



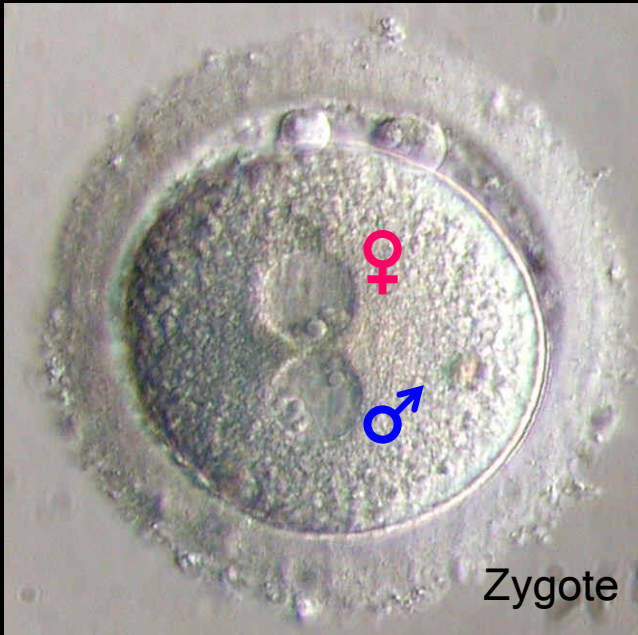
Nuclear and Chromatin Changes Accompanying Differentiation

Rui Pires Martins, PhD

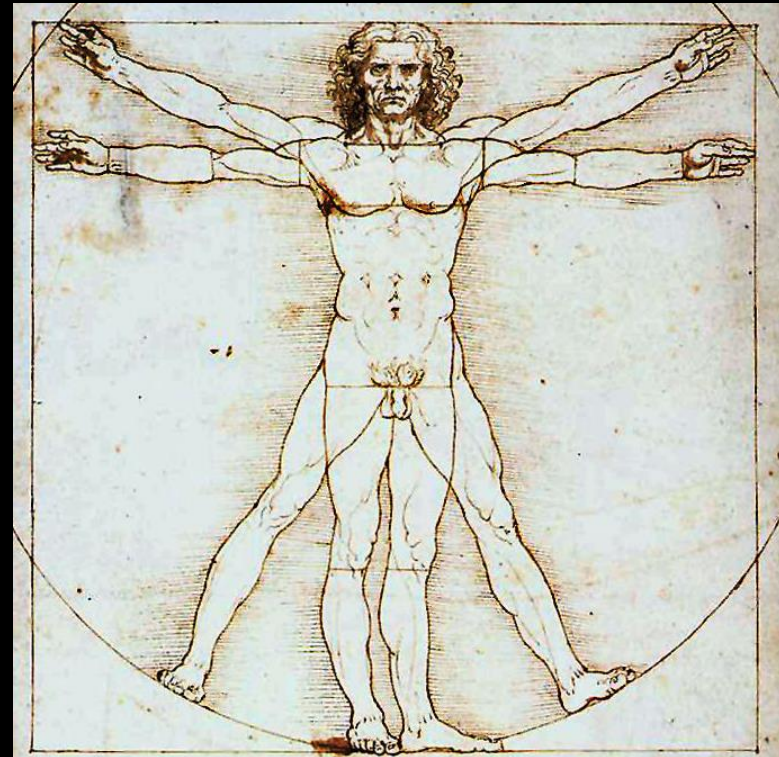
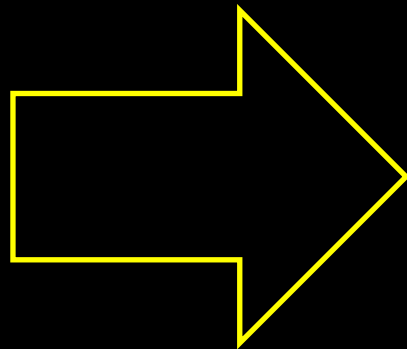
Outline of today's talk...

- Brief introduction to:
 - a. development/differentiation
 - b. mechanotransduction
- The nucleus, from the inside-out.
- Chromatin architecture
- Changes to chromatin and nuclear architecture that accompany differentiation in ES cells (including original work)





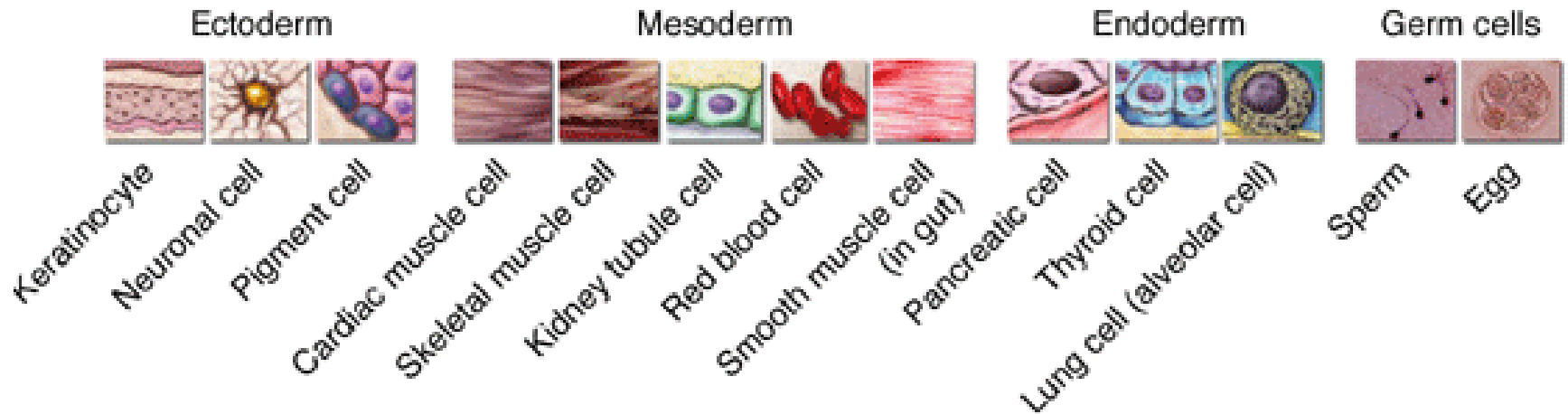
1 cell



≈50-100 trillion (10^{14}) cells



There are potentially hundreds to thousands¹ of different cell *fates* or types in the Human Body

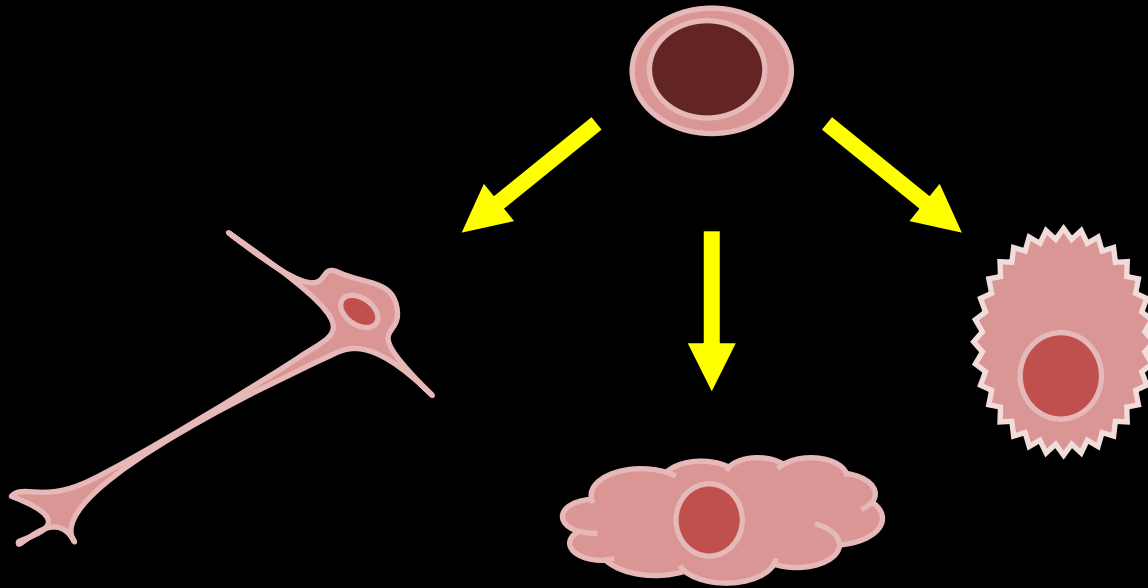


1. Cytokines & Cells Online Pathfinder Encyclopaedia. (<http://www.copewithcytokines.de/cope.cgi?key=cell%20types>)



Differentiation

is the progression from a naïve or unspecialised cellular state, to a more specialised state.



Potency

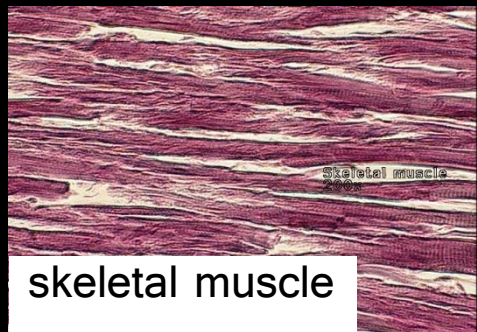
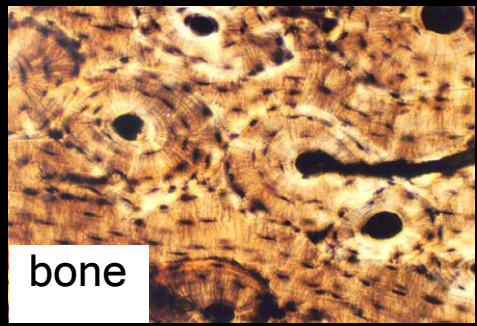
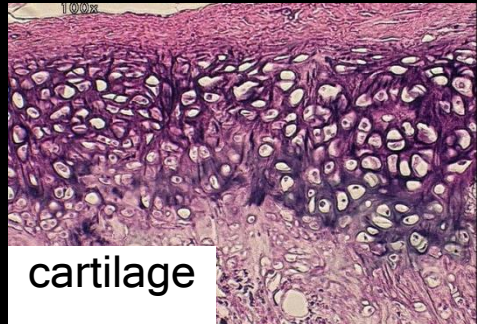
describes the developmental potential of a cell. Stem cells that can adopt multiple fates are said to be **pluripotent**.





DIFFERENTIATION

REPROGRAMMING



D E V E L O P M E N T A L P O T E N T I A L C O N T I N U U M

TOTI POTENCY / PLURIPOTENCY

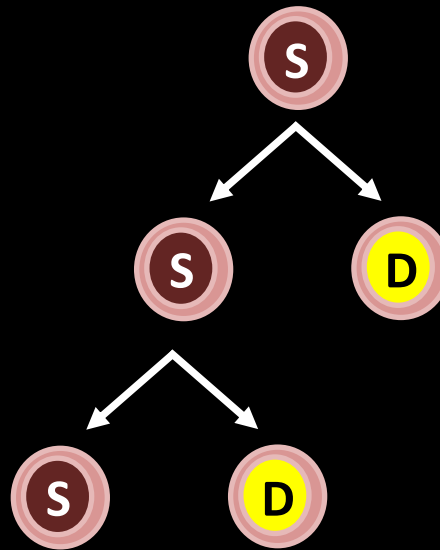
TERMINAL DIFFERENTIATION



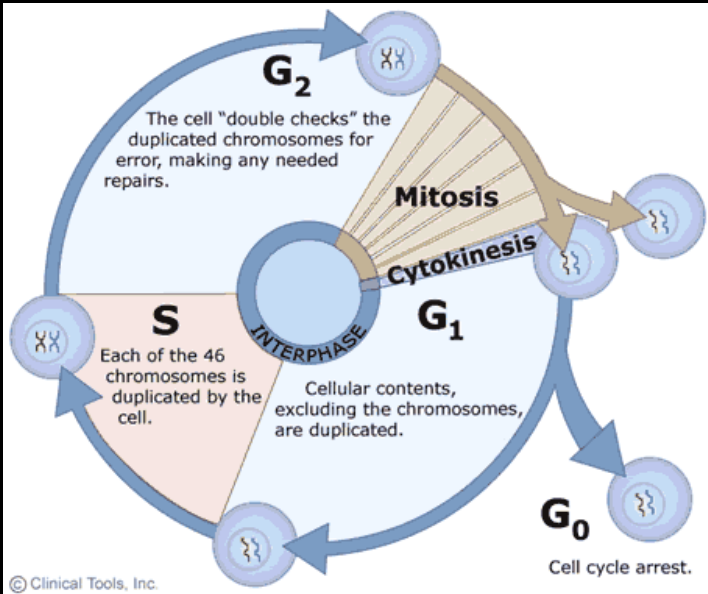
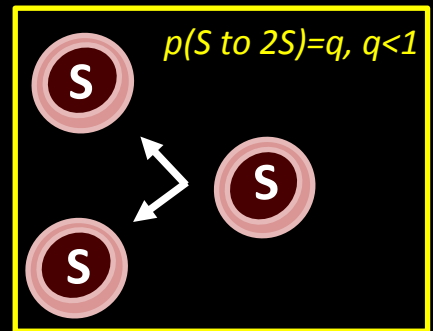
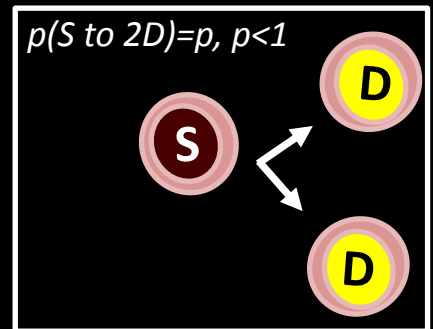
Self-renewal EXPAND

Embryos begin as a single cell and develop into multicellular organisms. Through mitotic cell division, cell numbers can expand exponentially, but self-renewal ensures that a stem cell population is maintained.

