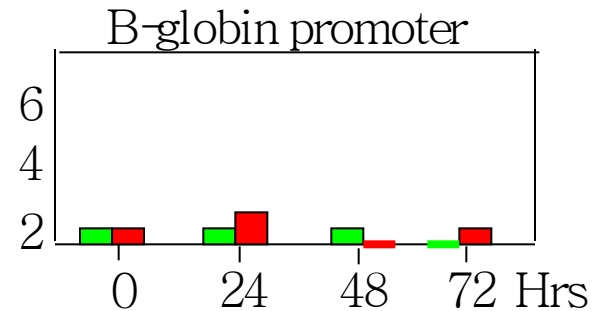
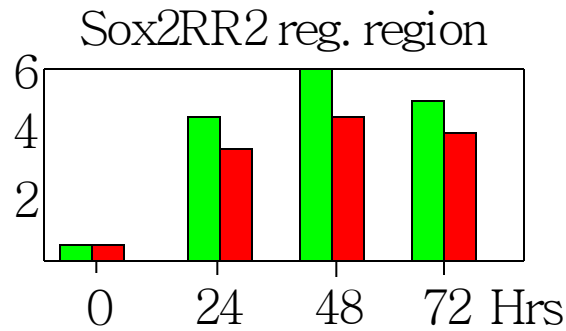
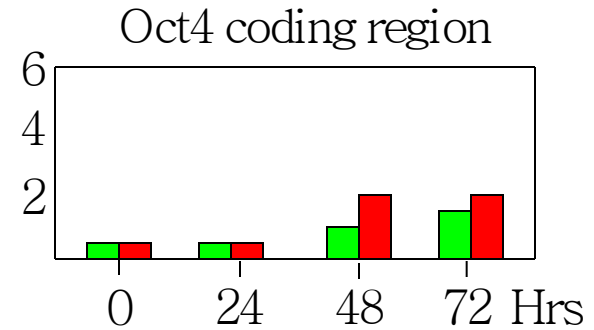
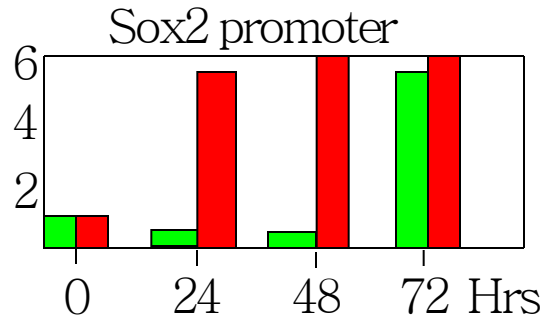


DAPI



Histones in gene control regions are methylated – Chip analysis

Nuclei from retinoic acid treated ES cells

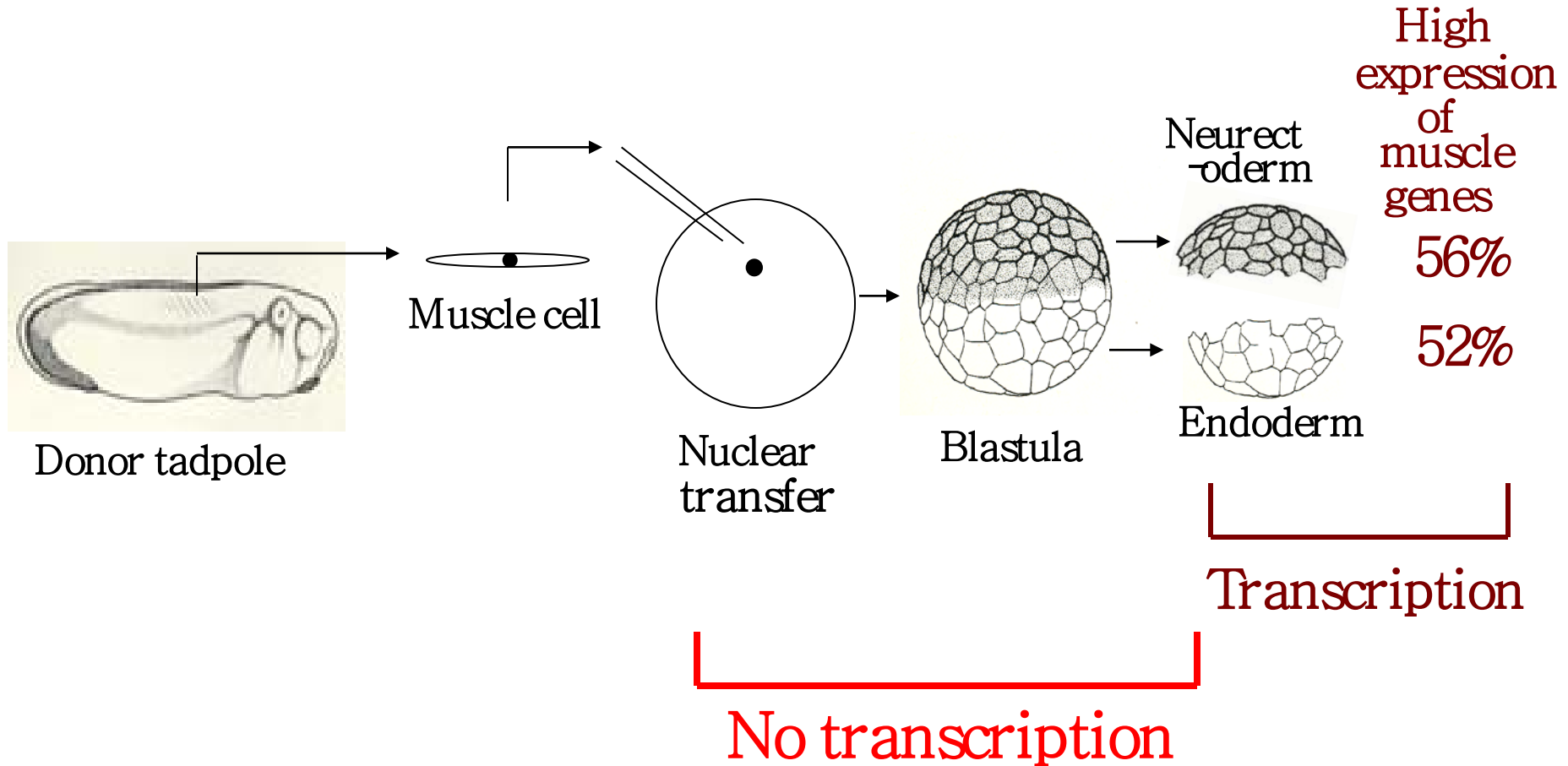


MacroH2A helps to explain resistance to reprogramming

MacroH2A is high on MEF-X:i resists reprogramming.
but absent from EPI-Xi: is reprogrammed.

MacroH2A is knocked down by inhibitory RNA,
and induces Oct4 and Sox2 in MEF-Xi cells

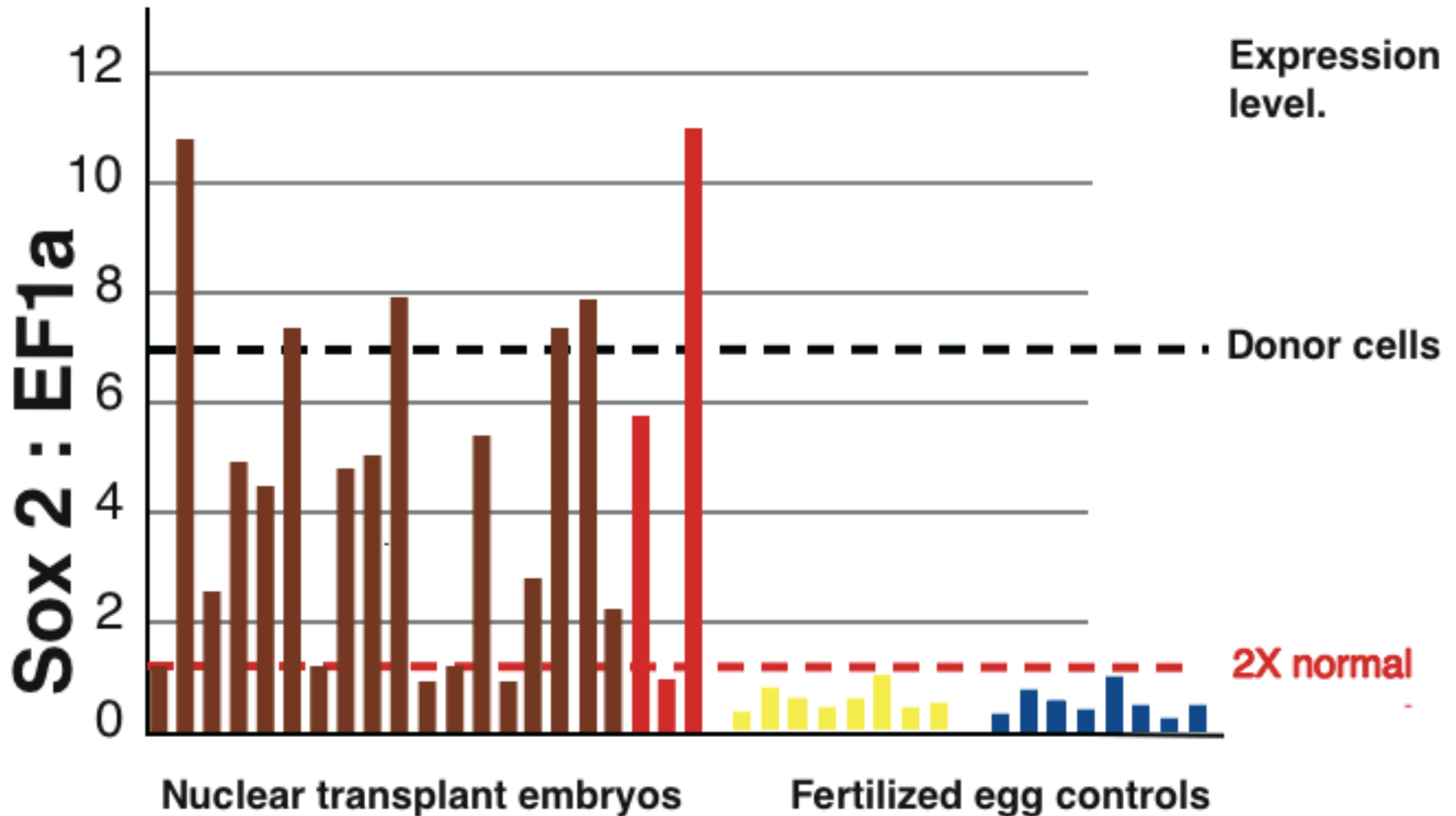
Epigenetic memory



The resistance of MEF Xi nuclei to reprogramming
by oocytes is not explained by
DNA methylation or by
histone H3K27 me

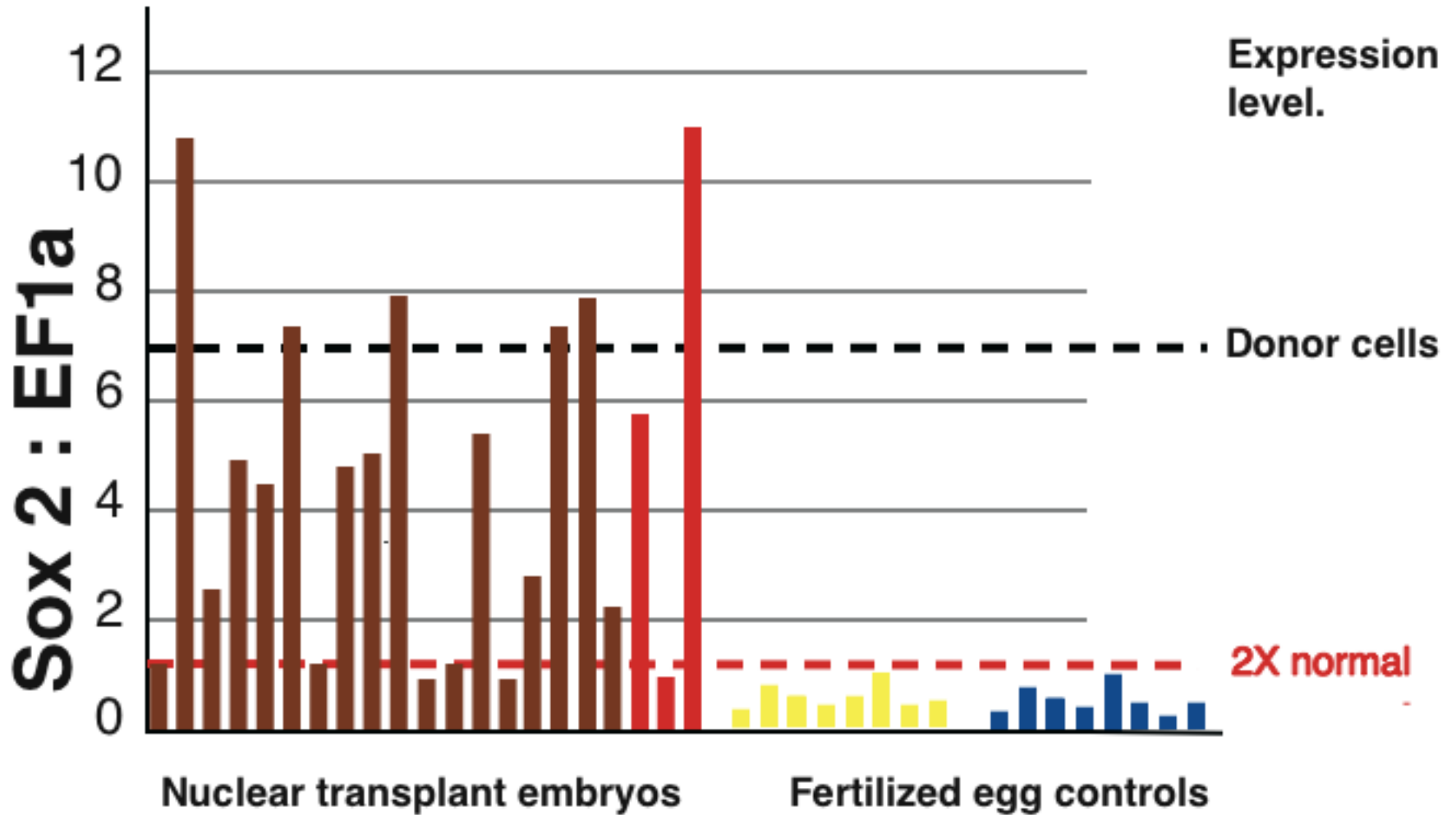
Memory of cell type gene expression persists through nuclear transfer.

Neurectoderm nuclei (neural marker Sox2)
and Sox2 expression in endoderm nuclear transfer cells.



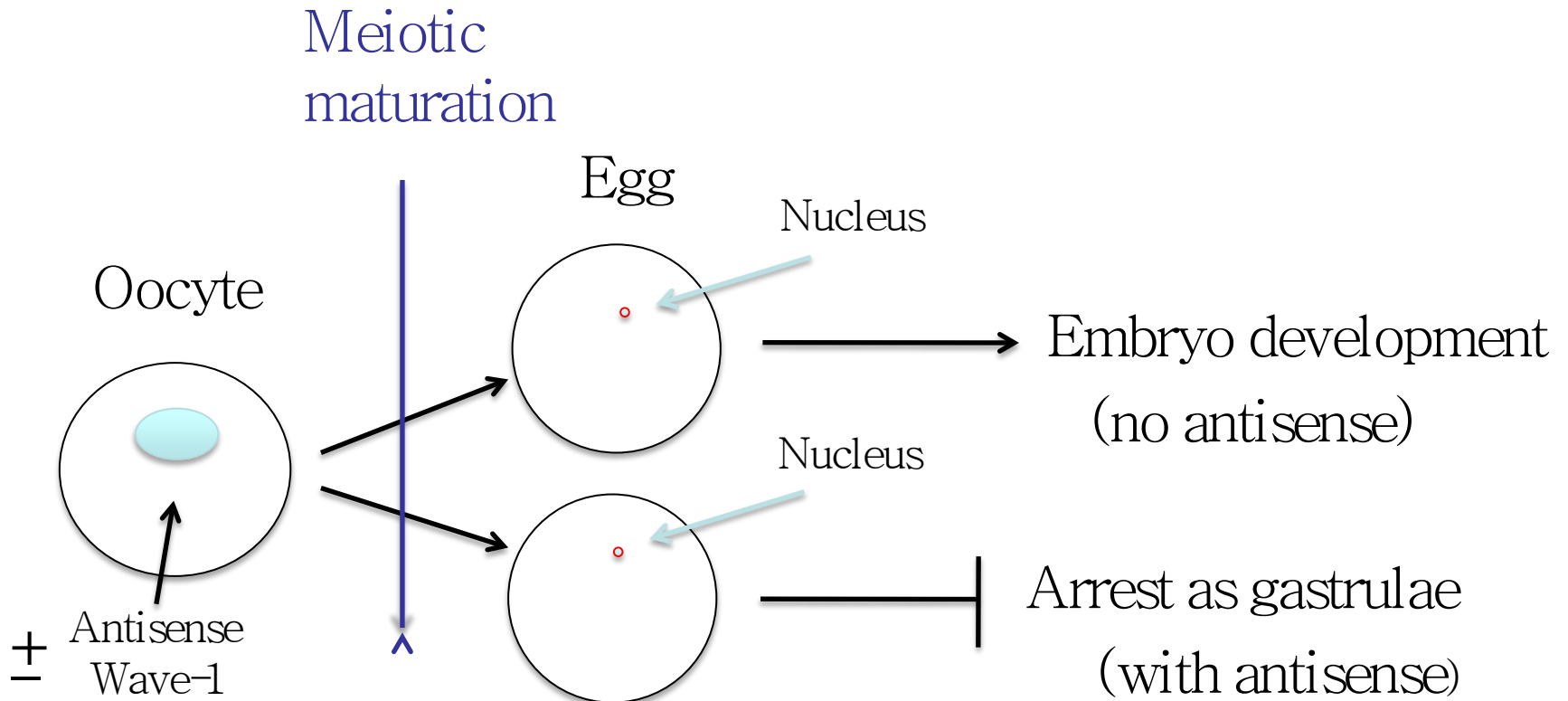
Memory of cell type gene expression persists through nuclear transfer.

Neurectoderm nuclei (neural marker Sox2)
and Sox2 expression in endoderm nuclear transfer cells.



WAVE-1 is required for zygotic genome activation and embryonic development

(Wiskott-Aldrich syndrome)

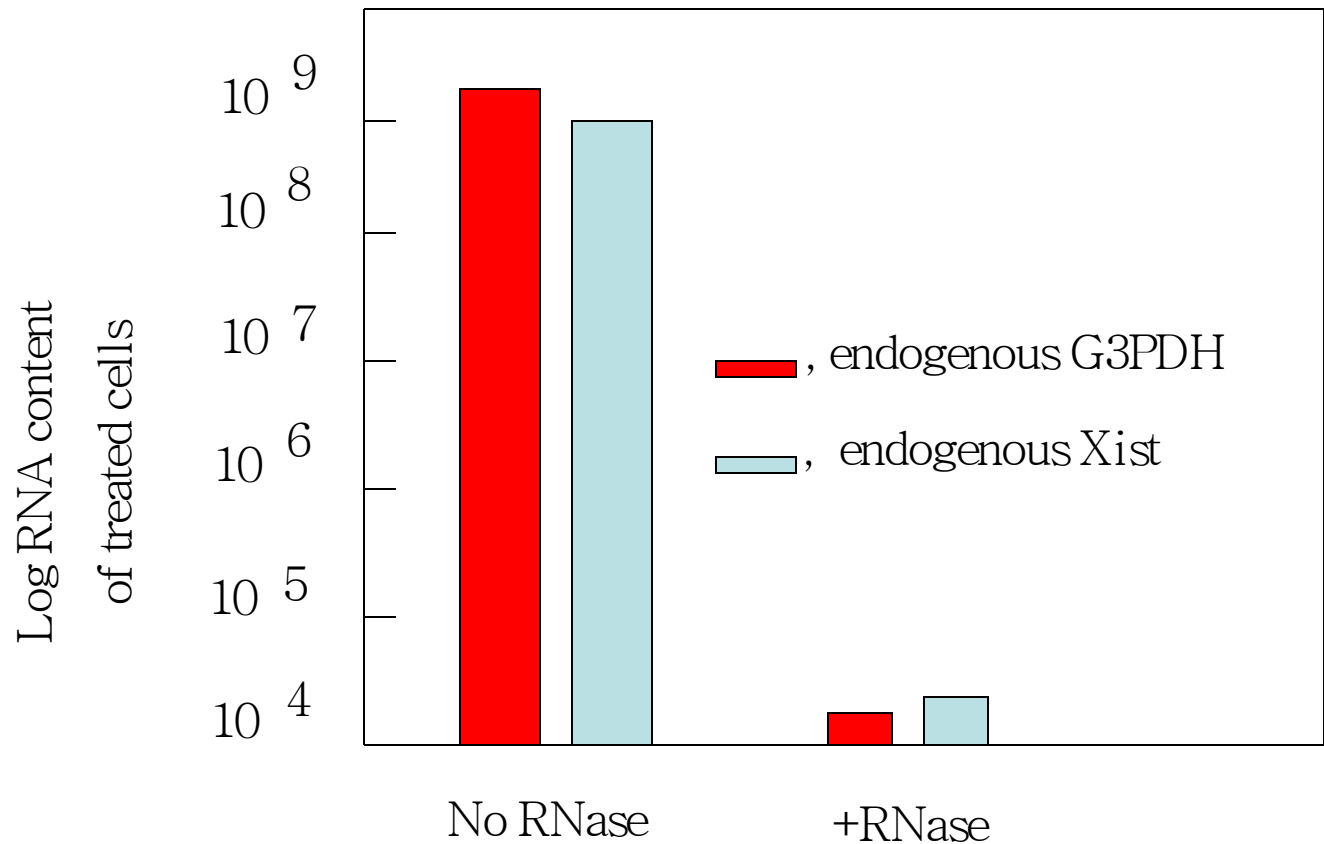


Histone modifiers overexpression in the oocyte:

- H3K9 demethylase KDM4D efficiently removes H3K9Me_{2/3} from transplanted nuclei and leads to loss of HP1 alpha
- H2A deubiquitinases (USP16&21) reduce ubiquitinated H2A level in transplanted nuclei

RNA can be depleted from donor nuclei by RNase.

Permeabilized cells, containing RNA, are treated with RNase, then assayed for residual RNA.



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Present colleagues

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Vincent Pasque

Marta Teperek

Eva Hoermanseder

Stan Wang

Celia

MEDICAL RESEARCH COUNCIL

WELLCOME TRUST

CANCER RESEARCH CAMPAIGN

Dolly the Sheep

When Dolly was born at the Roslin Institute on 5 July 1996, she was the first mammal to be cloned from an adult cell. In her short life Dolly came to symbolise the future of cloning throughout the world. In 1997 the Roslin Institute agreed to donate Dolly to the National Museums of Scotland when she died, so that she could be preserved for future generations to see. She died on 14 February 2003.

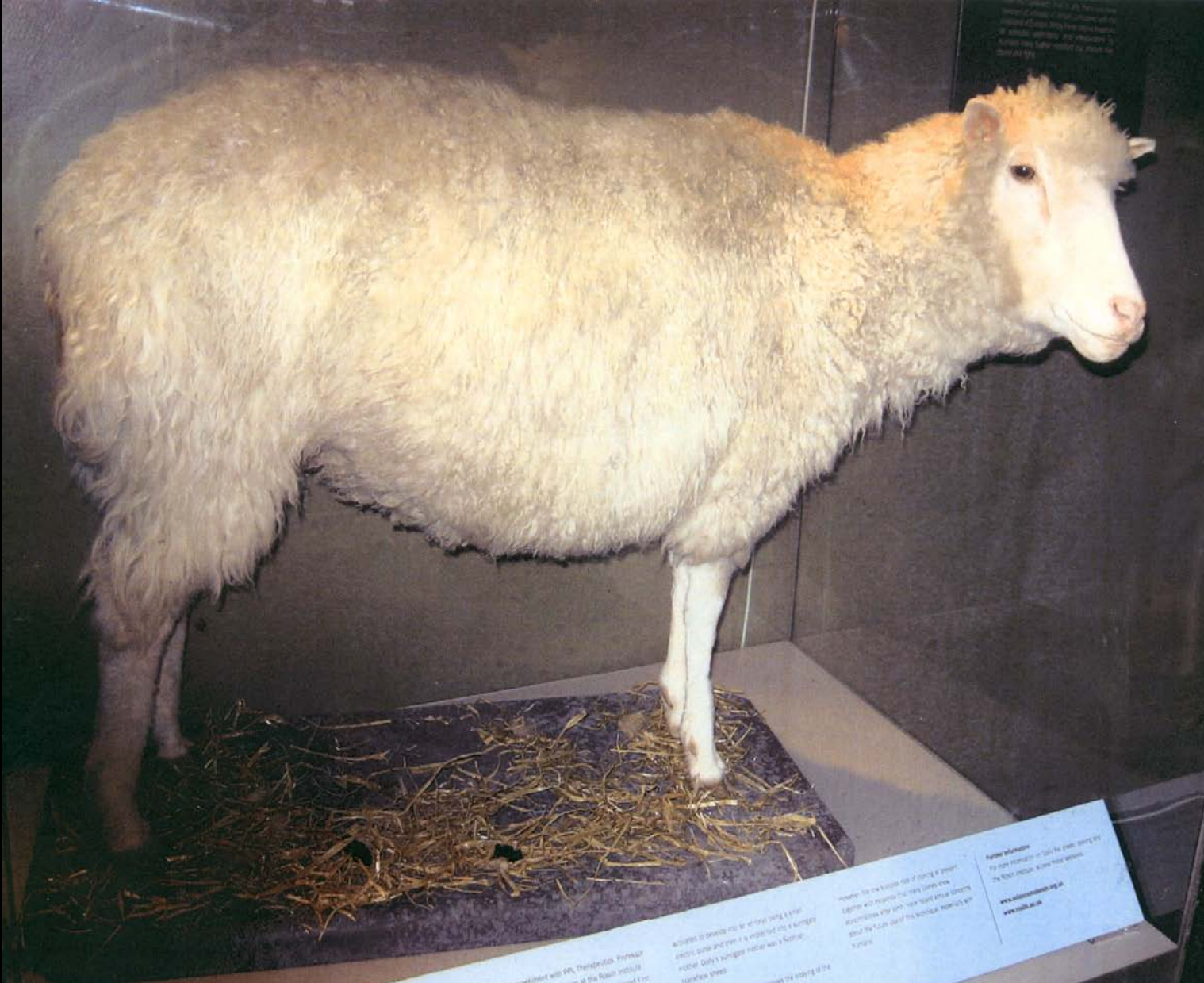
In a joint experiment with IAC, Theoretical Professor Ian Wilmut and his colleagues at the Roslin Institute used a cell from the udder tissue of a six-year-old Finn-Dorset ewe in order to create Dolly.