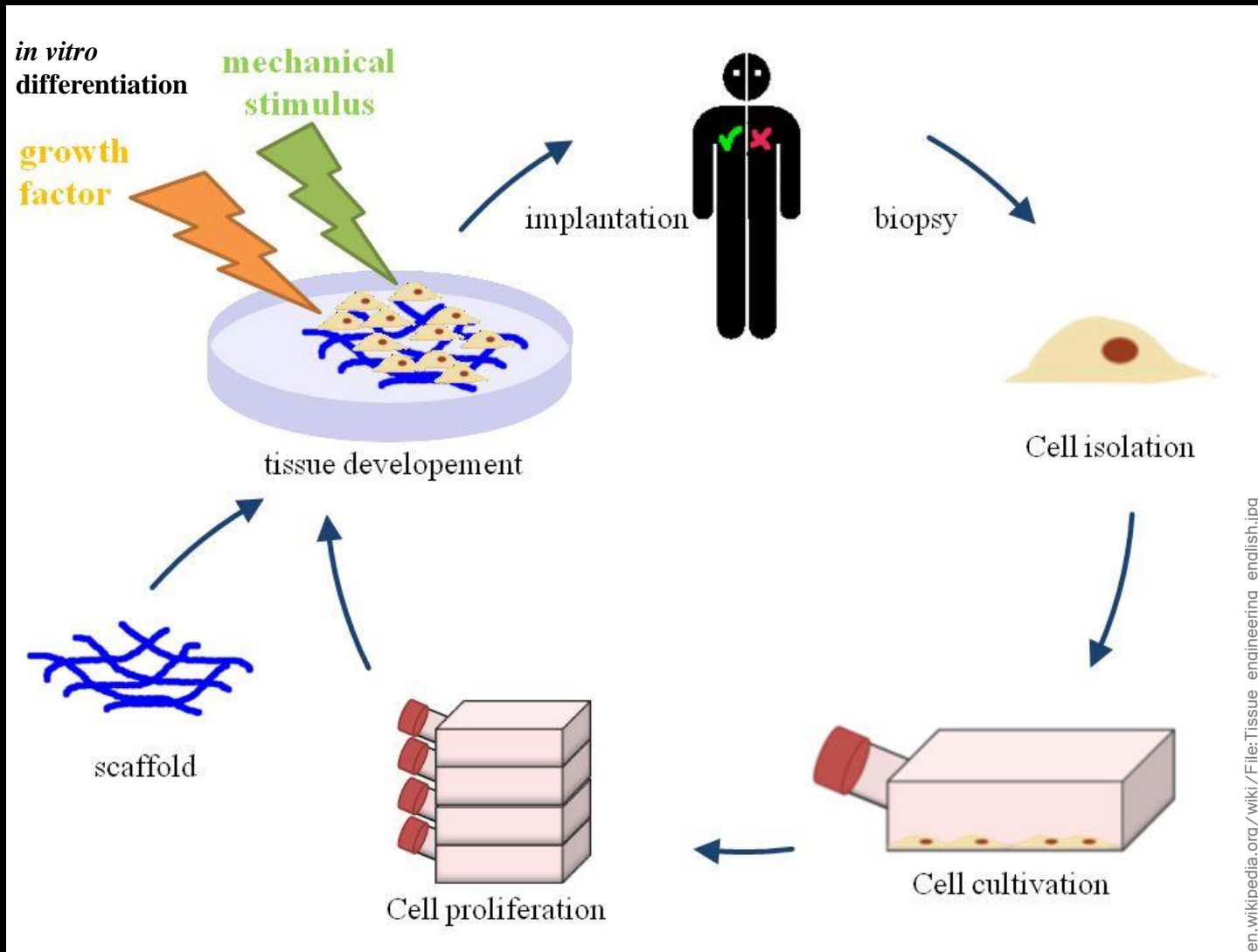
Asymmetric Division



Stochastic Mechanisms

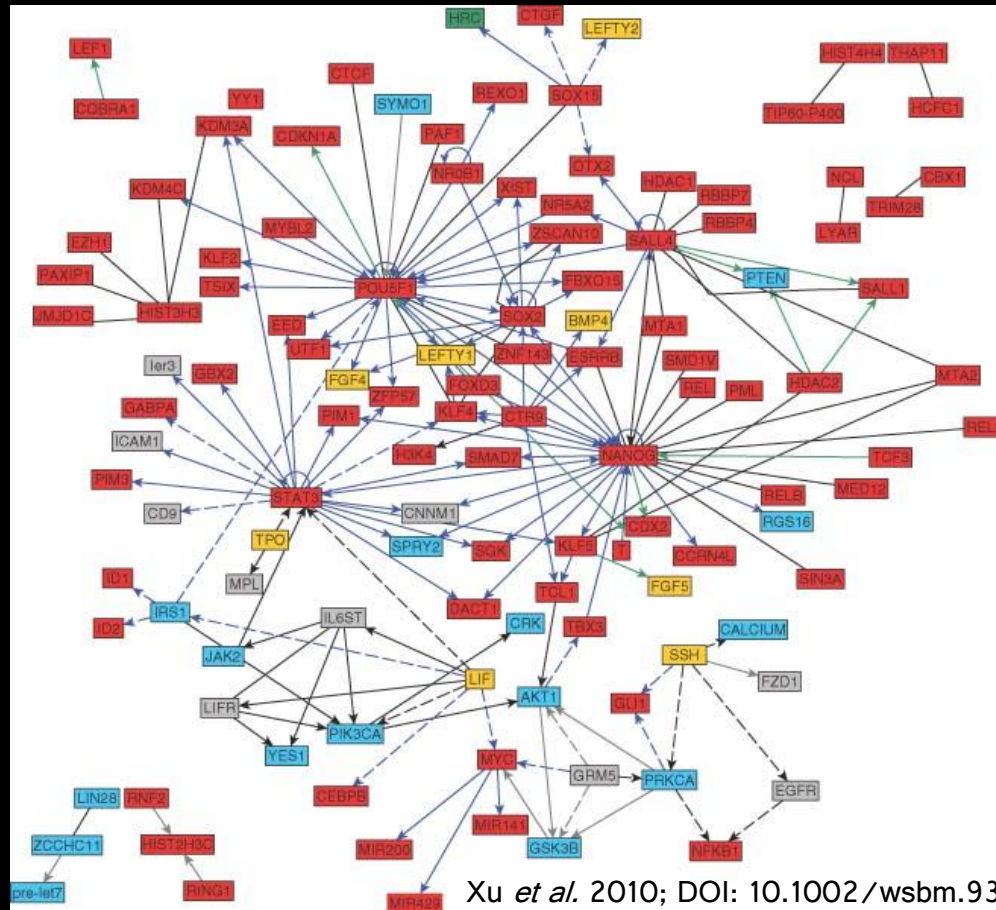


tissue engineering work-flow



Complex genetic/protein interaction pathways that govern the maintenance of pluripotency in stem cells are well characterised.

PLURIPOTENCY IN MESCS



HOWEVER, how mechanical signals feed into these and other differentiative circuits are very much open questions.



Mechanotransduction

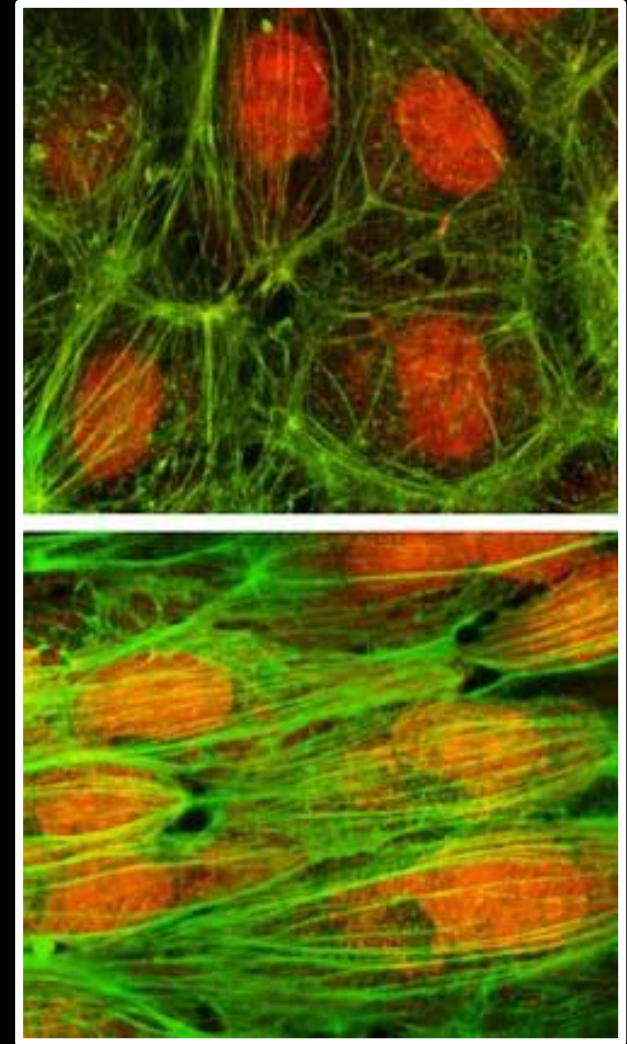
The process by which mechanical stimuli are converted to intracellular signals that control cellular physiology, and disease.

Sheer force on endothelial cells lining vessel walls

The force of blood flowing past an endothelial cell promotes changes in cell morphology: cells reorient their cytoskeletons in the direction of the blood flow, and cells send chemical signals to neighbouring endothelial cells that result in changes to local vessel stiffness.

Mechanotransduction may alter cellular physiology by stimulating an ion channel (e.g. Ca^{2+}) or activation of cellular signalling through cell surface receptors (e.g. integrins in the cell membrane)

Mechanical forces can also be transmitted along the various elements of the cytoskeleton on through to the nucleus.



Cytoskeleton (Actin), Nuclei (DNA)



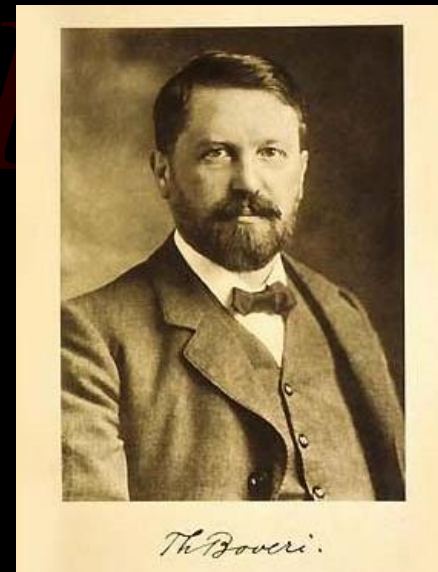
Theodor and Marcella Boveri

German and American cell and developmental biologists who worked in the late 1800s and early 1900s

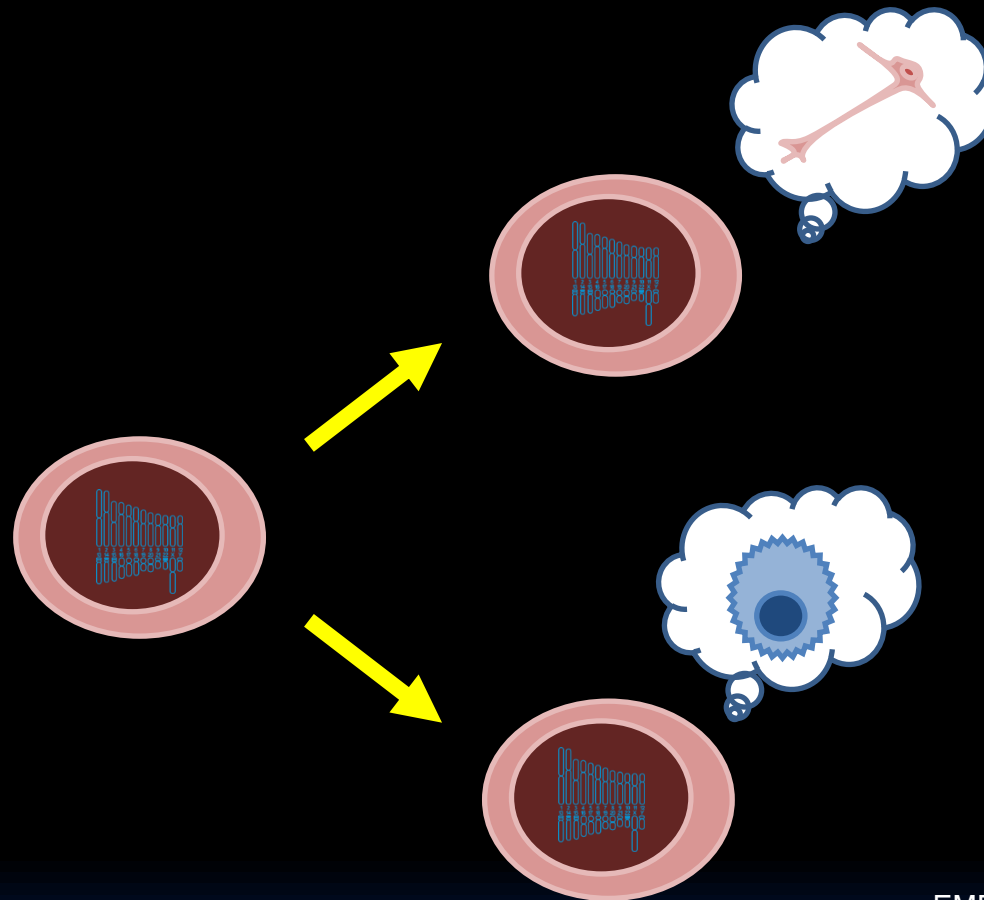
The Boveris' seminal work in roundworms and sea urchins helped confirm the nucleus, and indeed chromosomes as the source of heredity.

The prevailing wisdom governing development and differentiation was that only germ cells received a complete set of chromosomes as these needed to reproduce the whole organism. Differentiated cells of other lineages only received the genetic information pertinent to their function.

The Boveris' Chromosome Theory demonstrated that there was equal (genomic) contribution between parents (at fertilisation) and that all, not just a subset of the chromosomes were transmitted from germ cells to somatic cells during differentiation.



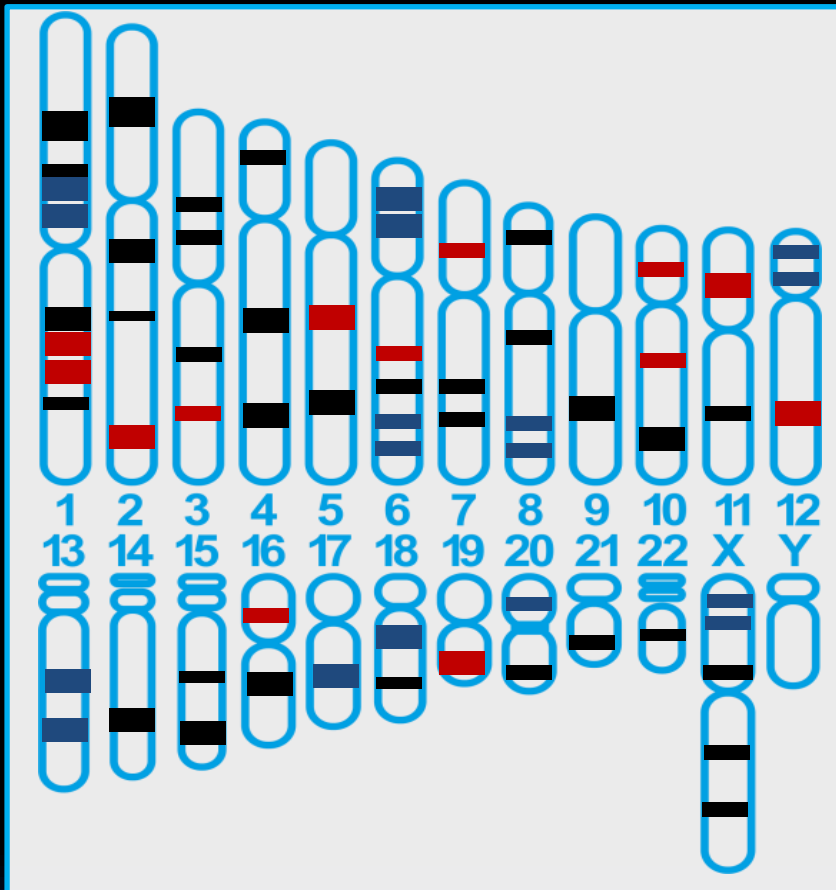
Fundamentally, **differentiation** operates at a genetic level. Genomically, cells in the body are clonal. What *differentiates* one cell type from another is the how each type utilises that genome.



Humans have 23 pairs of chromosomes, one inherited from each parent.

These are composed of 22 *autosomes*, and two sex chromosomes, X and Y.

These chromosomes code for approximately 21+K genes. That are arranged enigmatically through the genome. This is only 1% of the total information coded within.



Housekeeping Genes:

govern basic cell function, structure and metabolism. These generally function in ALL cell types.

Lineage Specific or

Determinant Genes: genes that impart *fate* ie. Turn on a genetic programme or contribute form or function to a cell lineage

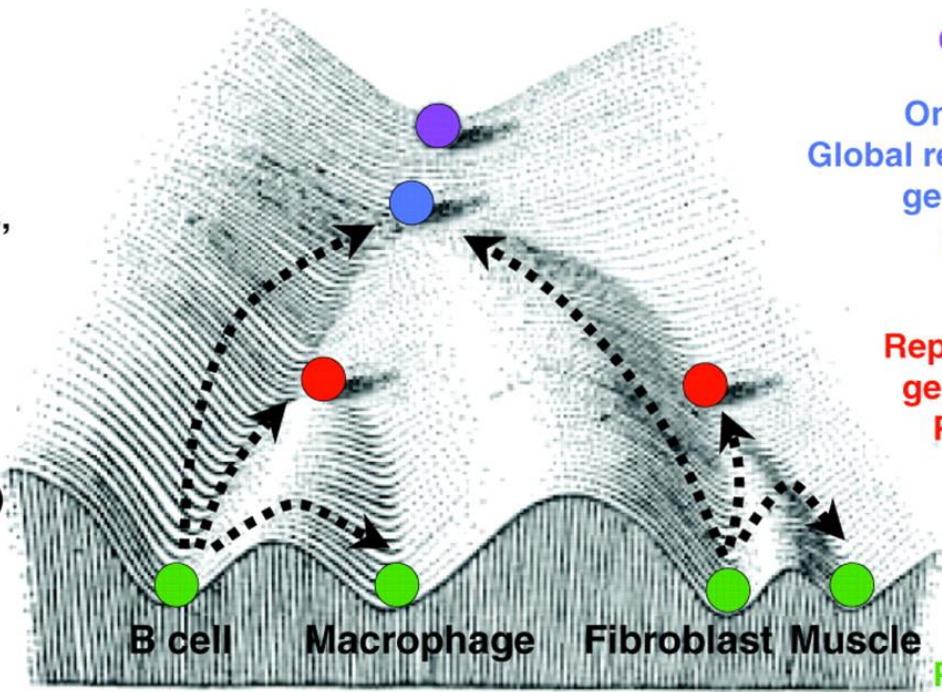


Waddington's Canalisation Concept that cell fate is progressively fixed/restricted by epigenetics

TOTIPOTENCY / PLURIPOTENCY
 DIFFERENTIATION
 TERMINAL

Developmental potential

- Totipotent**
Zygote
- Pluripotent**
ICM/ES cells, EG cells, EC cells, mGS cells, iPS cells
- Multipotent**
Adult stem cells (partially reprogrammed cells?)
- Unipotent**
Differentiated cell types

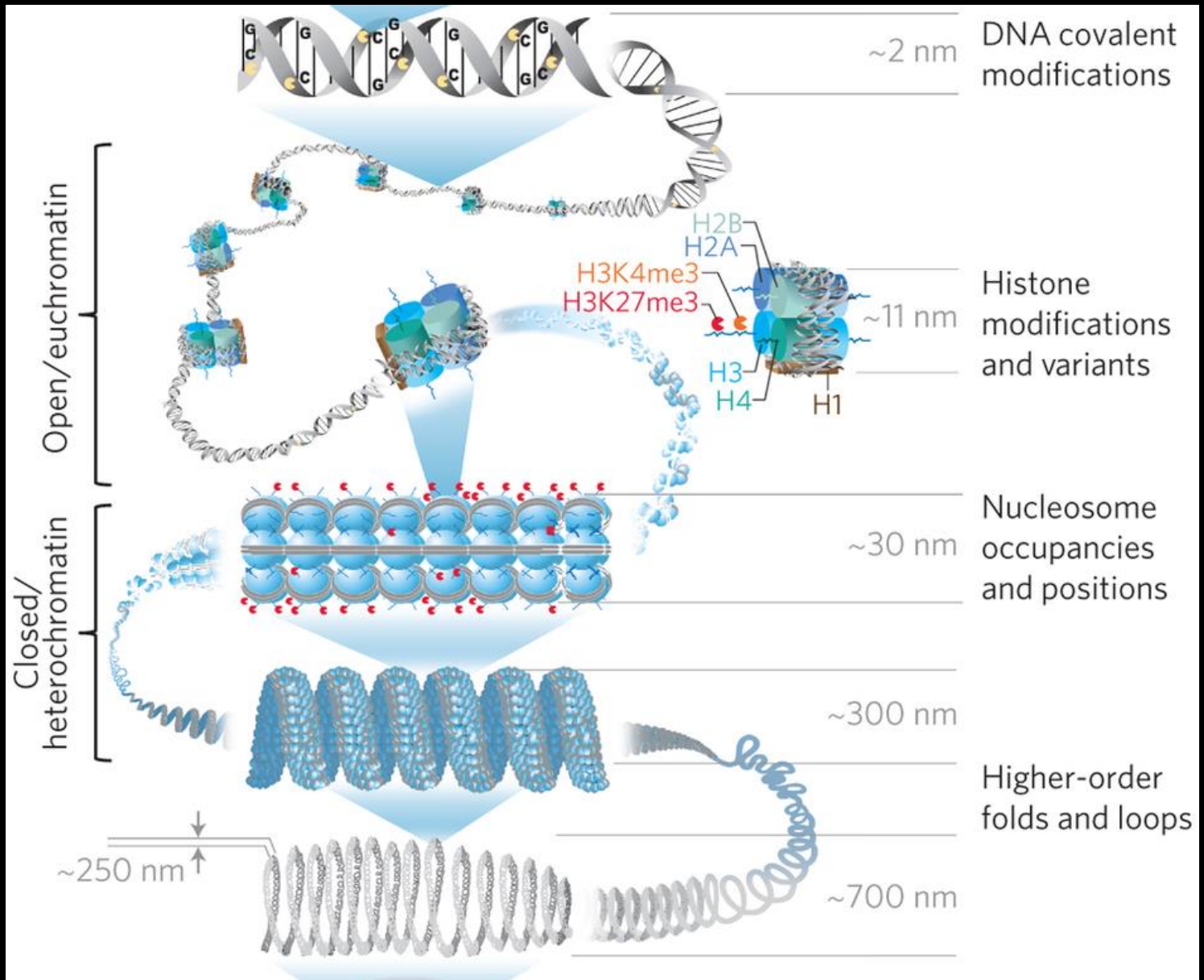


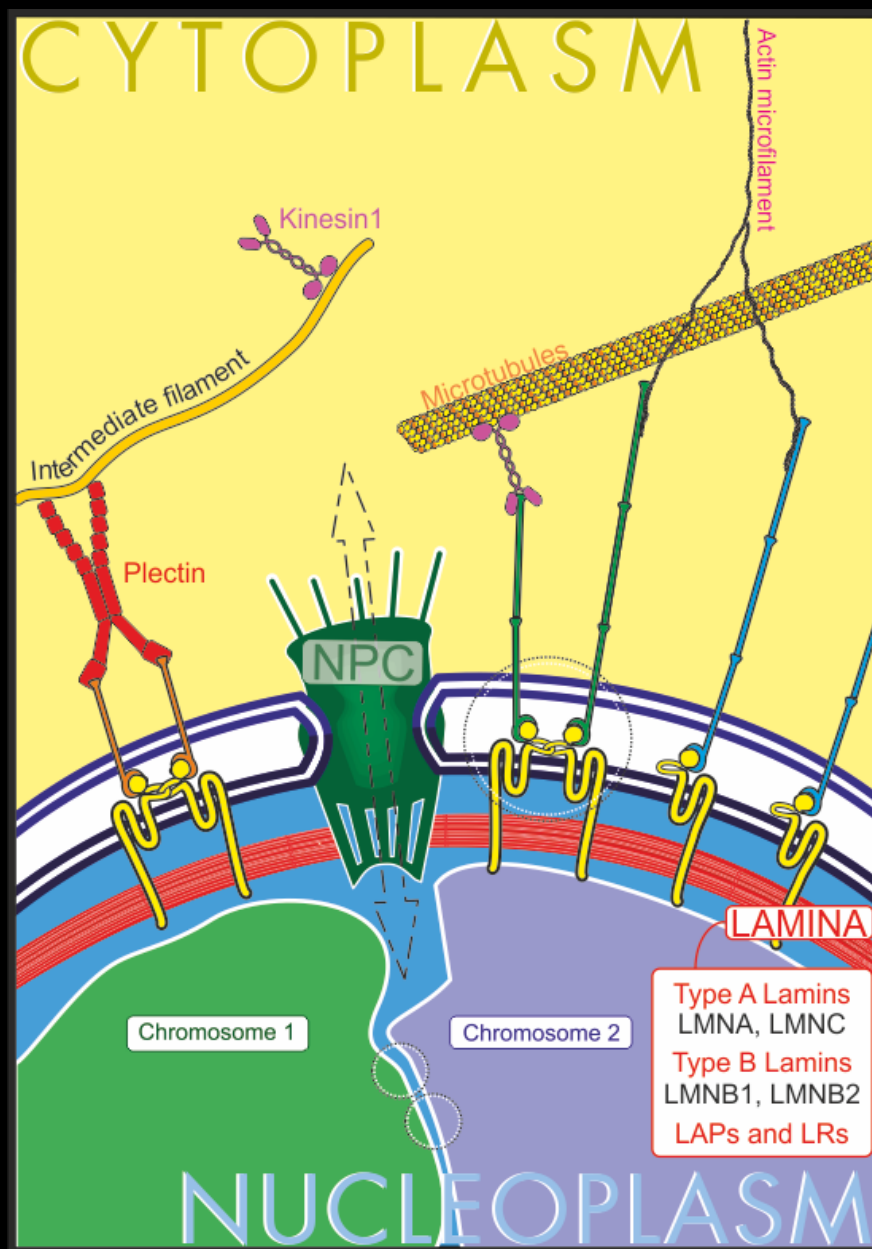
Epigenetic status

- Global DNA demethylation**
- Only active X chromosomes; Global repression of differentiation genes by Polycomb proteins; Promoter hypomethylation**
- X inactivation; Repression of lineage-specific genes by Polycomb proteins; Promoter hypermethylation**
- X inactivation; Derepression of Polycomb silenced lineage genes; Promoter hypermethylation**

Hochedlinger and Plath 2009; DOI:10.1242/dev.020867







The Nucleus

Is the largest compartment of the cell, it houses the genome (chromatin/chromosomes) and is the site of all genomic transcription.