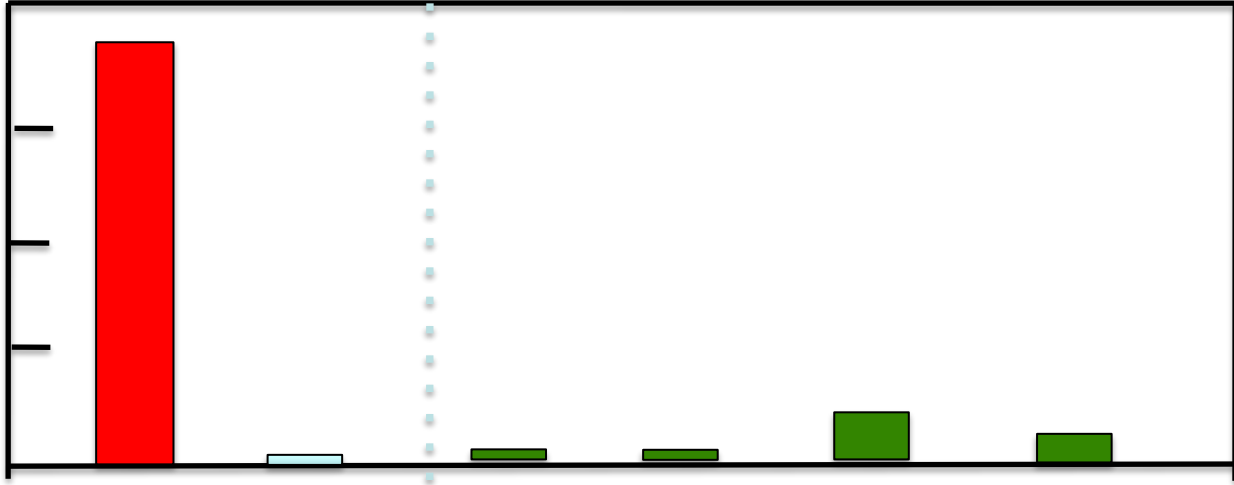
Cloning involves the removal of the nuclear DNA from an egg cell, so that it can be replaced by the nucleus from a donor cell. The reconstructed egg cell is

incubated to develop into an embryo using a special electric pulse and then it is implanted into a surrogate mother. Dolly's surrogate mother was a Scottish Blackface ewe.

Cloning from adult mammal cells is the ending of the very brief 'stem' phase and may not have the potential of early embryonic stages.

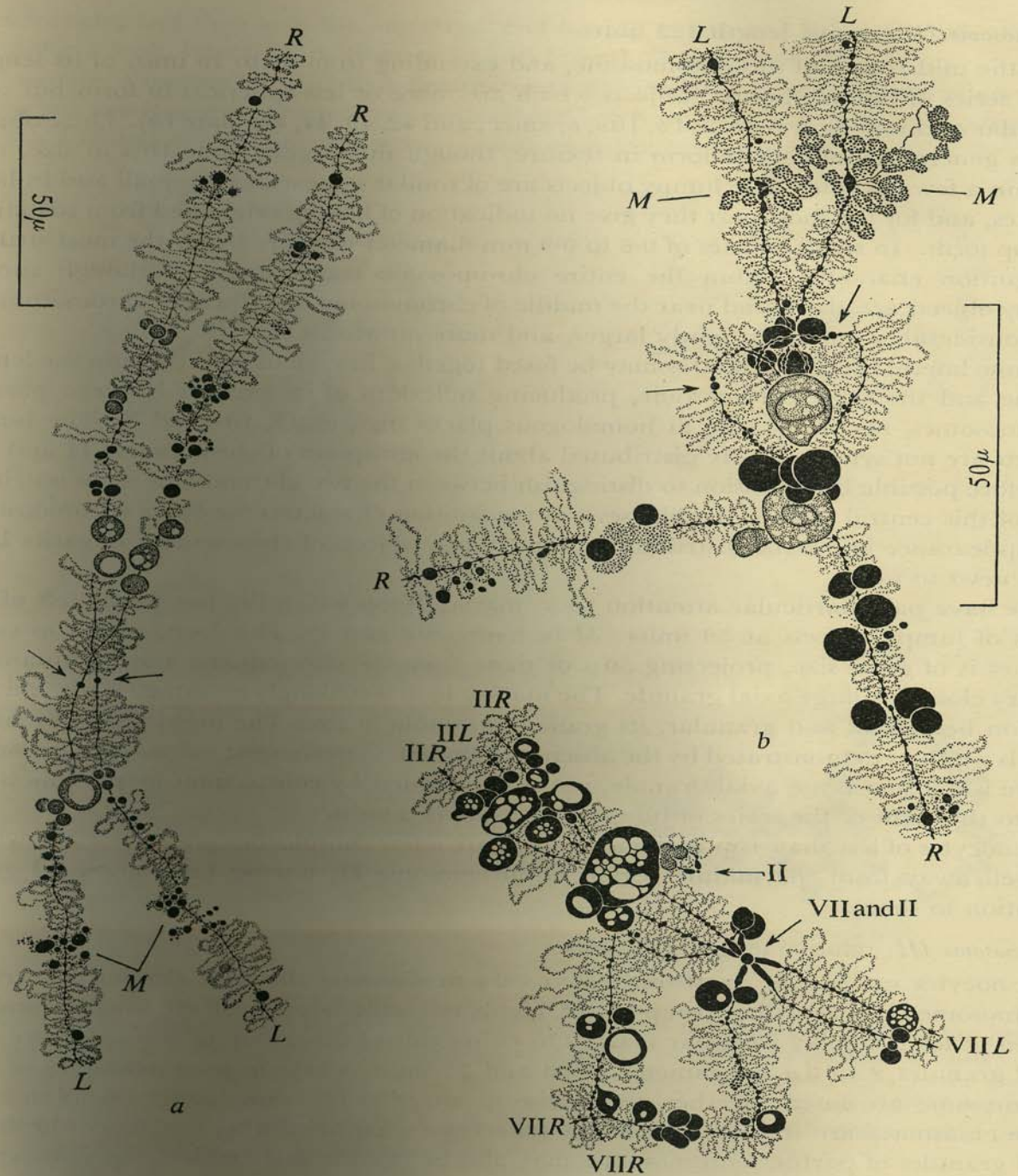
Further information
For more information on Dolly the sheep, visit the Roslin Institute's website:
www.roslin.ac.uk



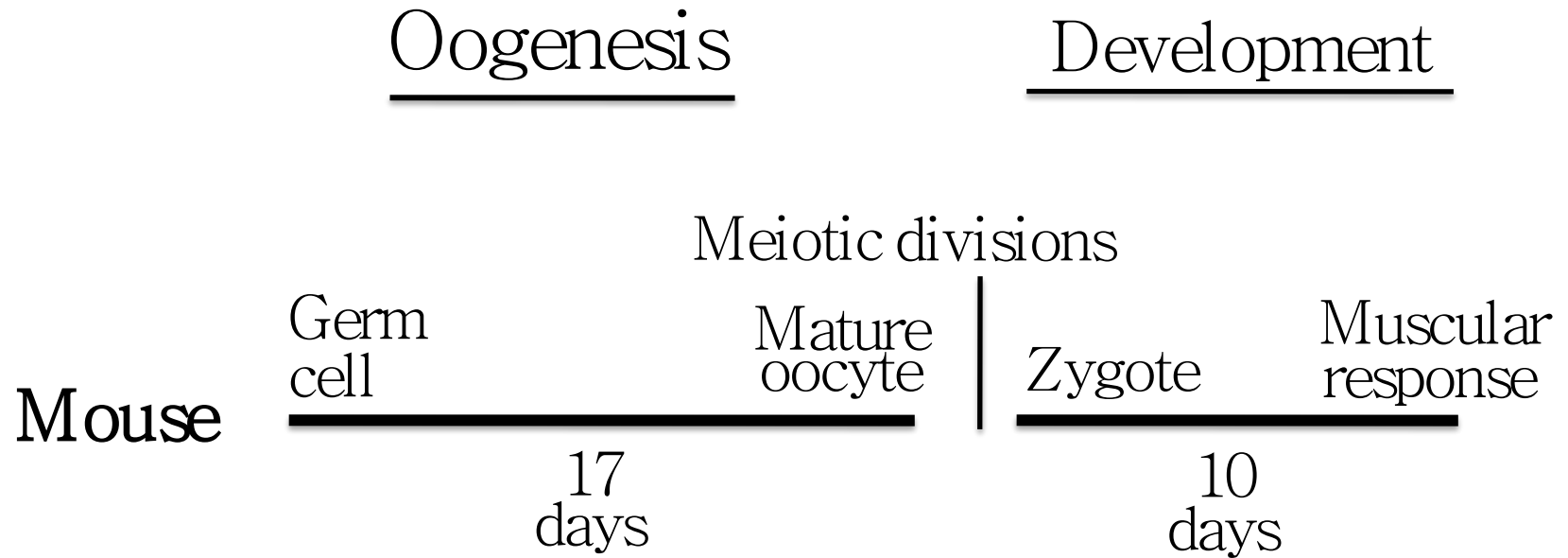


Newt chromosome II.