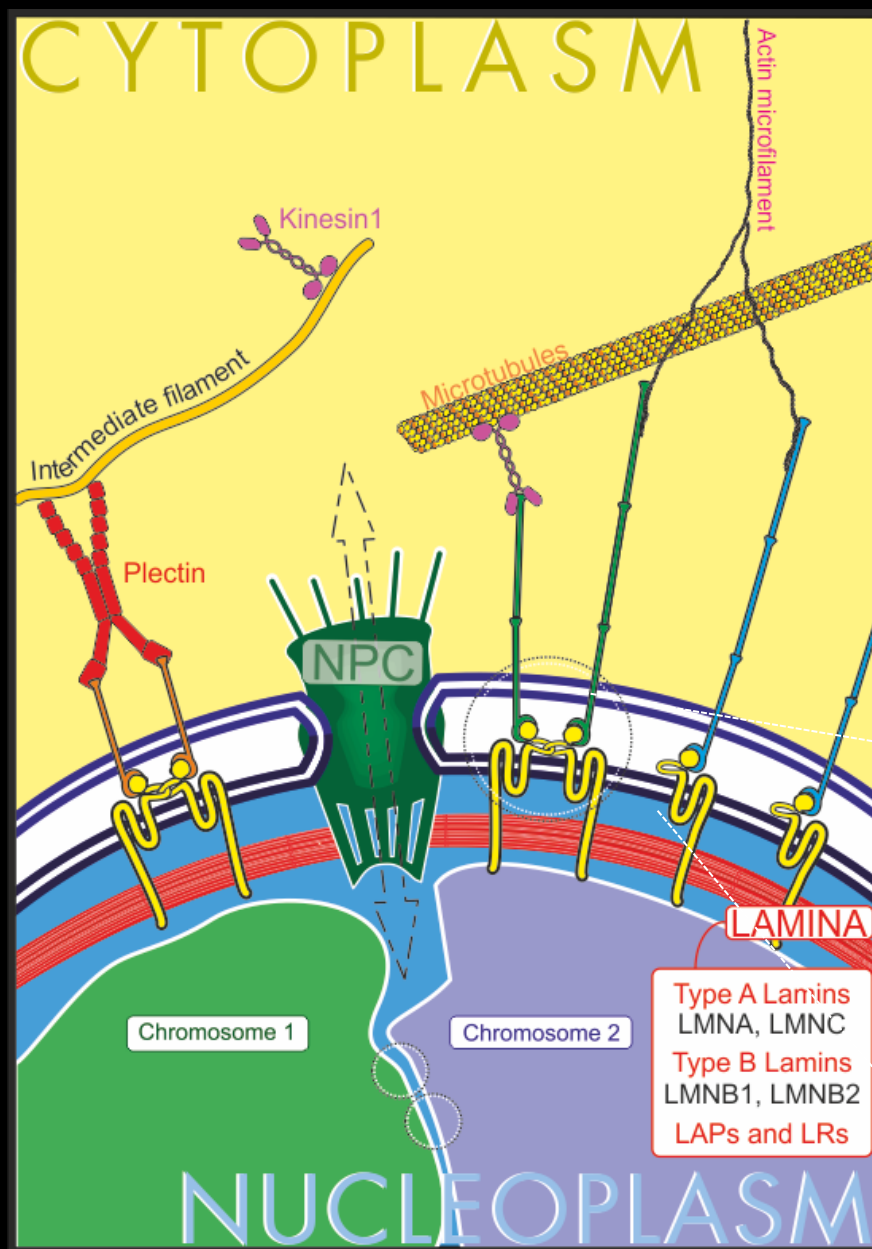
It is surrounded by the nuclear envelope, composed of:

- The nuclear membrane, semi-permeable double lipid bi-layer
- large nuclear pore complexes control trafficking in and out of the nucleus
- the nuclear lamina, the major support structure of the nucleus, composed of intermediate filaments

The LINC complex traverses the nuclear envelope and connects the cytoskeleton to the nuclear lamina.

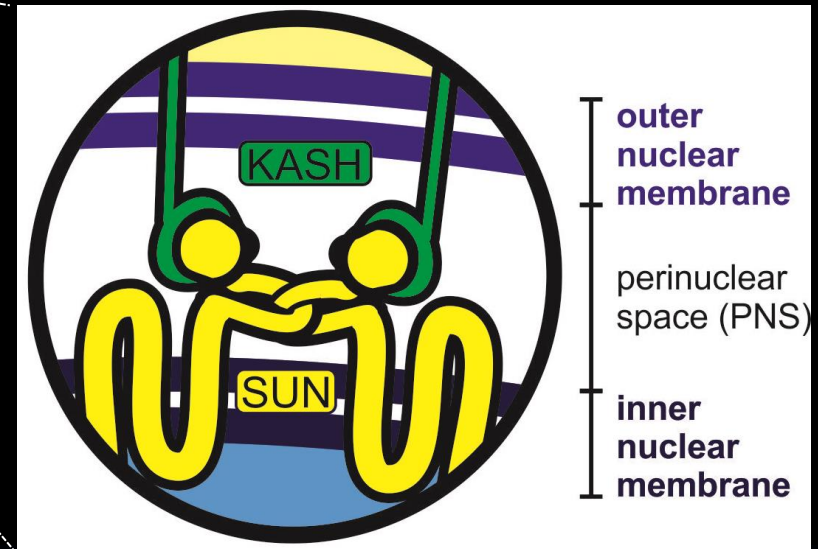
- It involved in cell polarisation and nuclear positioning
- Force transmission through the cell is dependent on an intact LINC complex





The LINC complex

- SUN-domain proteins on the inner nuclear membrane that connect the lamina
- KASH-domain proteins (Nesprins) on the outer membrane that bind to elements of the cytoskeleton



Cytoskeleton

The major support structure in the cell with three principal components

- **Actin** rapidly polymerises into **microfilaments** that are cross-linked to form **filaments** that are mobilised in the cell to apply forces to neighbouring cells or its environment, or to induce cell migration/movement

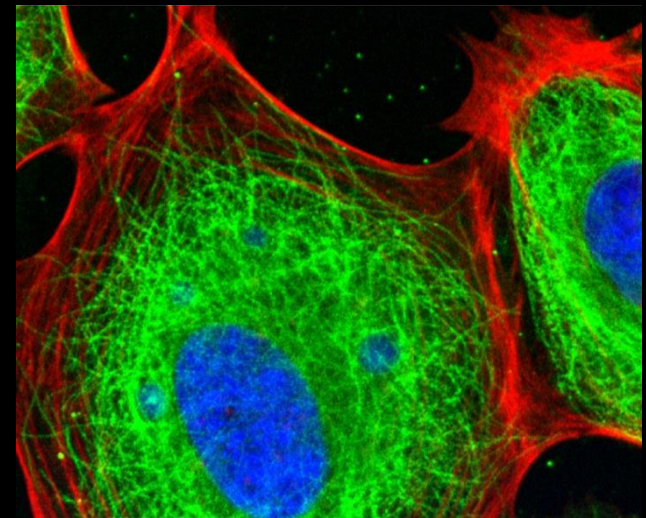
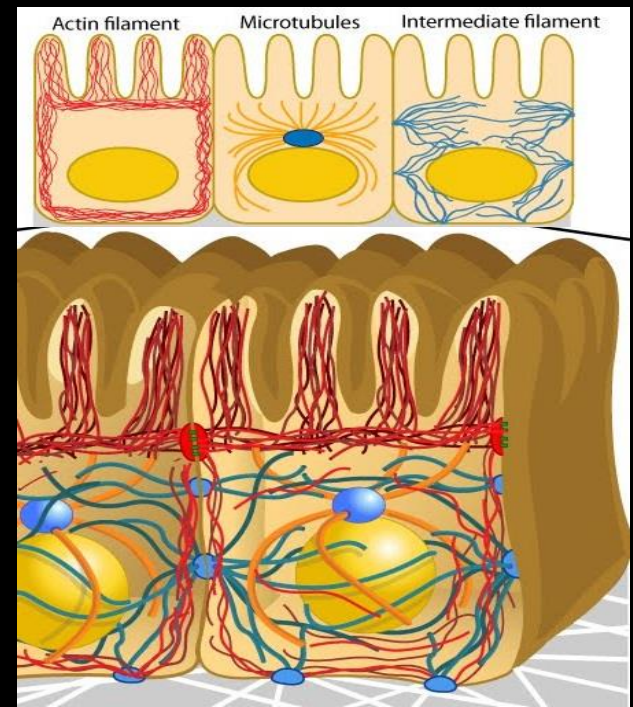
http://www.youtube.com/watch?v=y6Ro_gxl_HE

- **Microtubules** are polymers of α and β -tubulin that are involved in organelle localisation, cell division, motility (cilia, flagella) and mechanosensation (?)

<http://www.youtube.com/watch?v=2-L-Ts6fsks>

- **Intermediate filaments** form fibrous structures that are flexible but resistant to fracture or extension and connect to neighbouring cells through desmosomes.

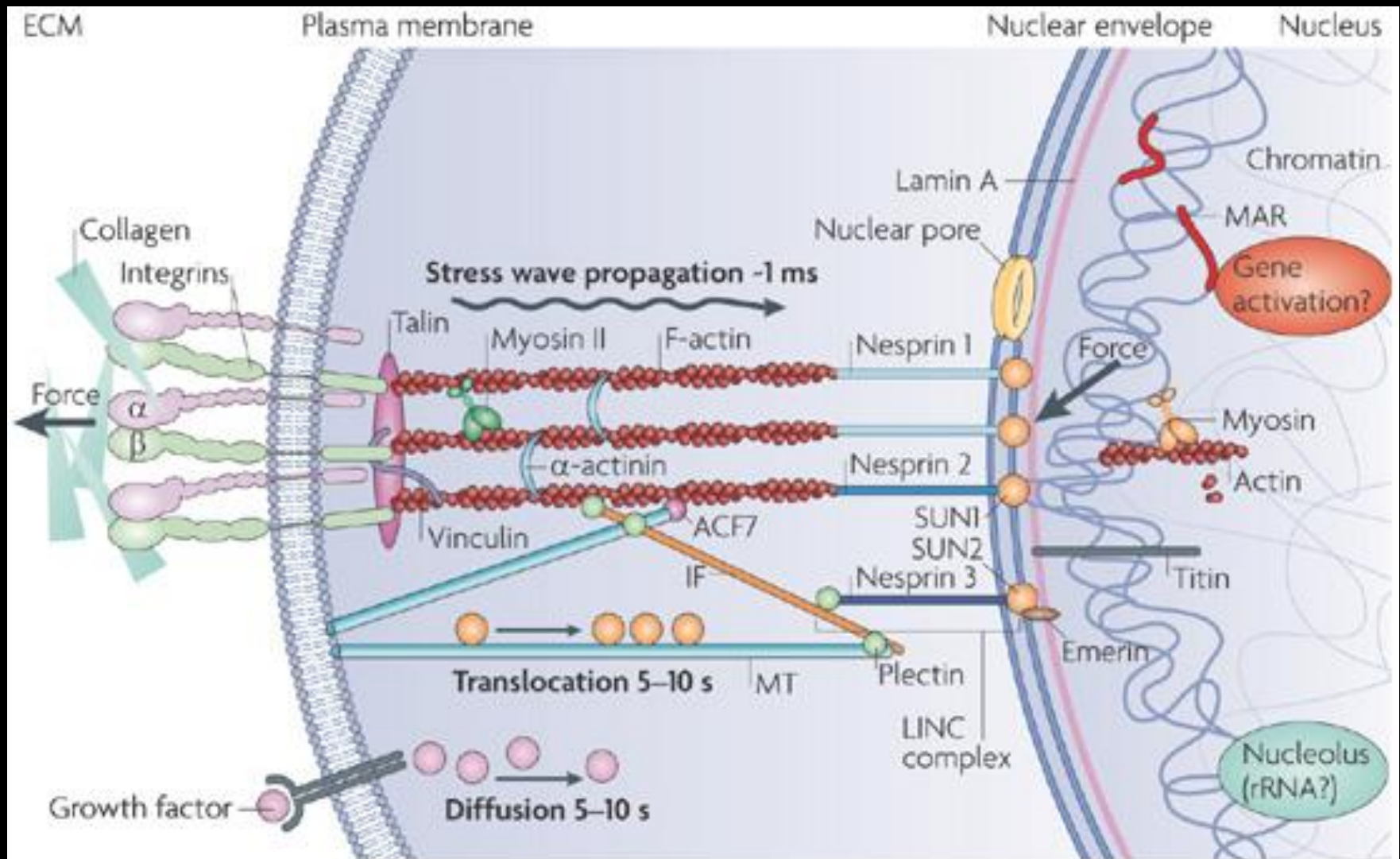
<http://www.youtube.com/watch?v=FoDniO676Dw>



Actin Tubulin Nuclei (DNA)



Mechanotransduction on the nucleus



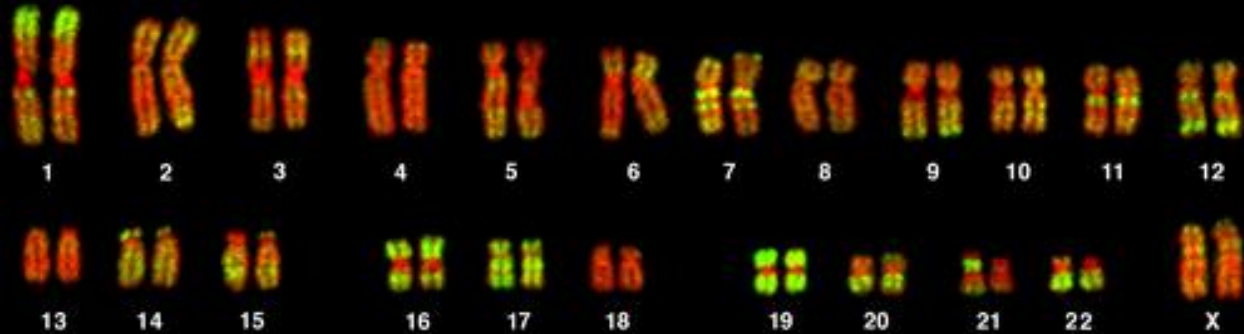
Nature Reviews | Molecular Cell Biology

Wang et al. 2009 doi:10.1038/nrm2594

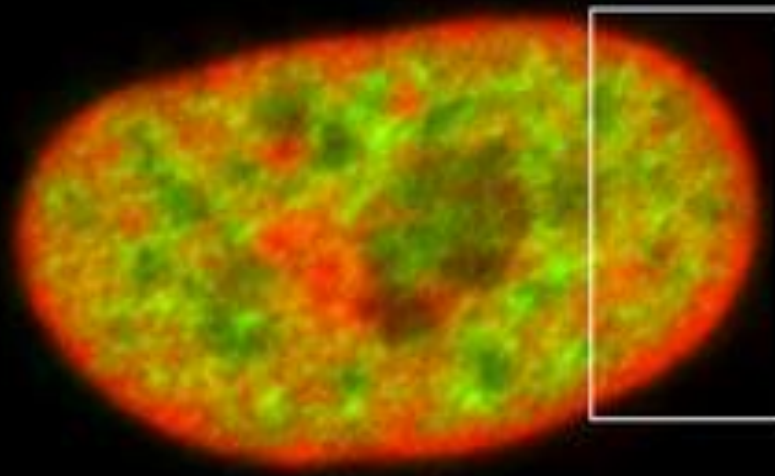
EMBRYO PHYSICS | 26 FEB 2014



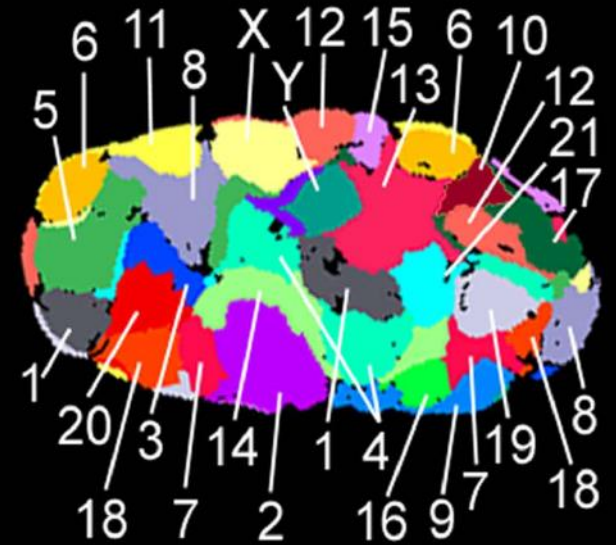
Chromosomes decondense after cell division



metaphase chromosomes from a female lymphocyte (Karyotype: 46, XX)



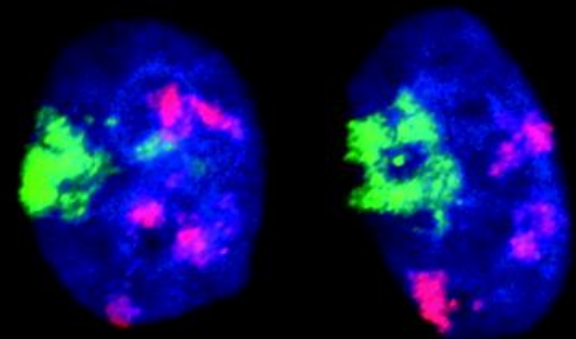
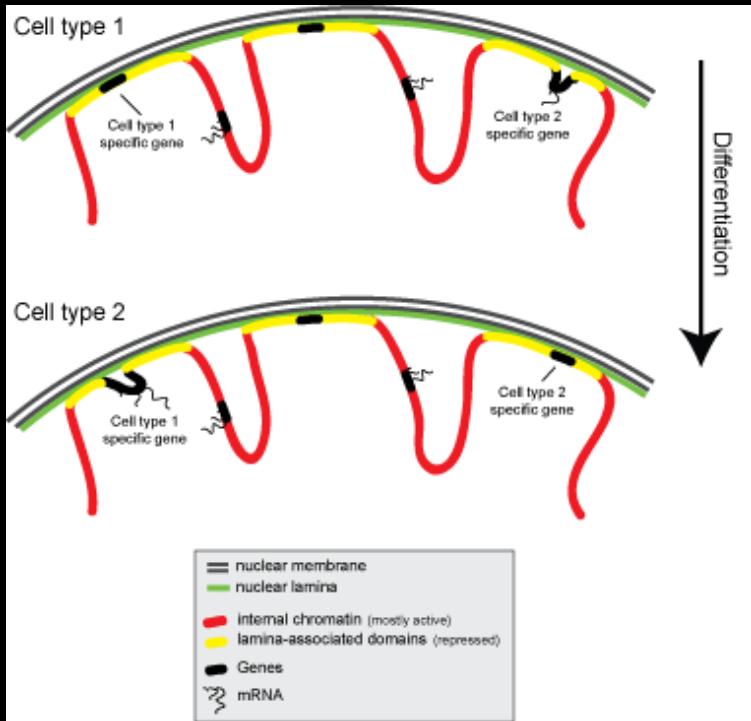
decondensed chromosomes from a female lymphocyte



Bolzer et al. PLoS Biol. 3(5):e157. doi:10.1371/journal.pbio.0030157



- The organisation of the genome within the (interphase) nucleus is non-random
- Chromosomes occupy specific *Territories* within the nucleoplasm
- Radial 3D arrangement (ie. localisation towards the nuclear interior vs. periphery) has been associated with gene activity and can be clonally inherited.
- Proximity to the nuclear periphery is associated with developmental gene silencing.
- Changes in intranuclear 3D localisation of specific chromosome territories (and their adjacent “chromosome neighbourhood”) have been correlated with different cell states
- Mechanism is still unclear, but both active (molecular motors) and passive movements involving a cell division have been suggested

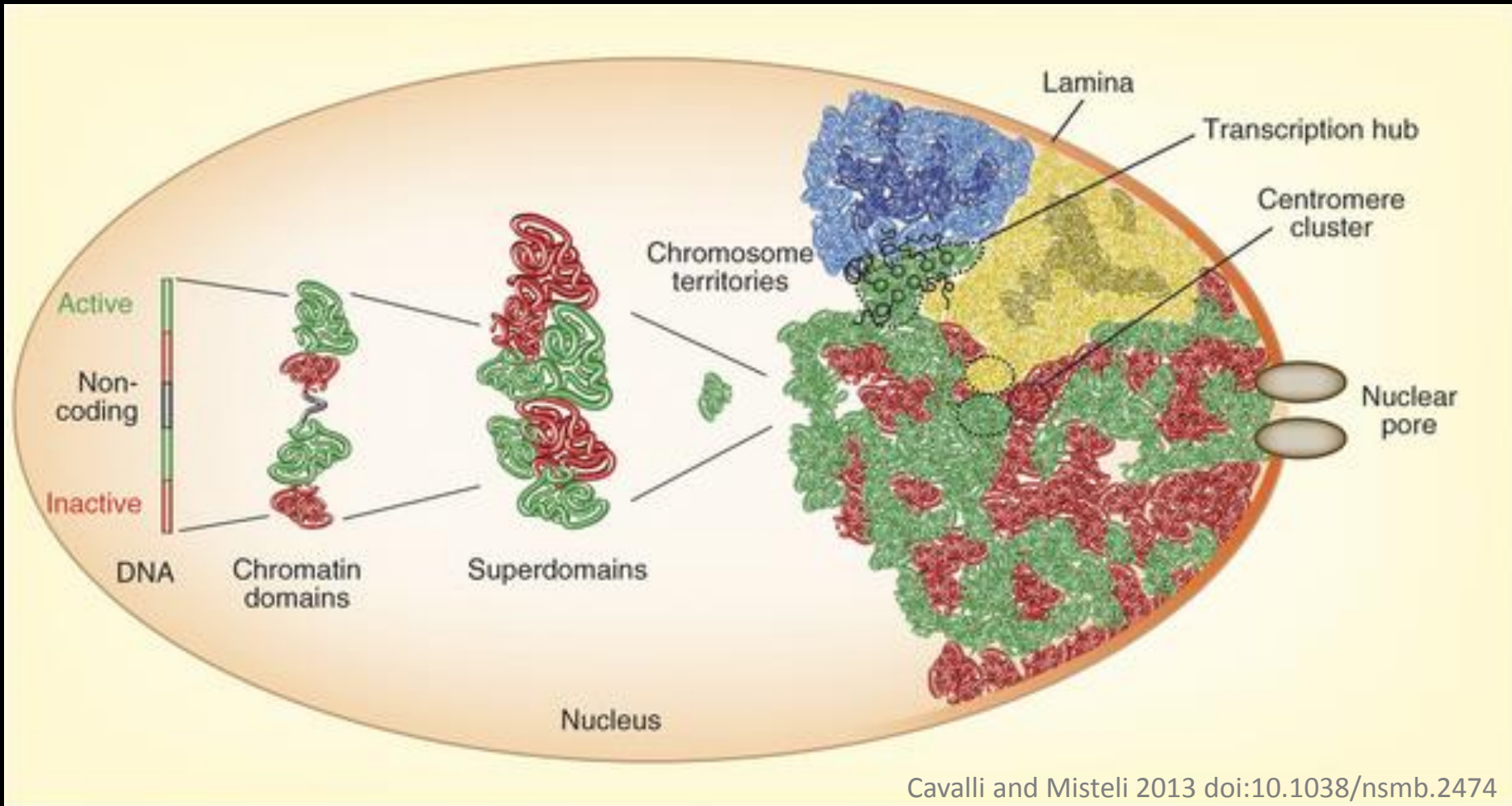


Chr10 Chr7

Walter et al 2003 doi: 10.1083/jcb.200211103

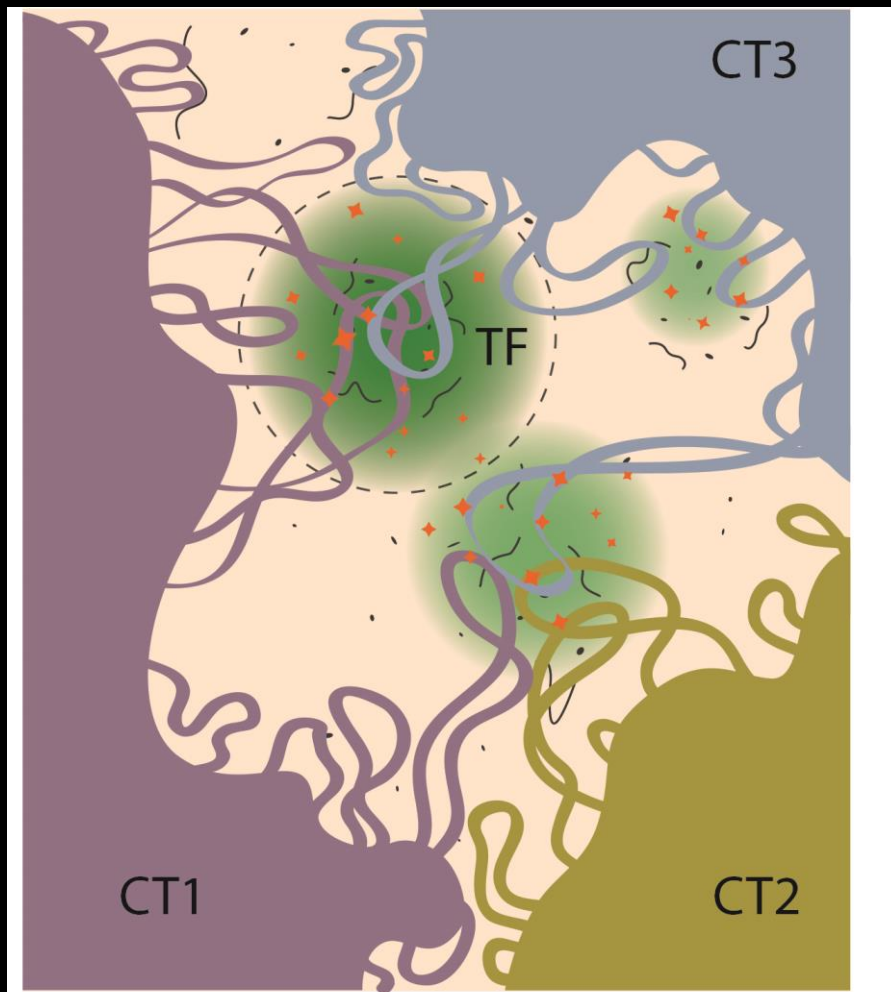


Chromatin architecture: Chromosome domains and neighbourhoods facilitate the silencing or transcription of co-regulated genes that are genomically distant...