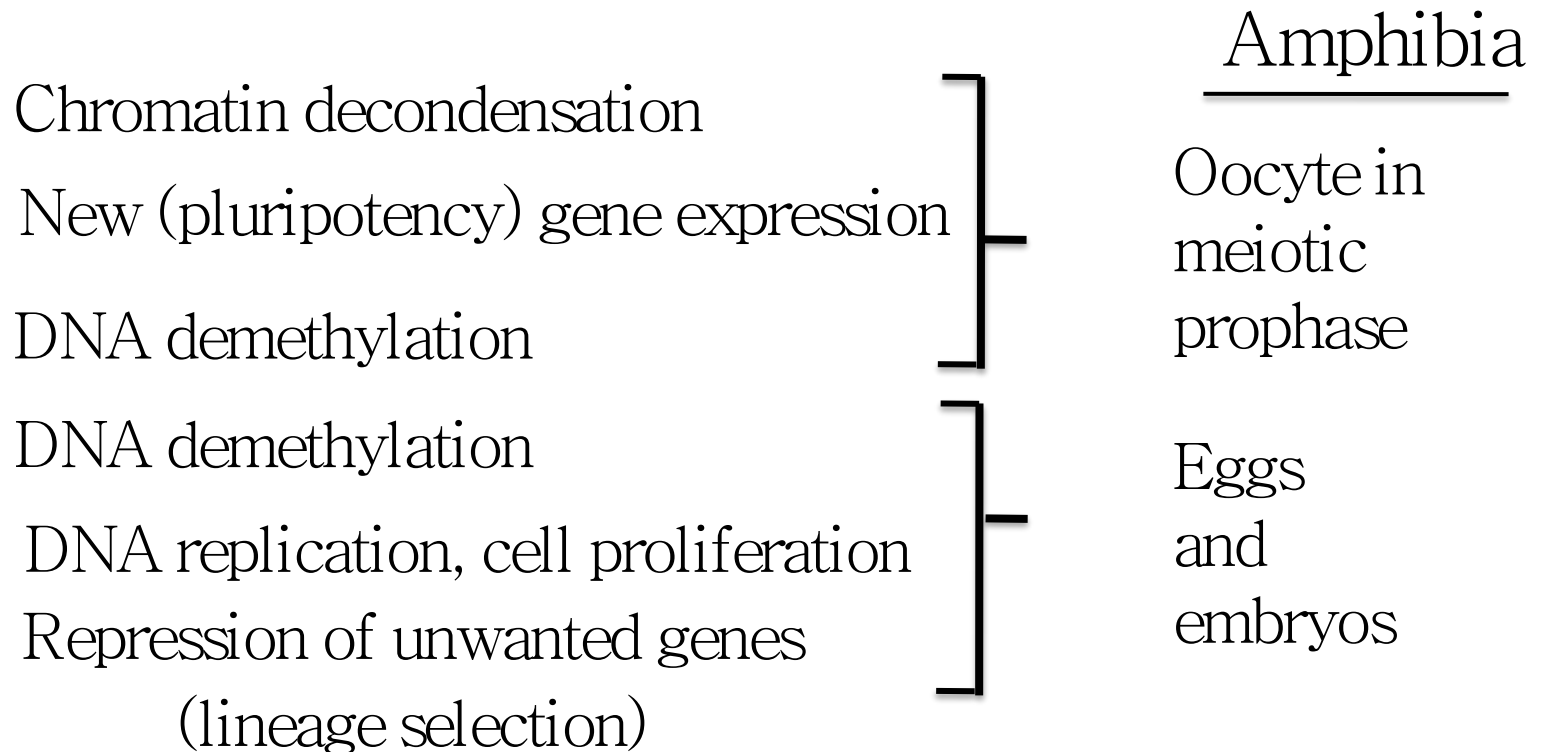
**Amphibian
lampbrush
chromosomes**



Oogenesis and development in the mouse



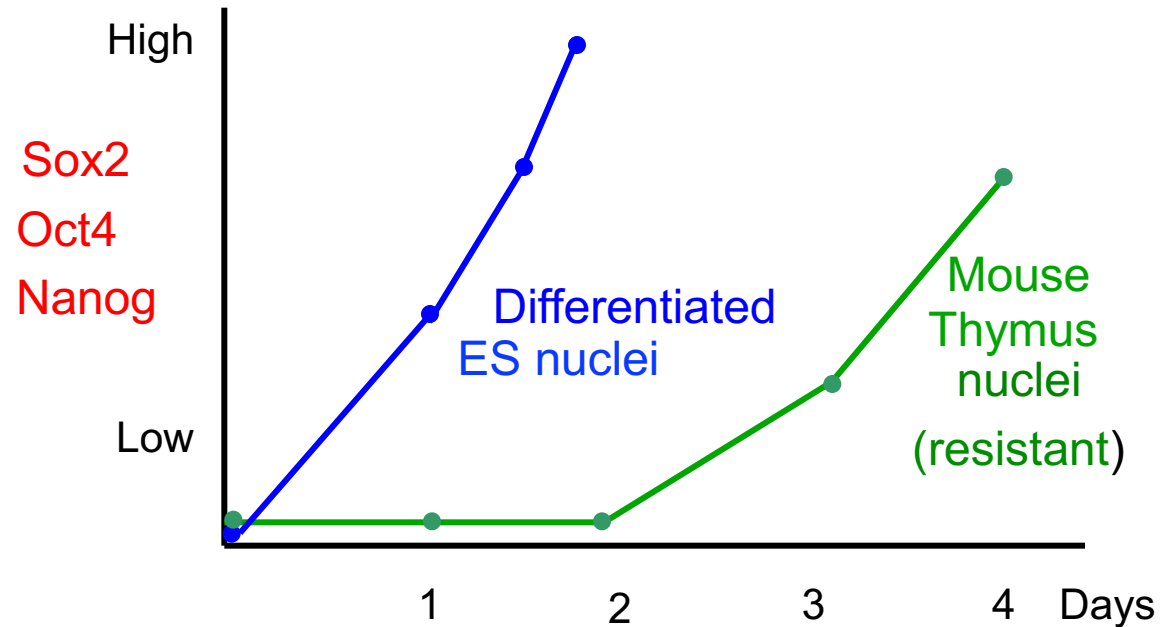
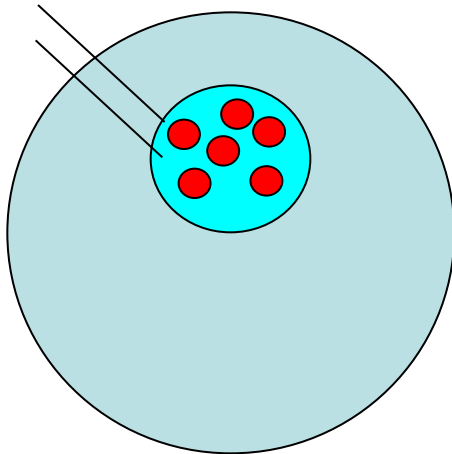
Major events in nuclear reprogramming



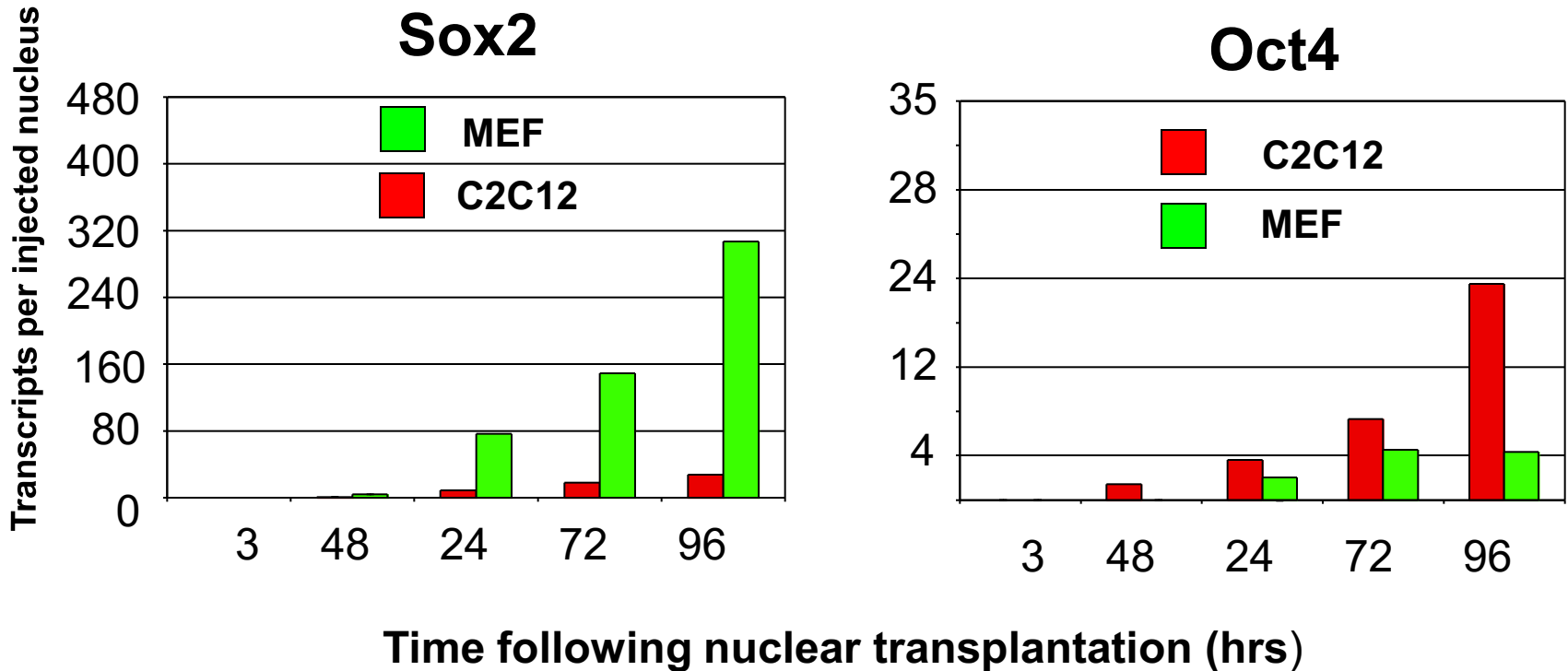
Stem cell genes are rapidly activated
in mammalian nuclei transplanted to *Xenopus* oocytes

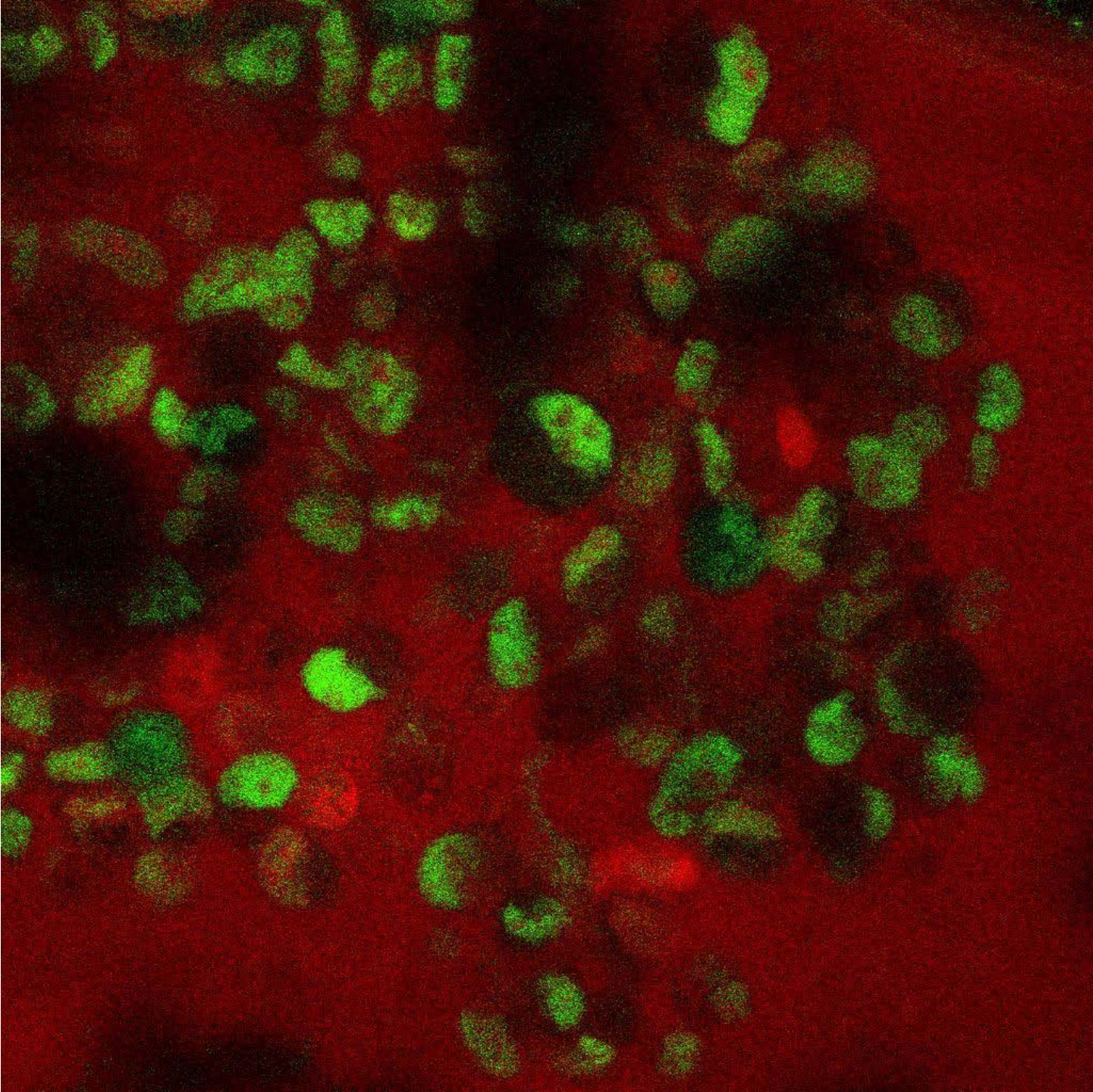
Nuclei of most differentiated cells resist reprogramming.

Mouse/human
somatic cell
nuclei

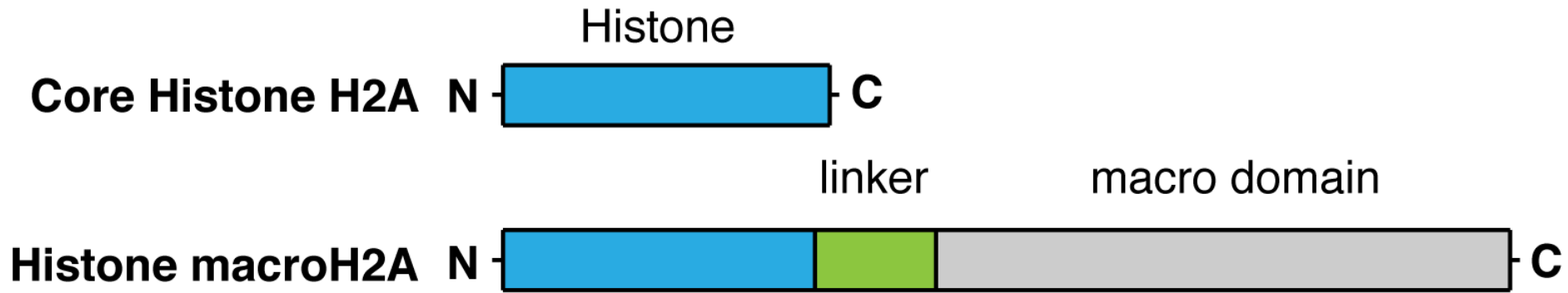


Resistance to reprogramming is pronounced when comparing different donor cell-types. [by up to 50X]





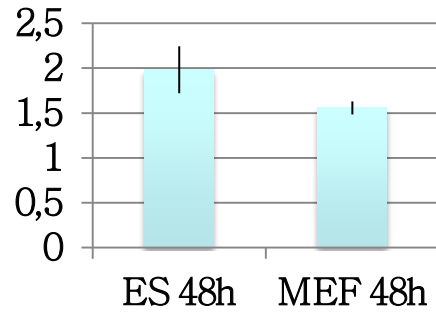
Histone variant macroH2A



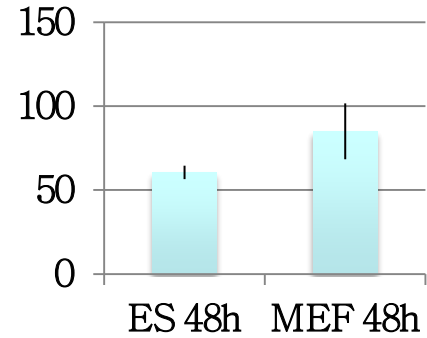
- macro domain = $2/3$ of macroH2A
- vertebrate-specific variant
- ‘hallmark’ of vertebrate heterochromatin