Cavalli and Misteli 2013 doi:10.1038/nsmb.2474





TF - transcription factory

◆ RNA polymerase

— mRNA transcript

CT - chromosome territory

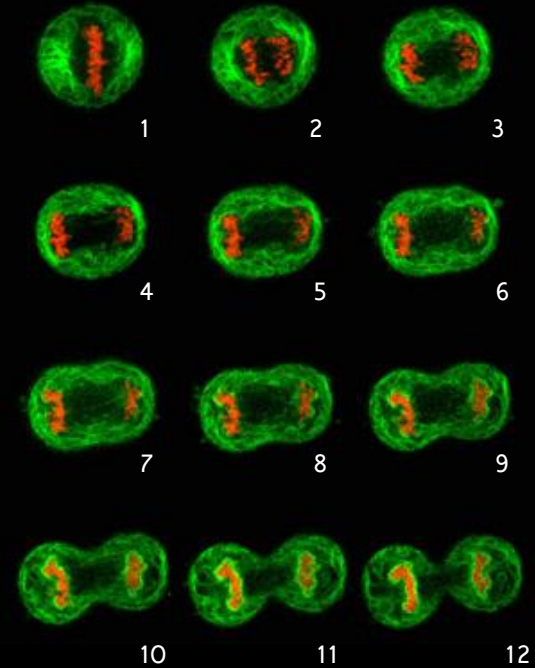
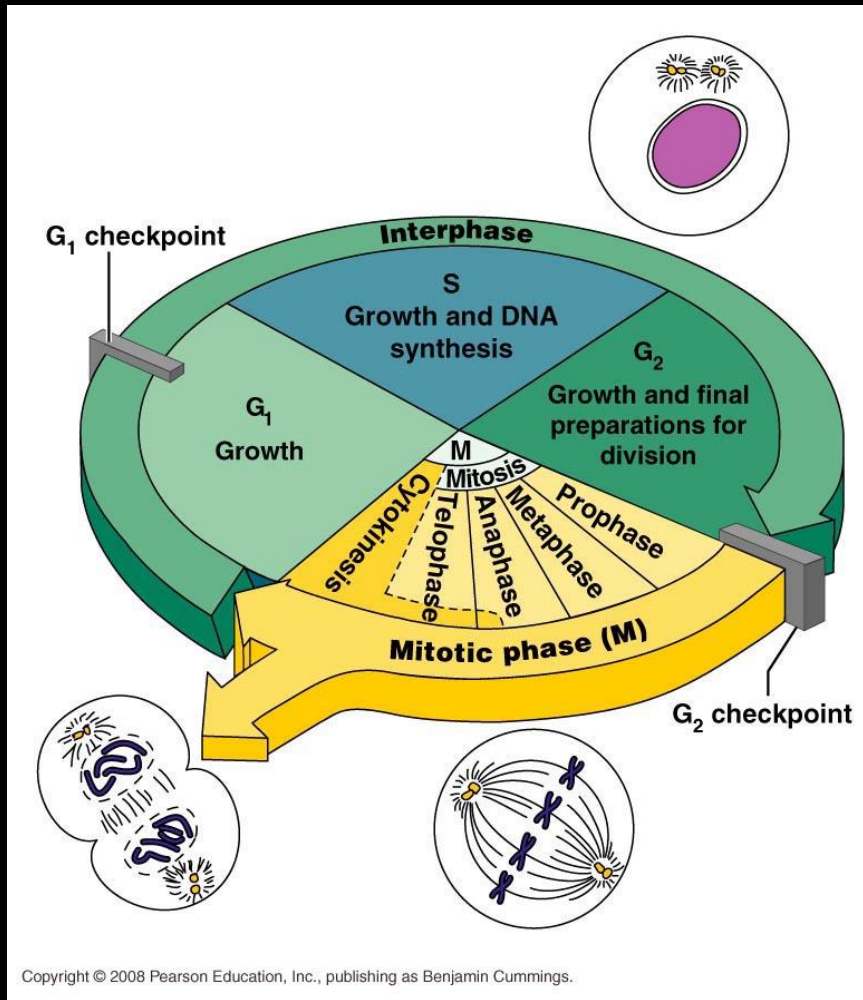
— chromatin loop

<http://www.mechanobio.info/topics/synthesis/go-0006351>

...placing them in a region (subnuclear domain) that is rich in the factors necessary to promote their intended regulation for that cell type/developmental state.



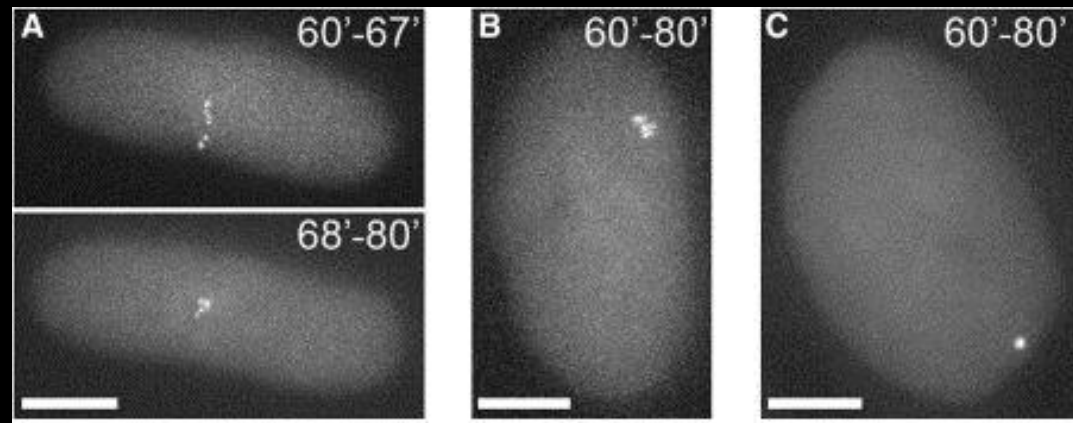
The Cell Cycle and Cell Division



Spindle microtubules (Tubulin), Nuclei (DNA)

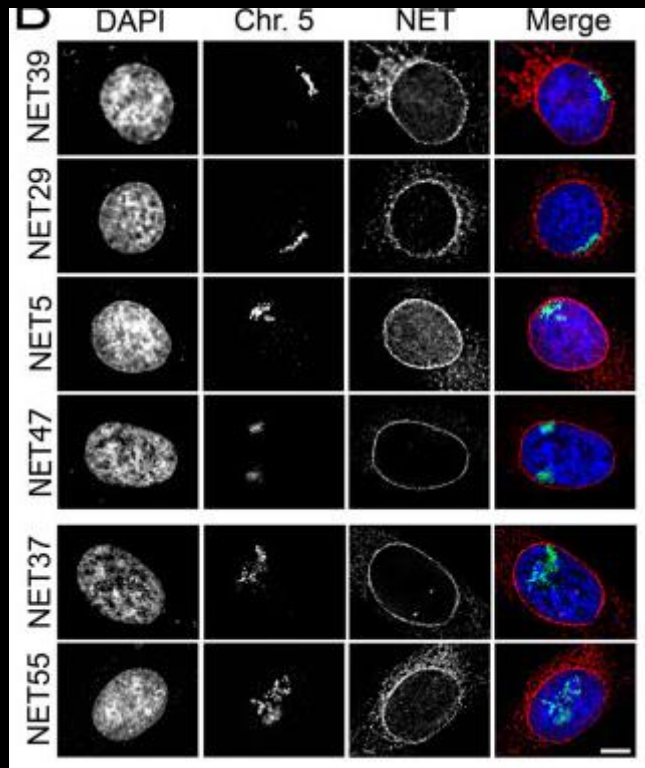


“Migration of an interphase chromosome site from the nuclear periphery to the interior 1–2 hr after targeting at transcriptional activator to this site. Spot redistribution is perturbed by specific actin or nuclear myosin I mutants.”



Chuang et al 2006 <http://dx.doi.org/10.1016/j.cub.2006.03.059>

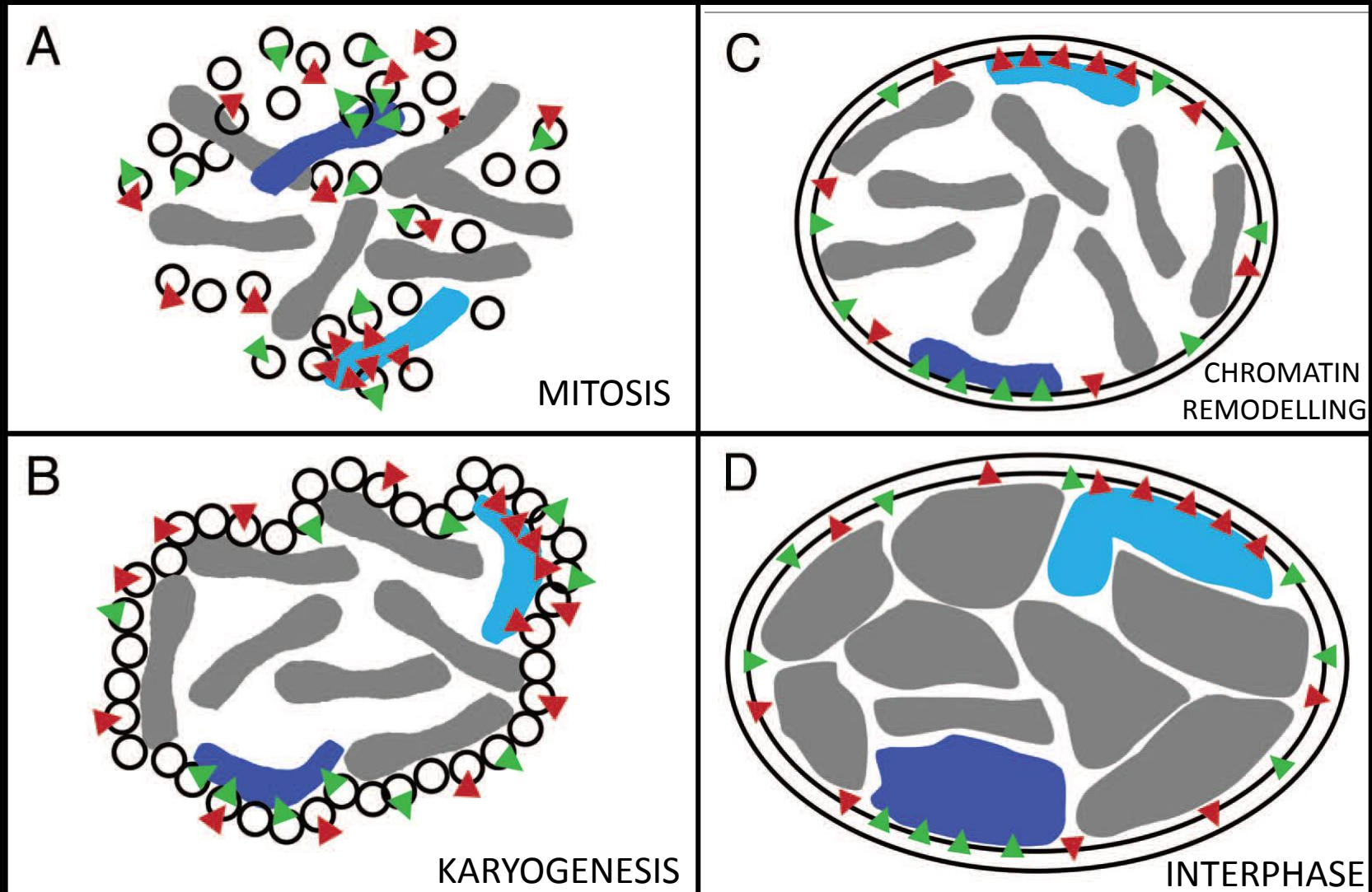
Zuleger et al 2013 doi:10.1186/gb-2013-14-2-r14



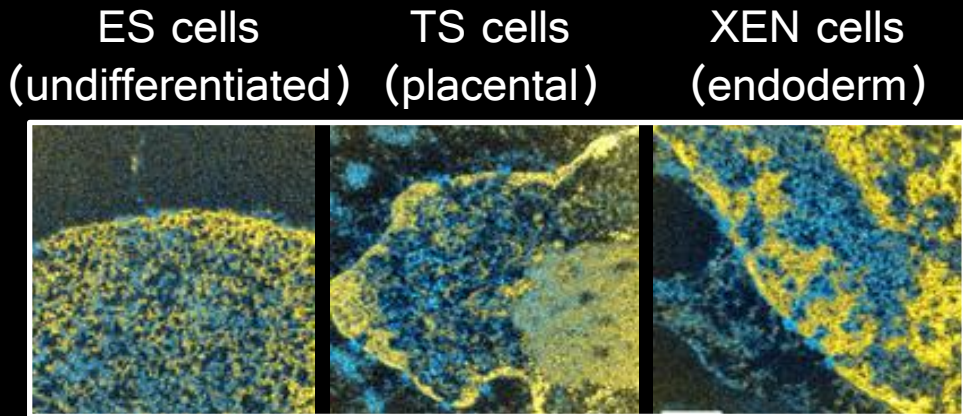
“The discovery of nuclear envelope transmembrane (NET) proteins that can modulate chromosome position and have restricted patterns of expression may enable dissection of the functional relevance of tissue-specific patterns of radial chromosome positioning.”



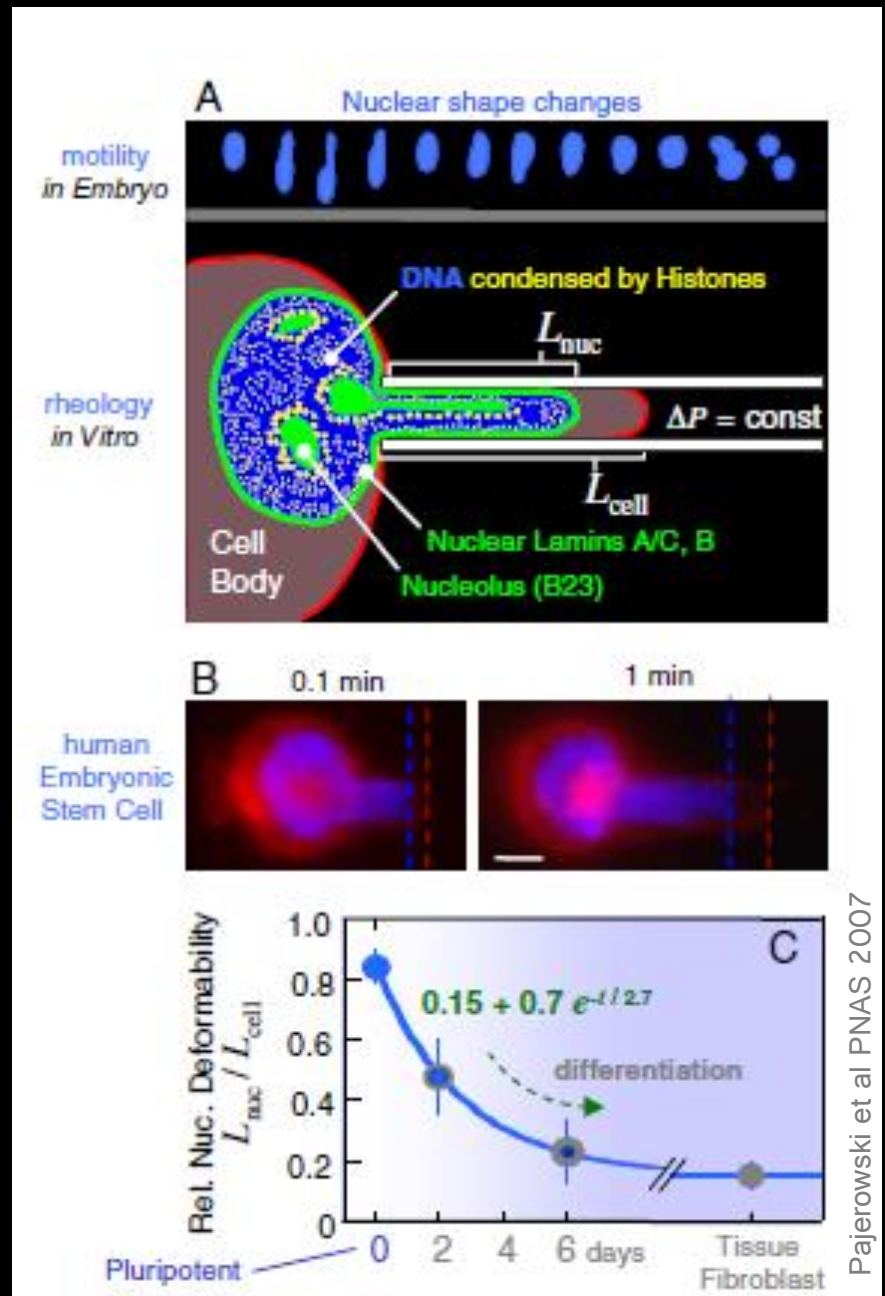
Speculative affinity mechanism for establishment of spatial chromosome organizational patterns.