Examples of genes with restricted expression in MEF nuclei after transplantation to *Xenopus* oocytes

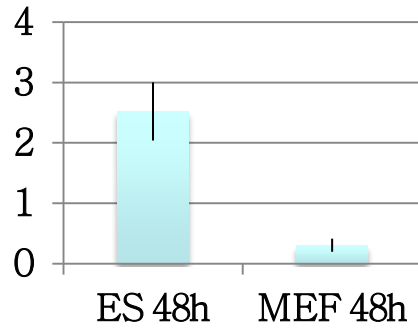
GAPDH (RIP)



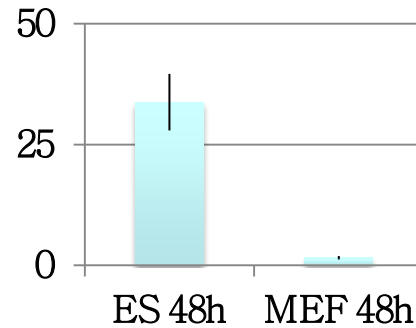
Gadd45 a (RIP)



Ooep (RIP)

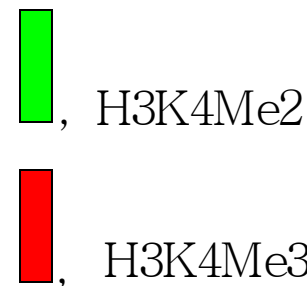
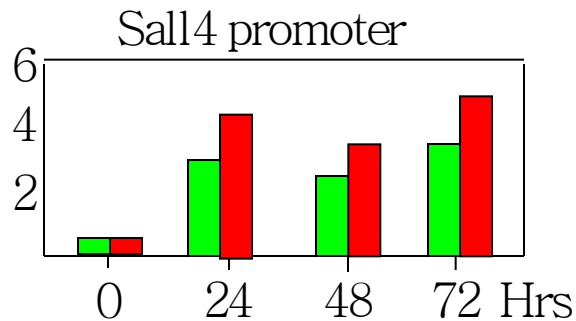
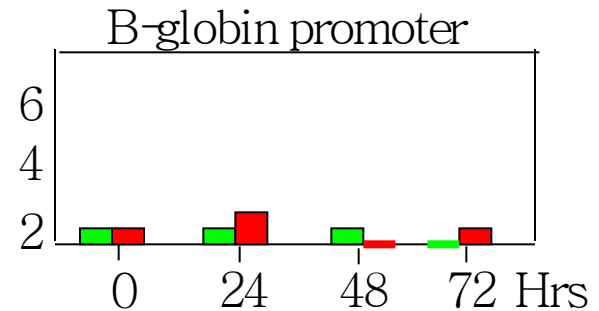
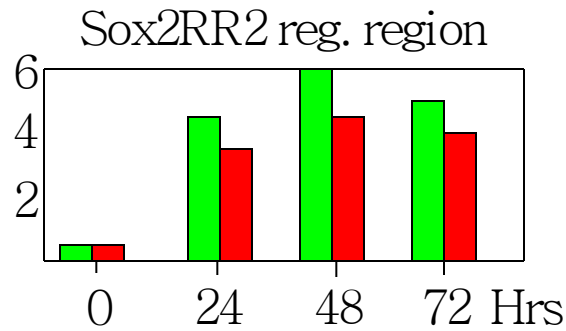
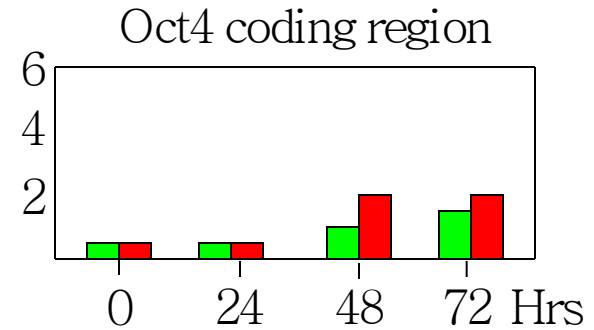
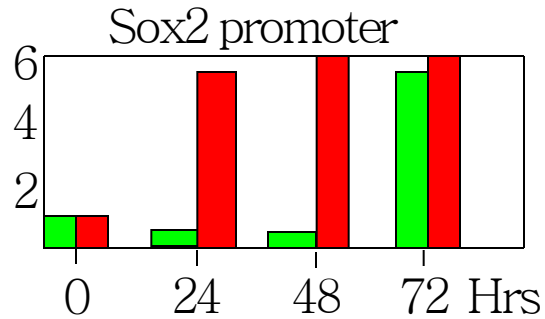


Prok2 (RIP)



Histones in gene control regions are methylated – Chip analysis

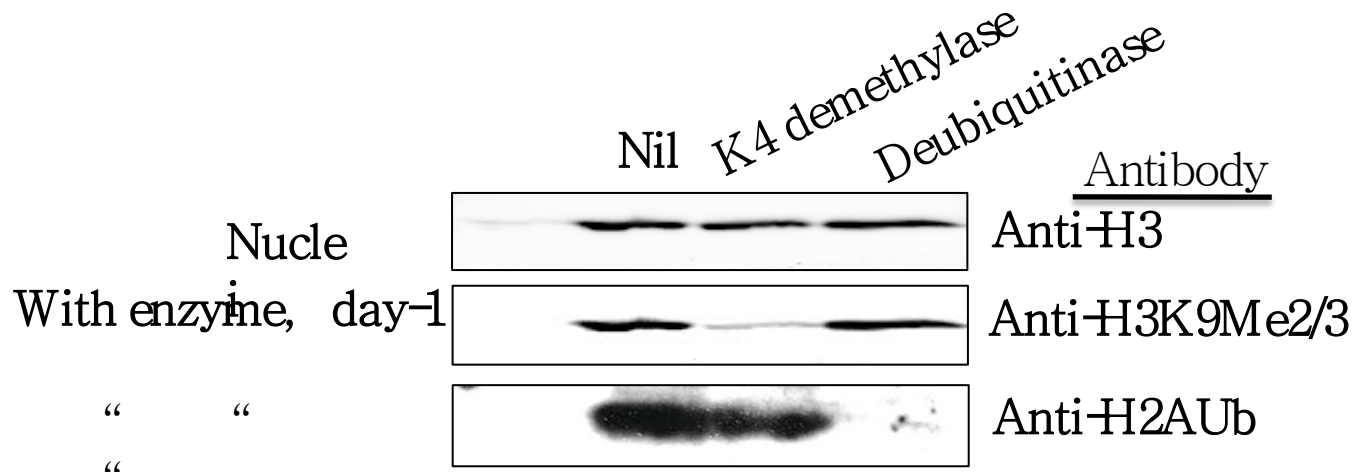
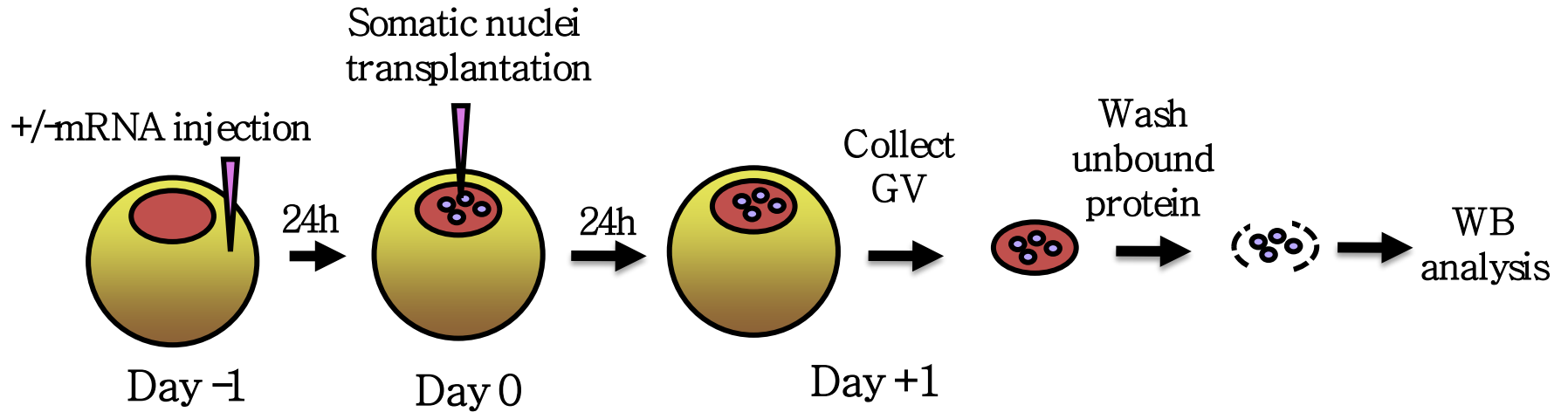
Nuclei from retinoic acid treated ES cells



Gene activation in somatic nuclei transplanted to oocytes is selective

		Number	%
Expressed in MEFs, But NOT in transplanted MEF nuclei	Repressed	7113	41
NOT expressed in MEFs, BUT in transplanted MEF nuclei	Activated	1176	9
Expressed in MEFs and in transplanted MEF nuclei	No change	3308	29

Histone modifiers overexpressed in the oocyte efficiently modify transplanted nuclear chromatin



Chromatin modifiers that alter the epigenetic state of transplanted nuclei

	enzymes	specificity
demethylase	Kdm1a	H3K4me2/1 H3K9me2/1
	Kdm2b	H3K36me2/1 H3K4me3
	Kdm3a	H3K9me2/1
	Kdm4d	H3K9me3/2
	Kdm5	H3K4me3/2
	Kdm6a	H3K27me3/2
deubiquitinase	Usp16	H2Aub
	Usp21	H2Aub
acetylase	Elp3	H3/H4
	Tip60	H3/H4

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Marta Teperek (Sperm progenitors)	Poland
Stan Wang	USA

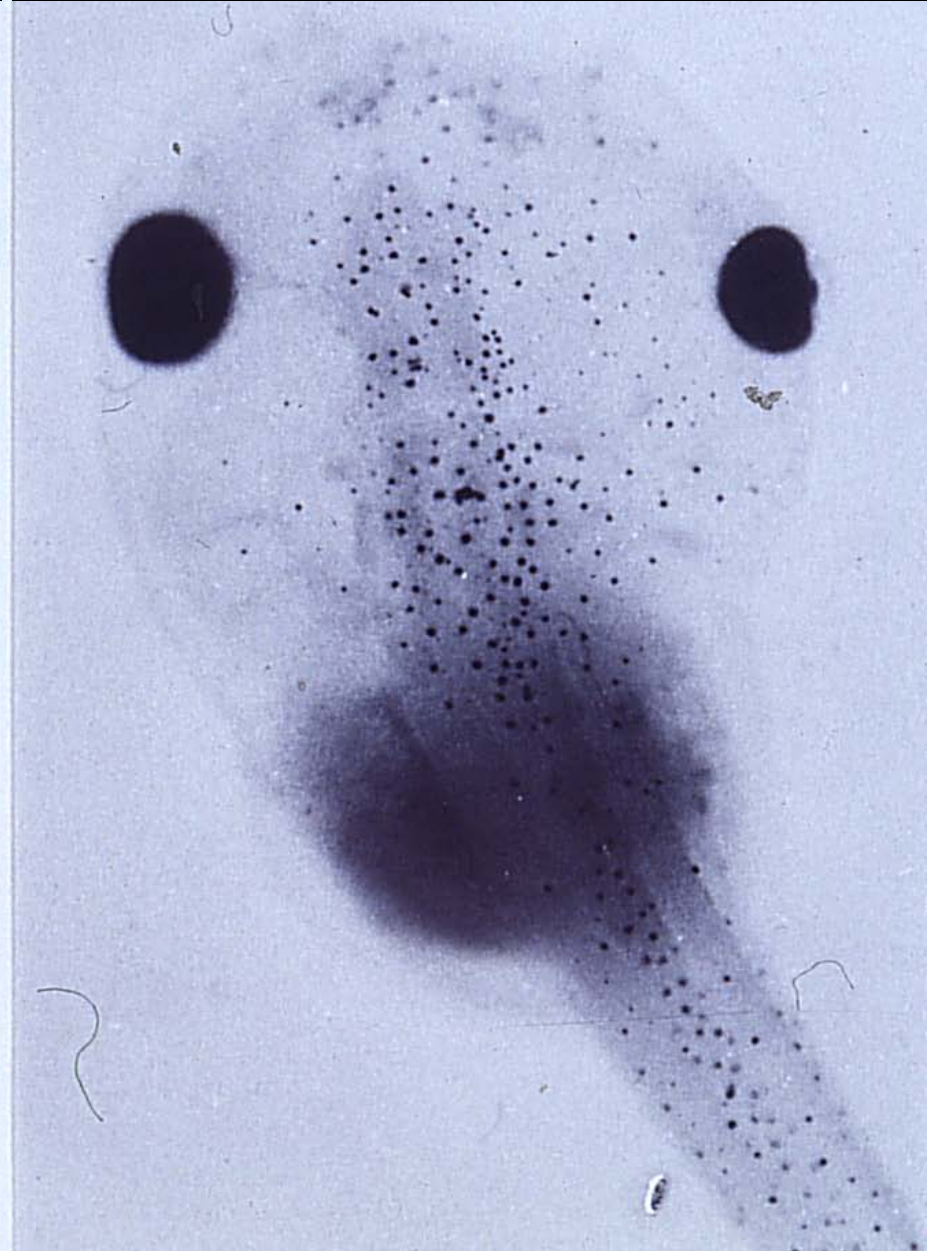
Welcome Trust

Medical Research Council

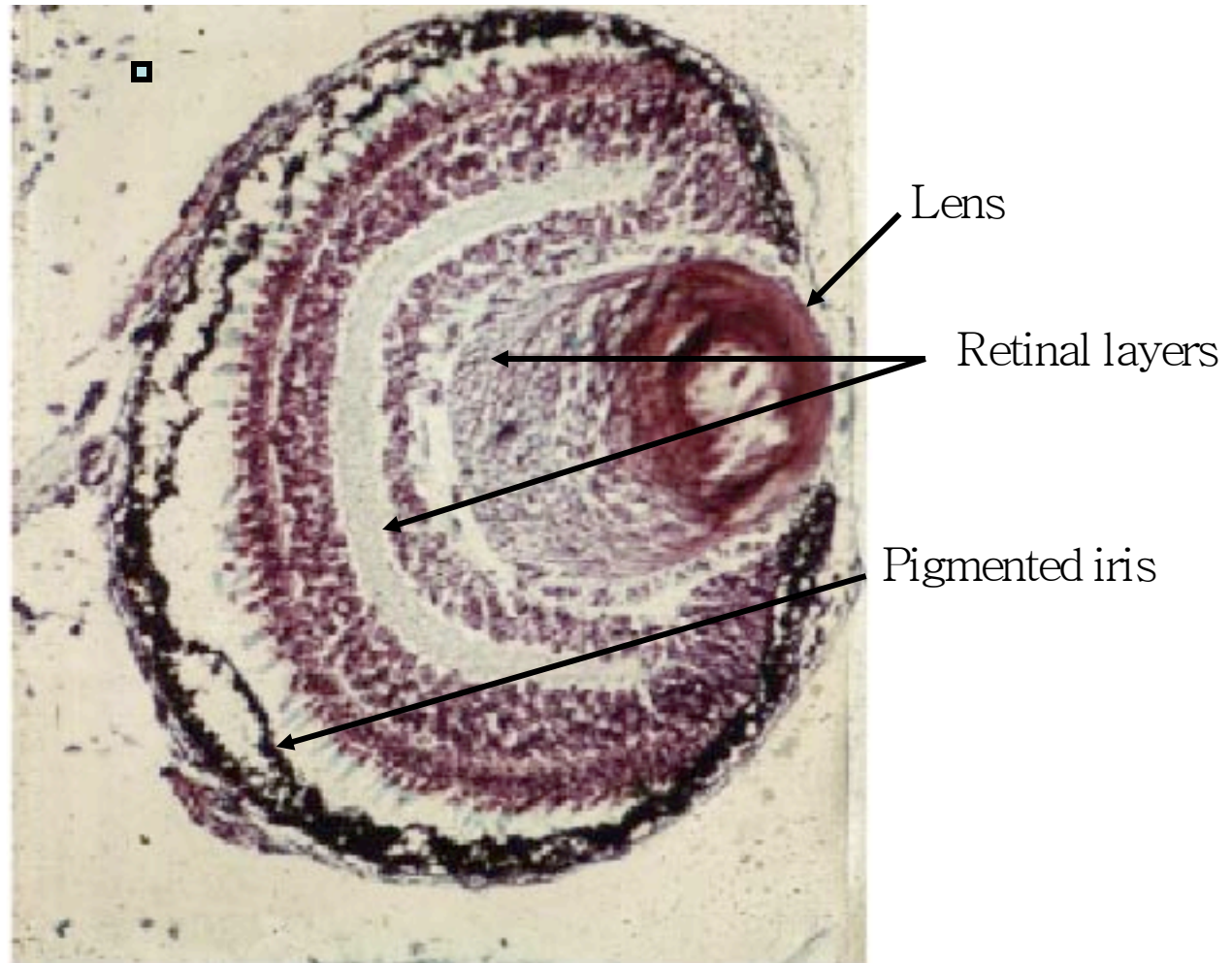
Tadpole from fertilized egg

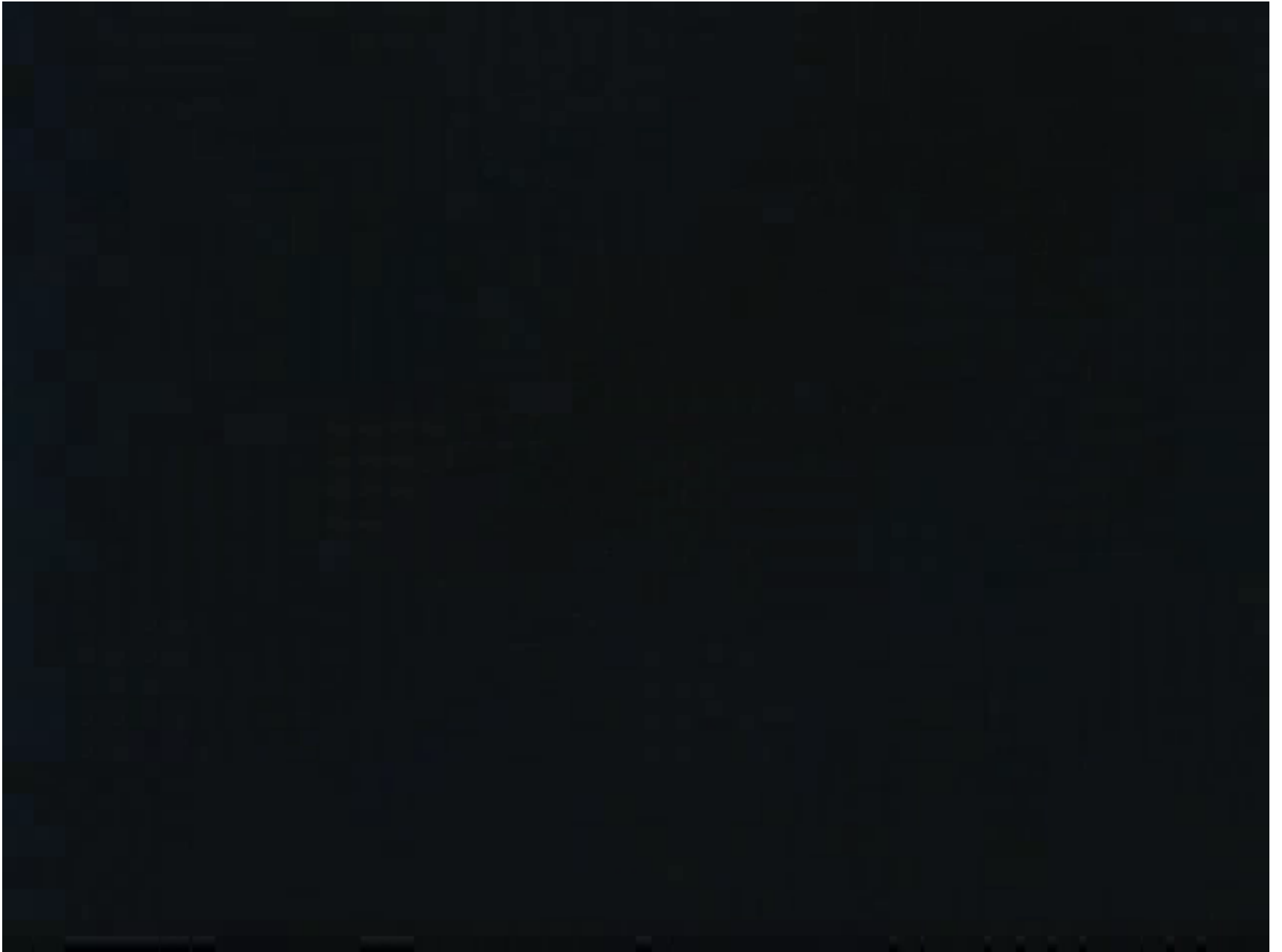


Tadpole cloned from a muscle cell

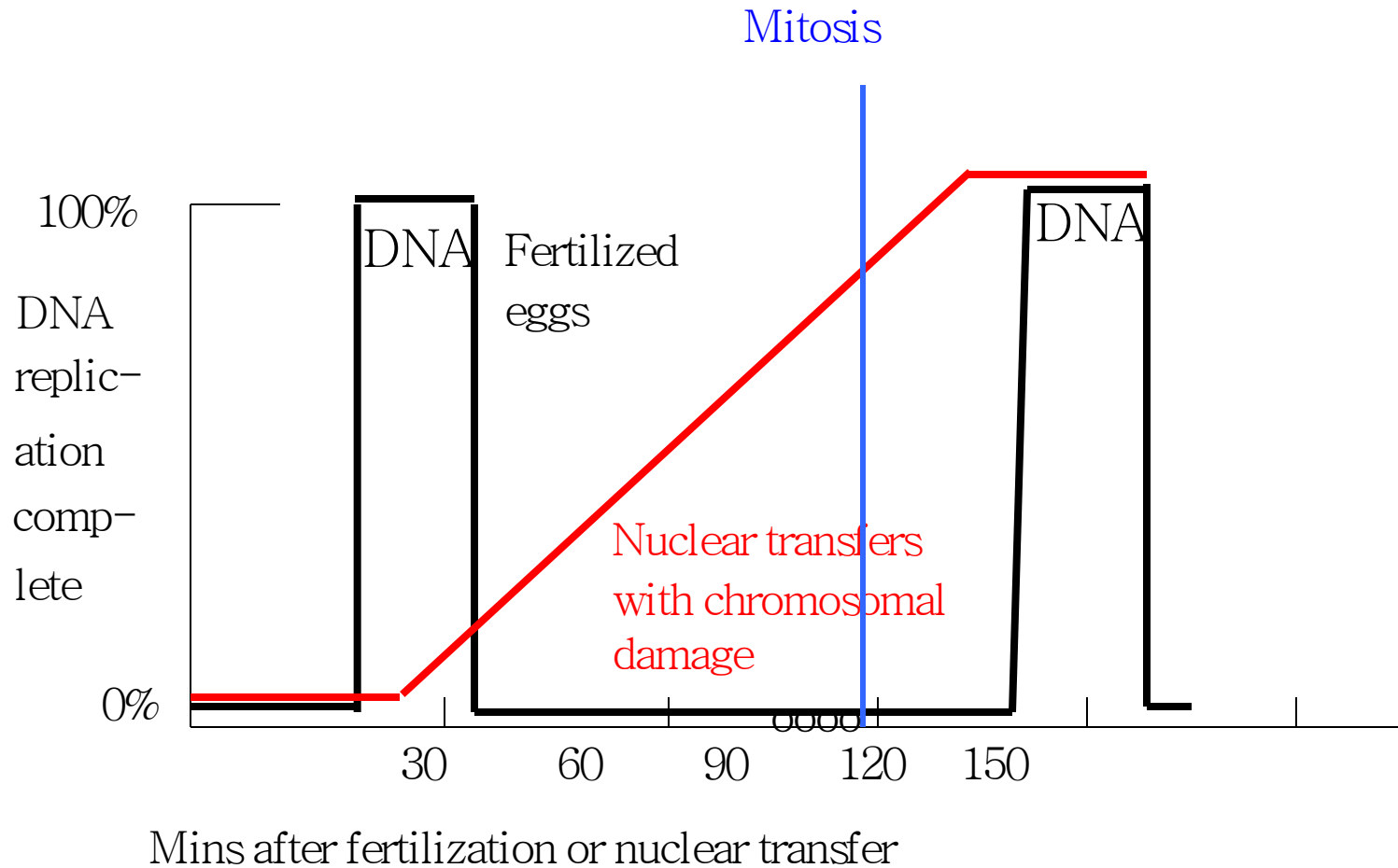


Normal eye
by cloning from a muscle cell

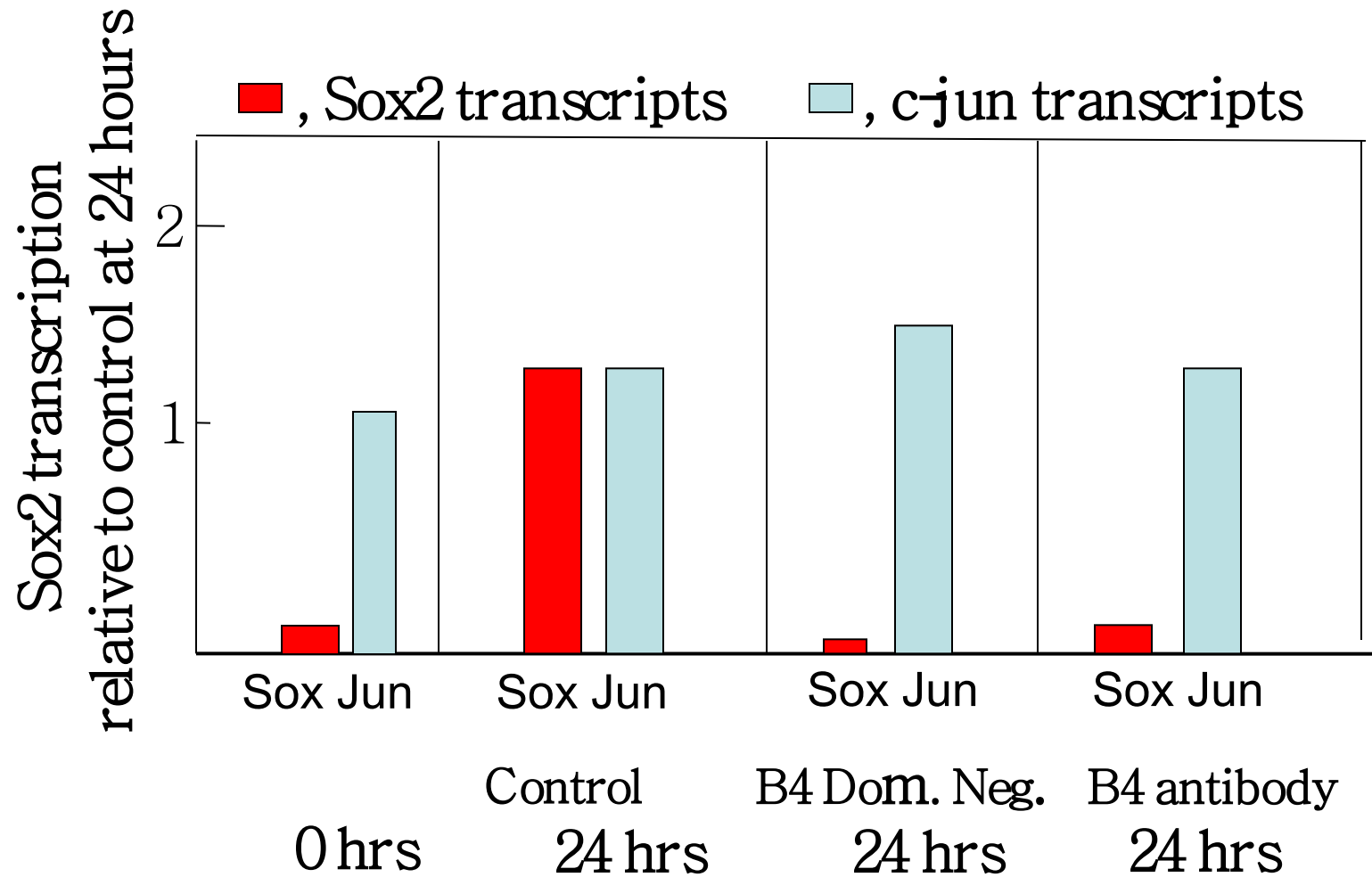




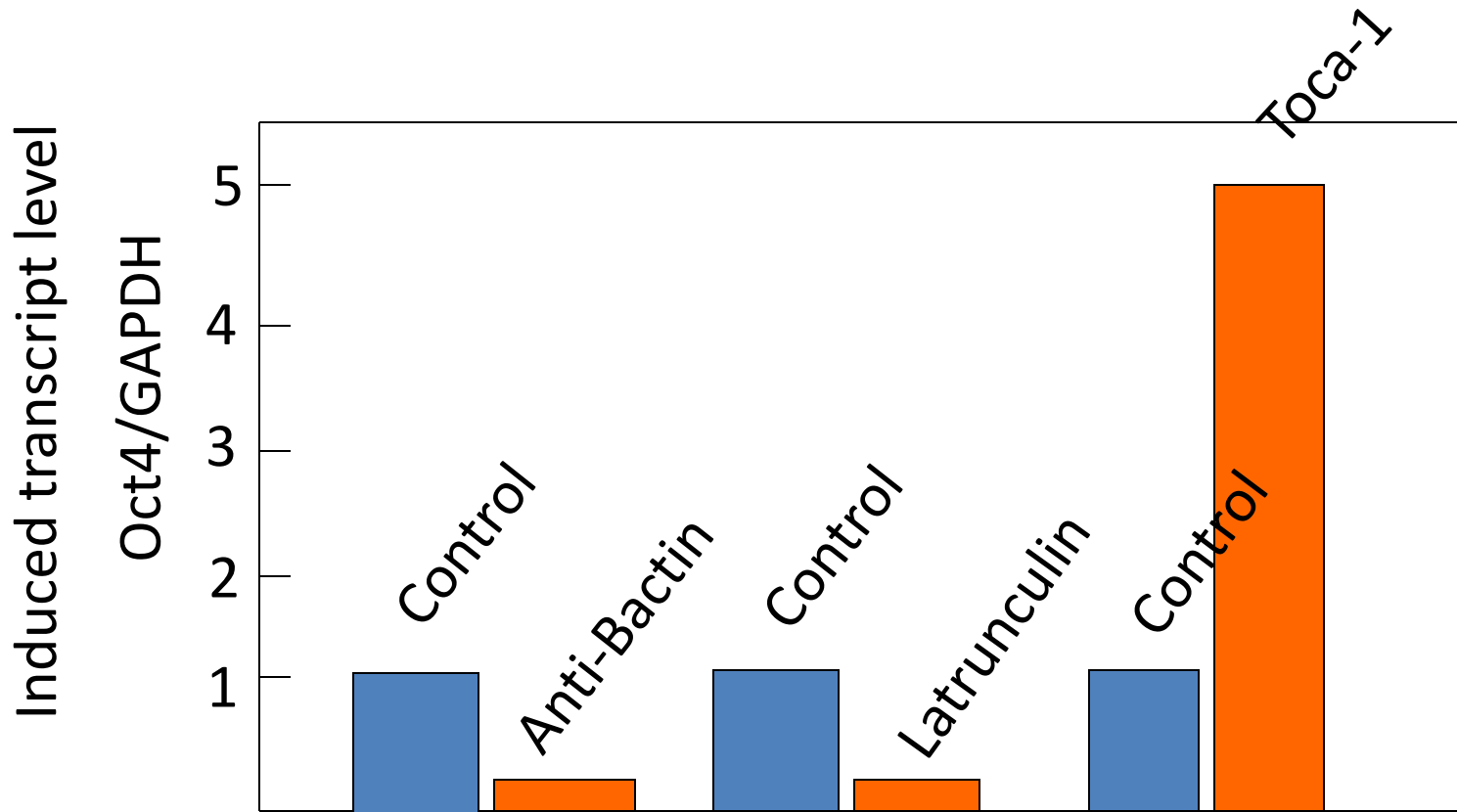