- As cells differentiate, nuclei also become more stiff (less deformable)
- As cells differentiate, the chromatin becomes more complex, and accumulations of highly complex DNA begin to form around the nuclear periphery



Ahmed et al . (2010 PLoS ONE 5(5): e10531

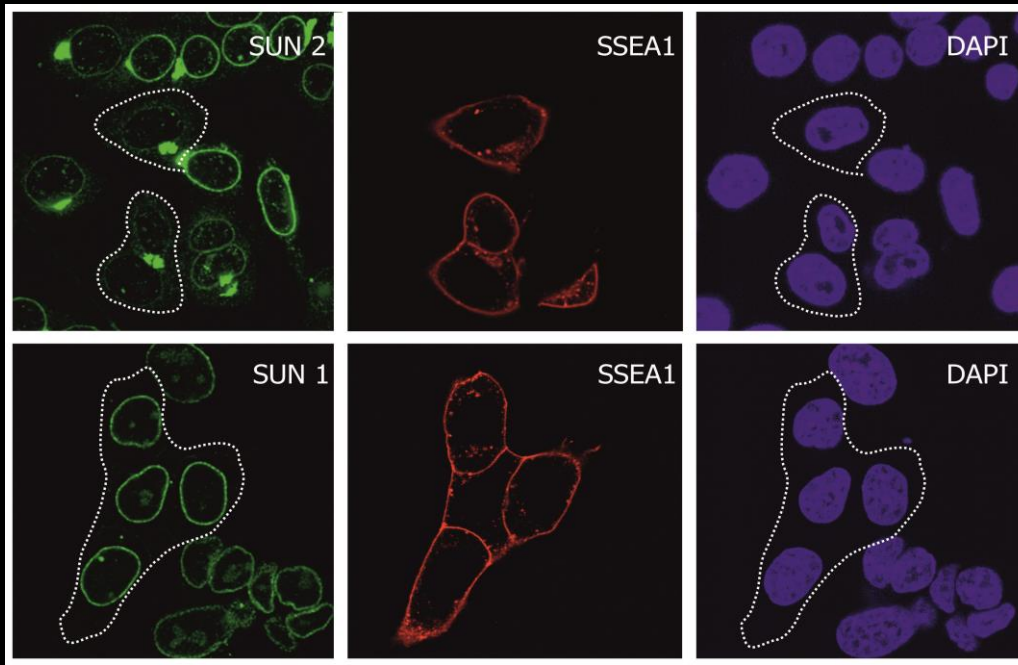


Pajerowski et al PNAS 2007

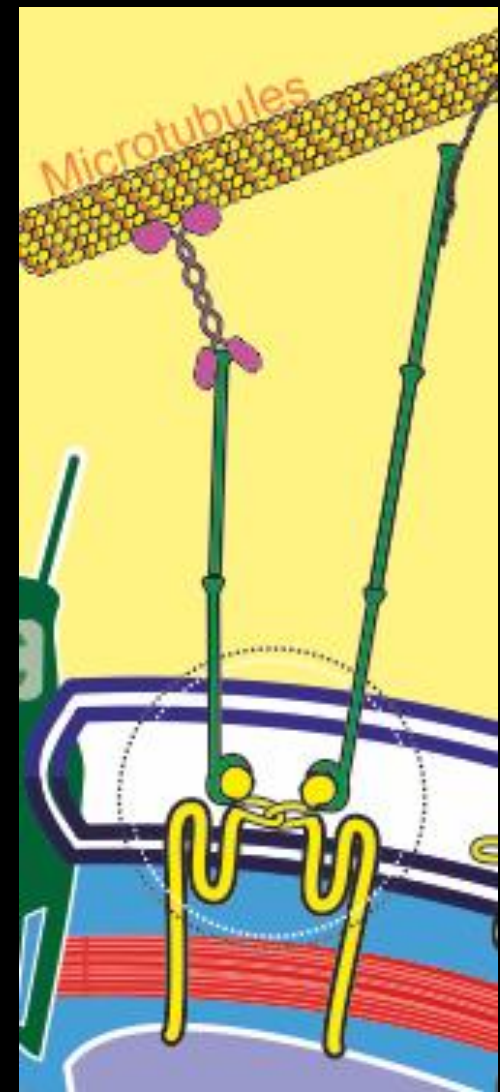


LINC complex & Differentiation

- SUN2 is part of the LINC, connecting the lamina (the major nuclear support structure) and the nuclear membrane and the cytoskeleton through the Nesprin proteins.
- Implicated in nuclear positioning and DNA anchorage
- SUN2 is only expressed after ES cells differentiate



(pluripotent ES cells)



Summary

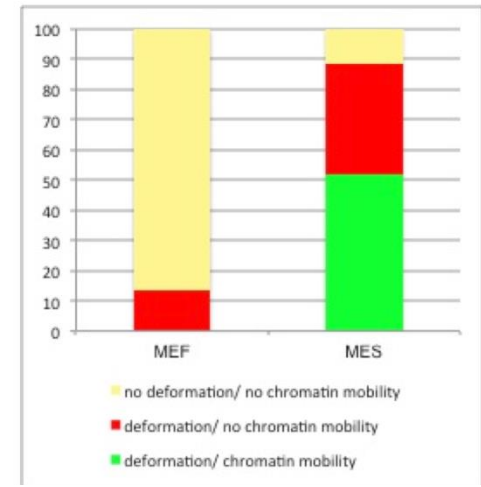
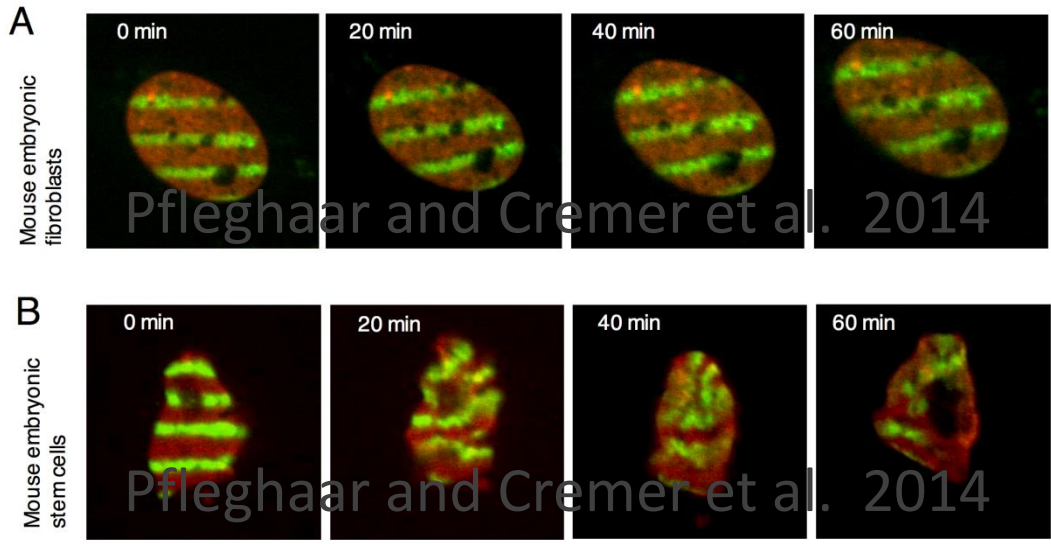
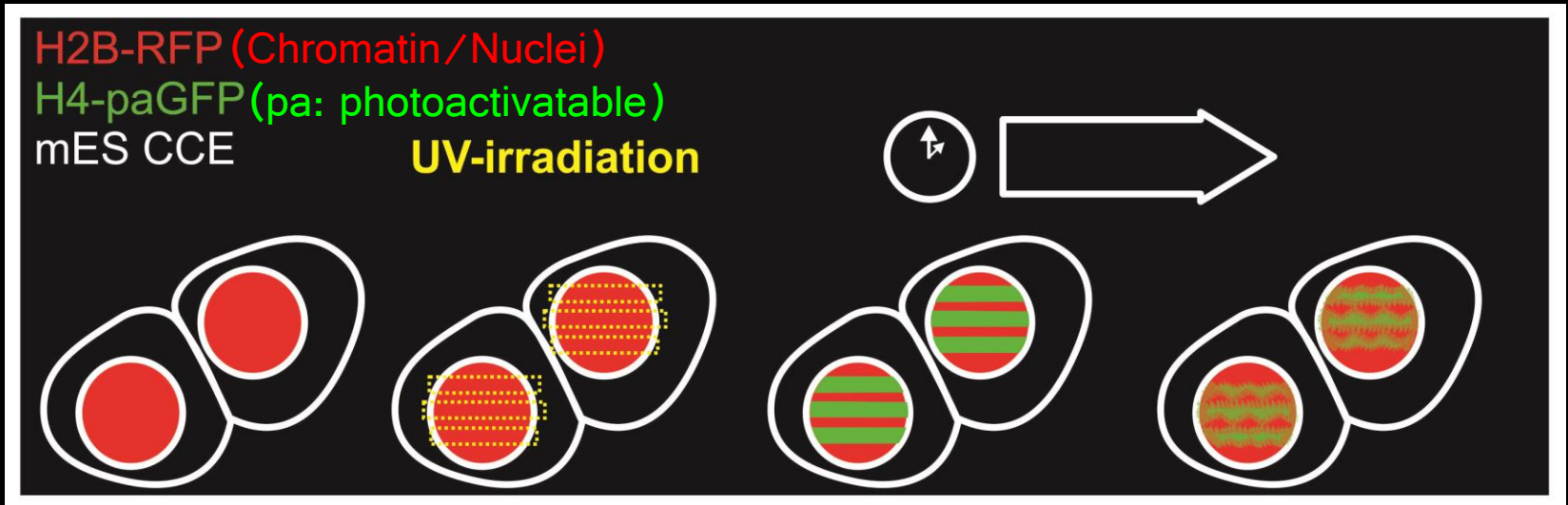
- Chromosomes occupy specific *Territories* within the nucleoplasm and 3D arrangements of neighbouring chromatin can be inherited between from mother cell to daughters.
- At differentiation the composition of the lamina and nuclear envelopes change as A-type Lamins, type-II Suns are up-regulated.
- Nuclei of embryonic stem cells stiffen when they differentiate. This is in part explained by changes to the lamina and envelope (Laminopathic cell data)

What about the chromatin, does it contribute structure to the nucleus?

(we'll revisit this in a moment...)

Q: How does nuclear deformability relate to chromatin organisation.

ES cells have more fluid chromatin (chromosomes)



But what is an ES cell?

And what does it mean, (epi)genomically to be a stem cell?

Is there more than one state to pluripotency?