DNA replication is retarded in Amphibian somatic cell nuclear transfers



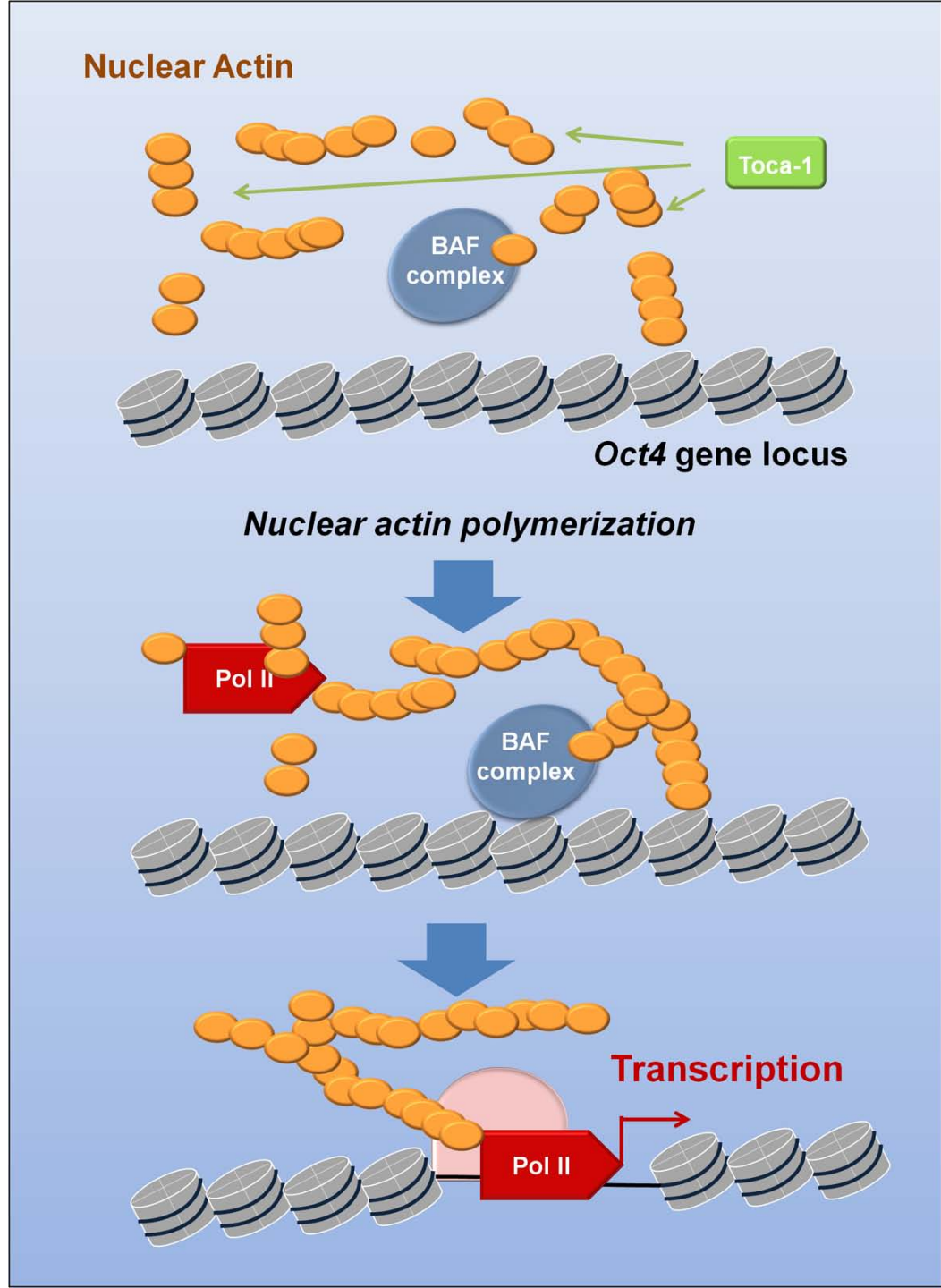
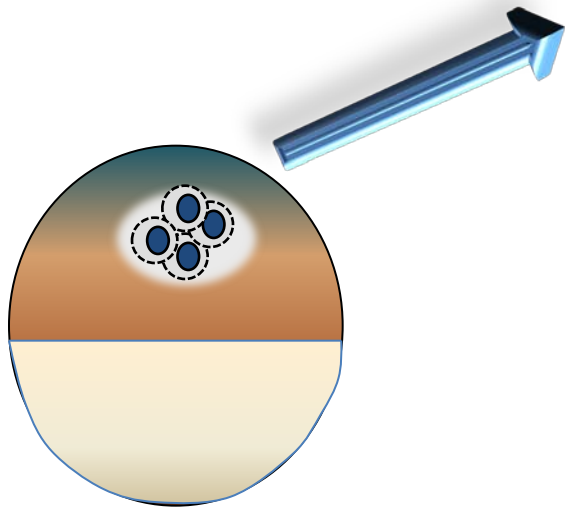
B4 histone is required for gene activation in oocytes



Transcription is enhanced by actin polymerization

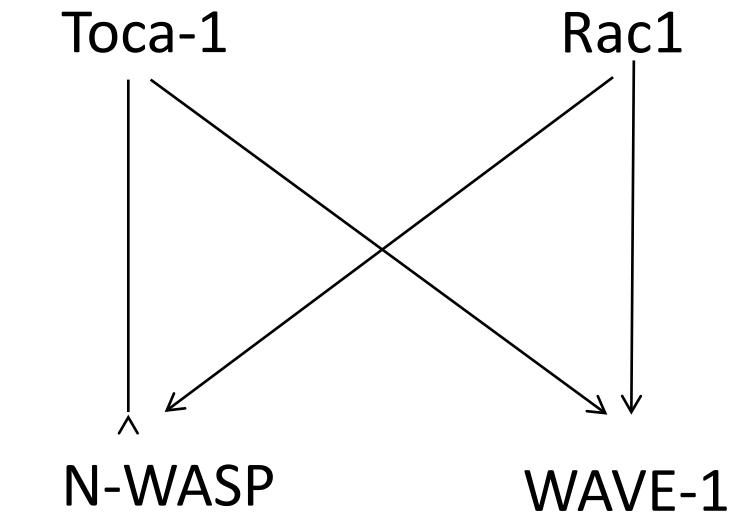


A model of nuclear actin function



Transcriptional activation is much enhanced by WAVE-1 (Wiskott-Aldrich syndrome)

Actin polymerization

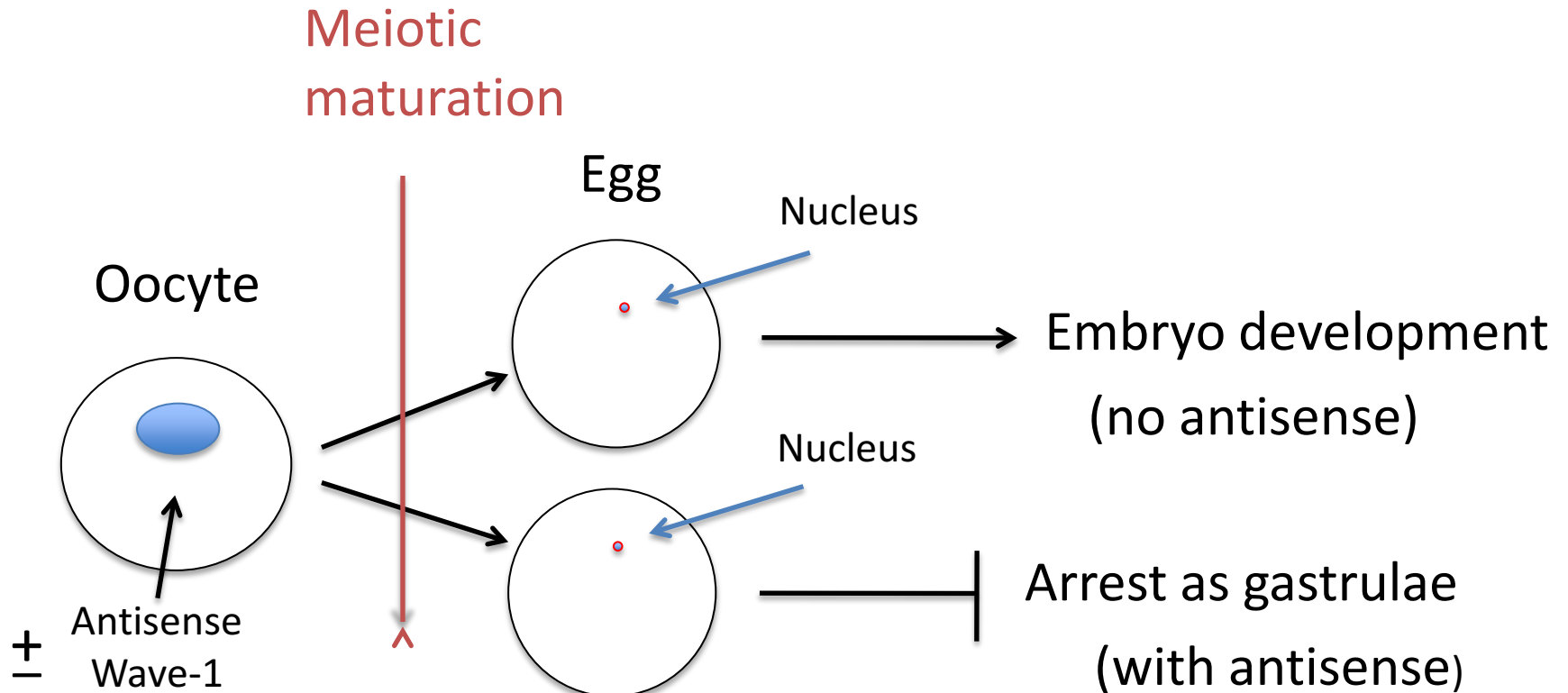


~~No effect on
Reprogramming~~

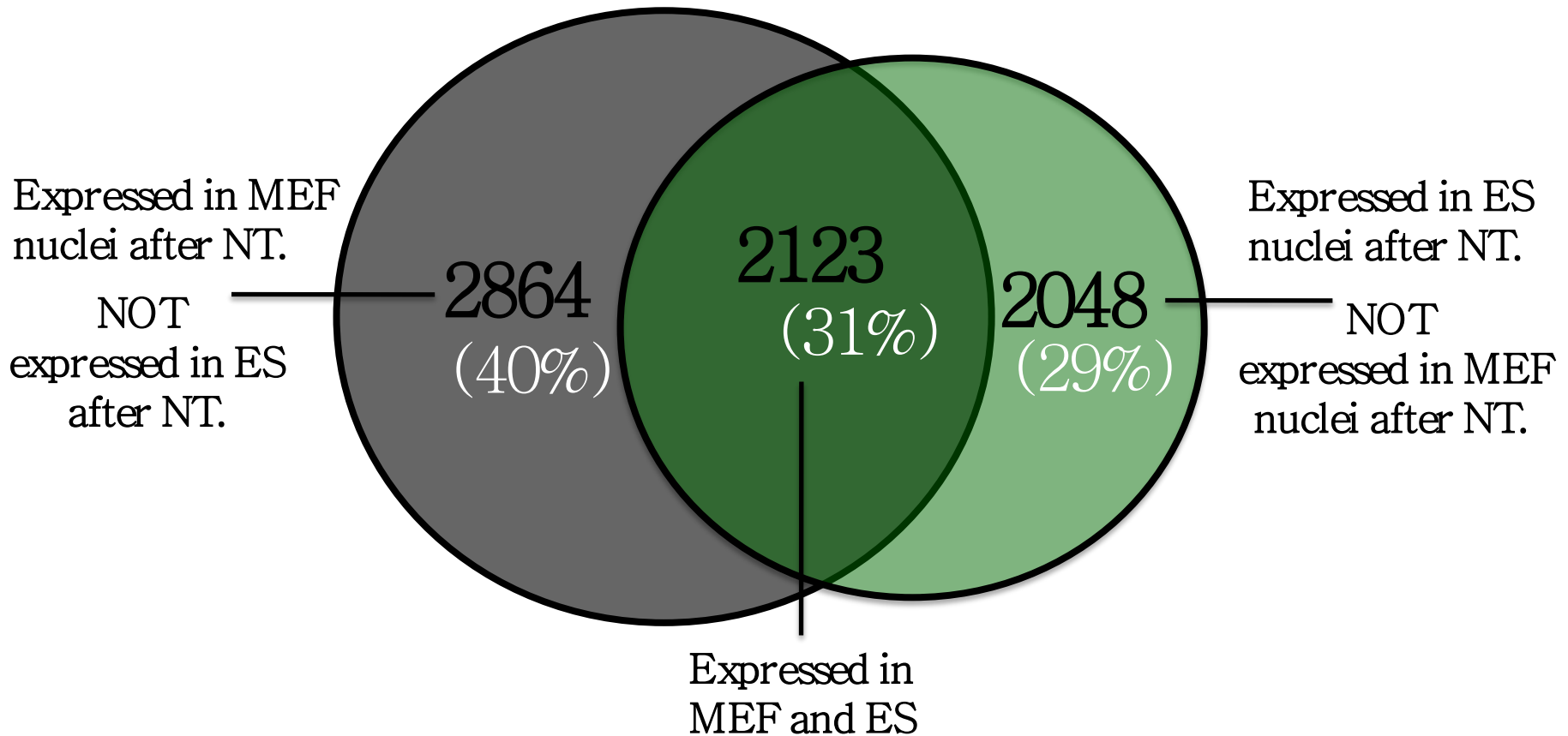
Enhances reprogramming seen by
pluripotency gene transcription in oocytes

WAVE-1 is required for zygotic genome activation and embryonic development

(Wiskott-Aldrich syndrome)

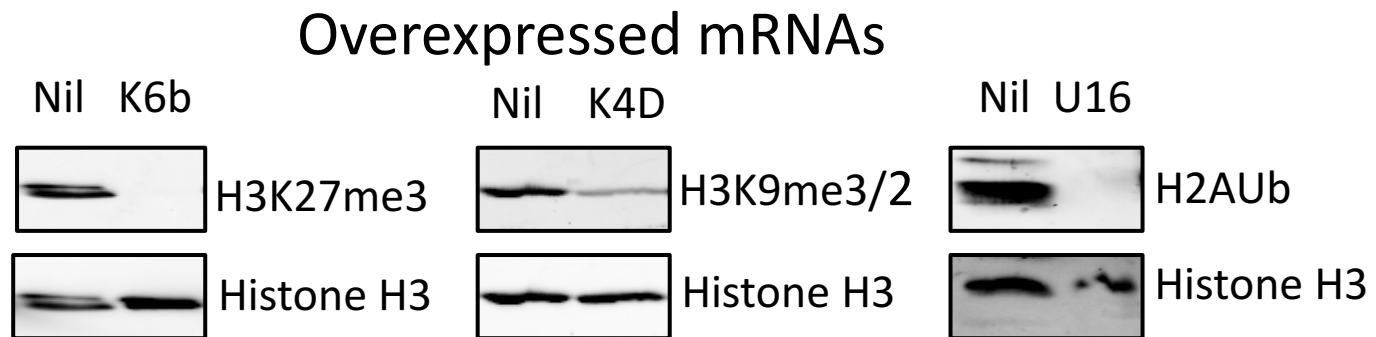


Genes with restricted expression in MEF or ES nuclei after transplantation to *Xenopus* oocytes



Histone modifications in nuclei can be changed after transfer to oocytes

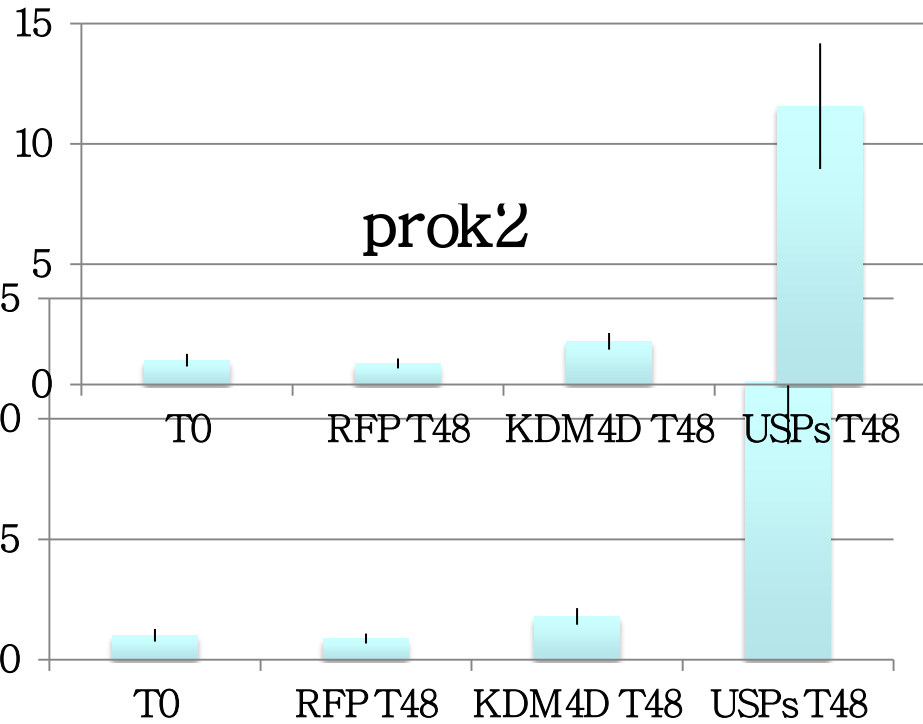
0 hour: mRNA injections. 24 hours: nuclear injections.
48 hours: reisolation of injected nuclei and Western analysis.



K6b, H3K27 demethylase. K4D, H3K9 demethylase. U16, H2A deubiquitinase.
Western blots to show loss of histone modifications 48 hours after mRNA injection.

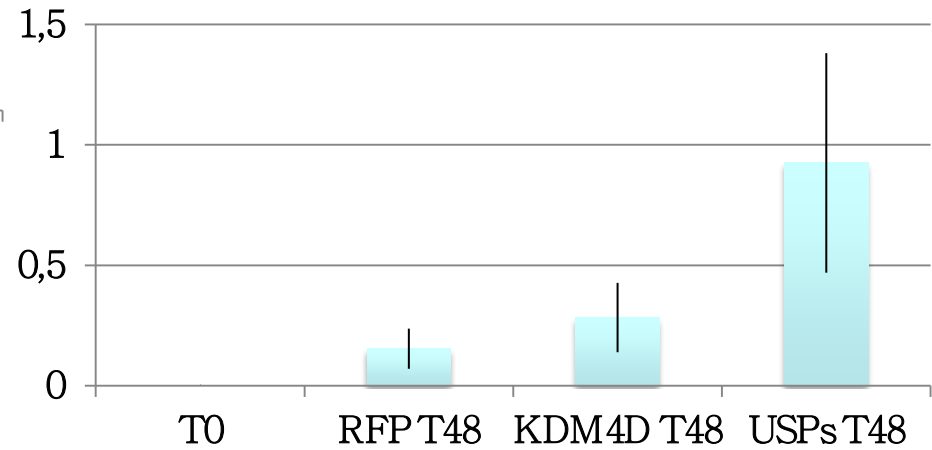
Overexpression of H2A deubiquitinases removes restriction

prok2



prok2

oep



Transcriptional reprogramming

CONCLUSIONS

Eggs and oocytes have a very high content of histone H3.3.

Histone H3.3 prolongs transcription of somatic nuclei in oocytes.