Cell

The Transcriptional and Epigenomic Foundations of Ground State Pluripotency

Hendrik Marks,¹ Tüzer Kalkan,² Roberta Menafra,¹ Sergey Denissov,¹ Kenneth Jones,² Helmut Hofemeister,⁵ Jennifer Nichols,^{2,3} Andrea Kranz,⁵ A. Francis Stewart,⁵ Austin Smith,^{2,4,*} and Hendrik G. Stunnenberg^{1,*}

¹Department of Molecular Biology, Faculty of Science, Nijmegen Centre for Molecular Life Sciences (NCMLS), Radboud University, PO Box 9101, 6500 HB Nijmegen, The Netherlands

²Wellcome Trust Centre for Stem Cell Research and Stem Cell Institute

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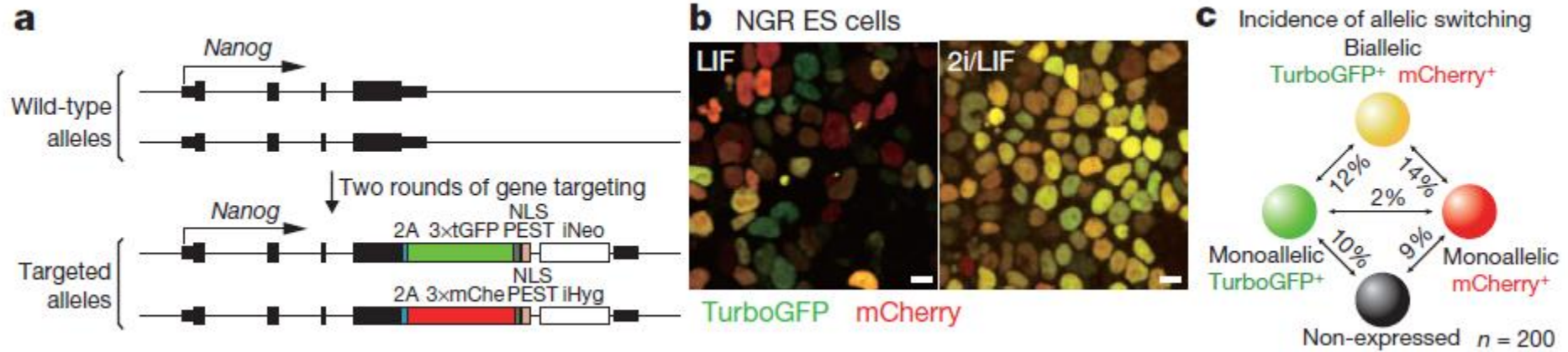
DOI 10.1016/j.cell.2012.03.026

SUMMARY

Mouse embryonic stem (ES) cells grown in serum exhibit greater heterogeneity in morphology and expression of pluripotency factors than ES cells cultured in defined medium with inhibitors of two kinases (Mek and GSK3), a condition known as “2i” postulated to establish a naive ground state. We show that the transcriptome and epigenome profiles of serum- and 2i-grown ES cells are distinct. 2i-treated cells exhibit lower expression of lineage-affiliated genes, reduced prevalence at promoters of the repressive histone modification H3K27me3, and fewer bivalent domains, which are thought to mark genes poised for either up- or downregulation. Nonetheless, serum- and 2i-grown ES cells have similar differentiation potential. Precocious transcription of developmental genes in 2i is restrained by RNA polymerase II promoter-proximal pausing. These findings suggest that transcriptional potentiation and a permissive chromatin context characterize the ground state and that exit from it may not require a metastable intermediate or multilineage priming.



Ground-State Pluripotency is mediated by bi-allelic *nanog* expression



Miyanari and Torres-Padilla 2012 doi:10.1038/nature10807

2060

Biophysical Journal Volume 103 November 2012 2060–2070

Chromatin Decondensation and Nuclear Softening Accompany *Nanog* Downregulation in Embryonic Stem Cells

Kevin J. Chalut,^{†‡Δ*} Markus Höpfler,^{†Δ} Franziska Lautenschläger,[†] Lars Boyde,[†] Chii Jou Chan,[†] Andrew Ekpenyong,[†] Alfonso Martinez-Arias,[§] and Jochen Guck^{†¶}

[†]Cavendish Laboratory, Department of Physics, [‡]Wellcome Trust/Medical Research Council Centre for Stem Cell Research, and [§]Department of Genetics, University of Cambridge, Cambridge, United Kingdom; and [¶]Biotechnology Center, Technische Universität Dresden, Dresden, Germany



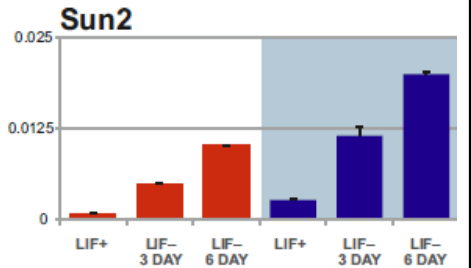
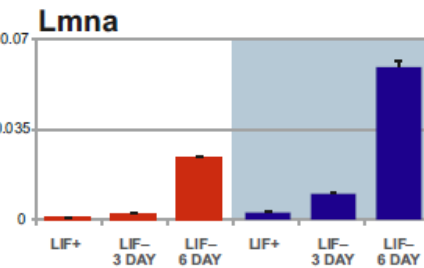
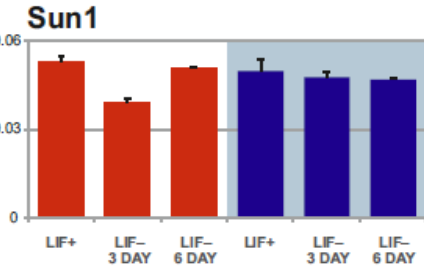
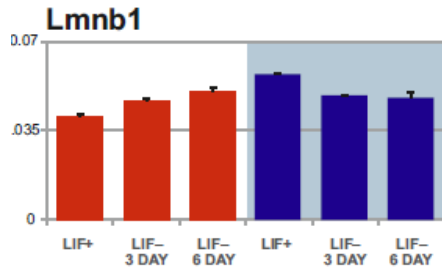
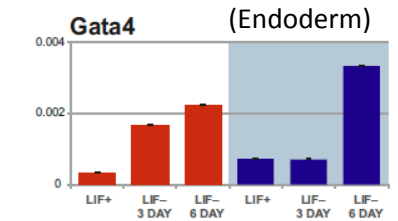
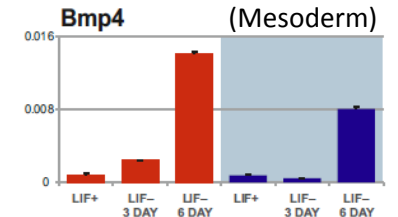
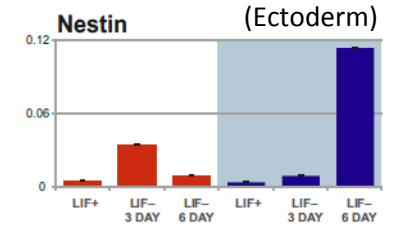
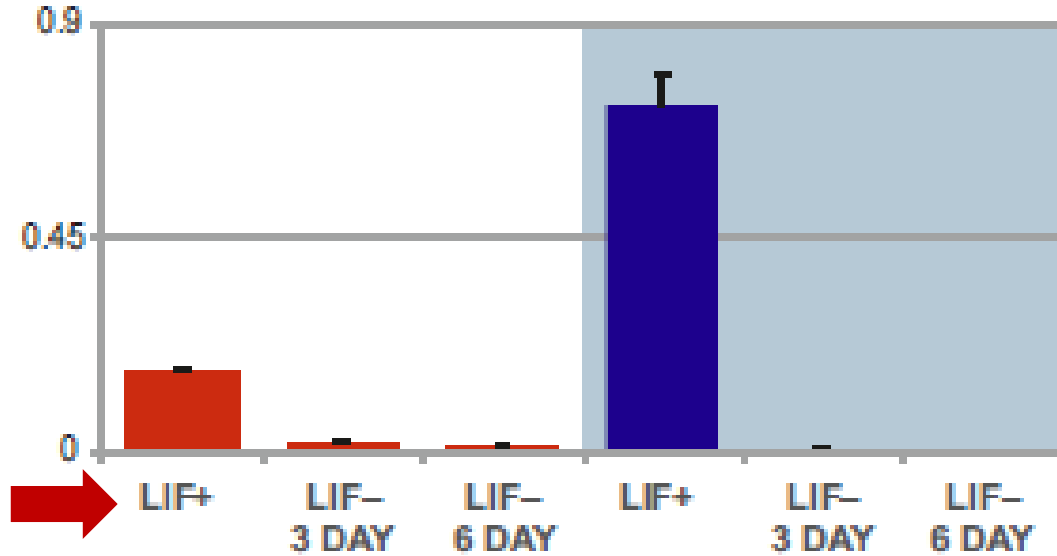


STD ESC CULTURE



2i: EMB GROUND STATE

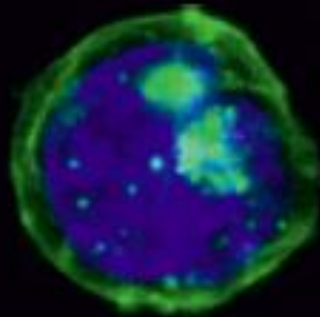
Nanog (Main Pluripotency Regulator)



- Pluripotency in ES cells has two phases that are regulated by the level of Nanog expression.
- Monoallelic (and stochastic)-Nanog cells are primed for differentiation, quite heterogeneous, but still capable of self-renewal. These cells have soft nuclei and highly motile chromatin, despite an increased amount of negative epigenetic marks.
- Bi-allelic Nanog cells are in the pluripotent ground state. In essence, the state defined as not being anything else (yet). Here they essentially self-renew, and have quite stiff nuclei.
- It has been suggested that the stiffness is contributed through specific repression of lineage-specification programmes.

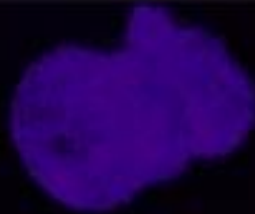


1. Cell selection



mESCs

Hoechst
 α SSEA1



MEFs

Hoechst

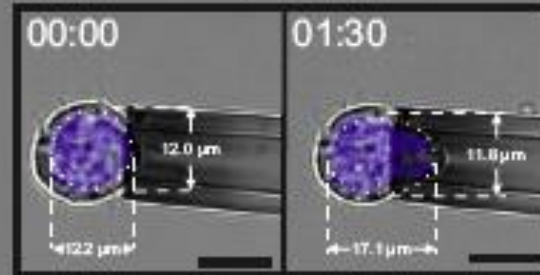


NSCs

Hoechst

2. Aspiration and imaging

1) Aspect Ratio (AR = x/y)



$AR_{00:00} = 1.02$

$AR_{01:30} = 1.45$

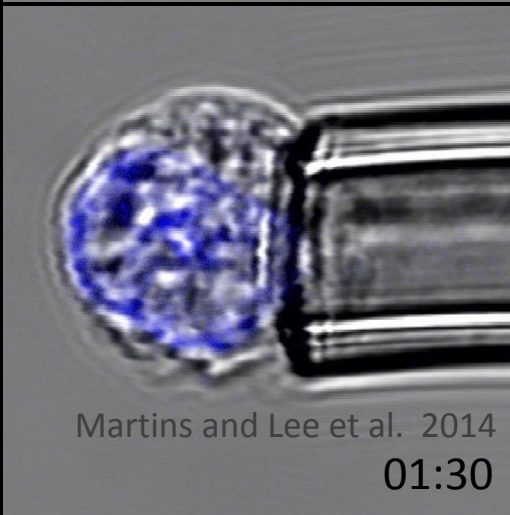
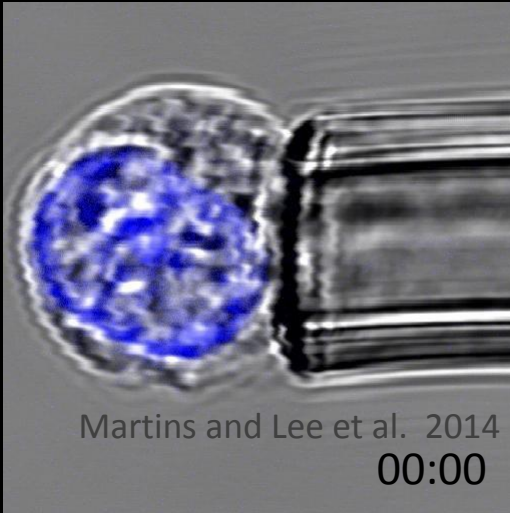
2) calculate aspiration pressure (AP) required to initially deform the nucleus



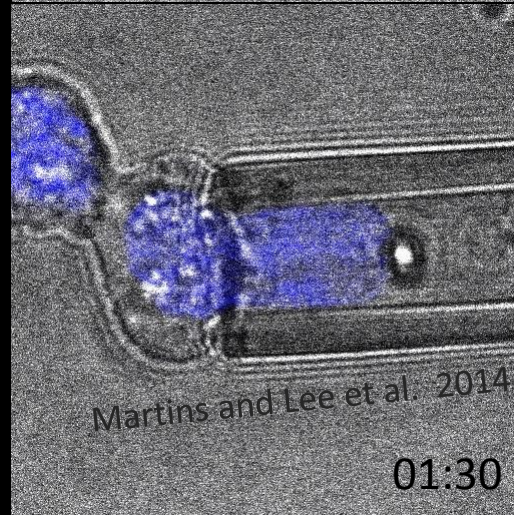
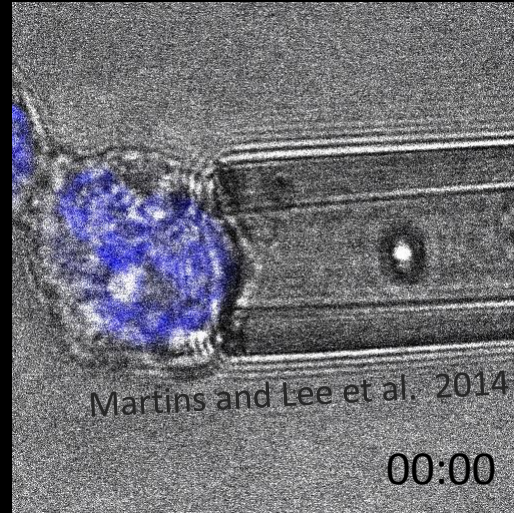
$$AP = (t_2 - t_1) \times 0.245 \text{ kPa/s} \\ = 27.5 \text{ kPa}$$



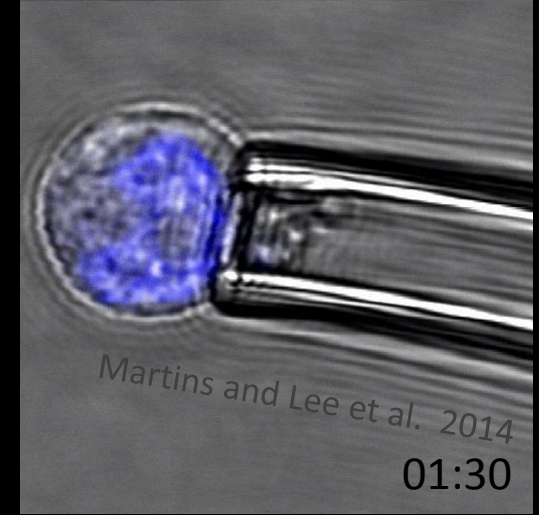
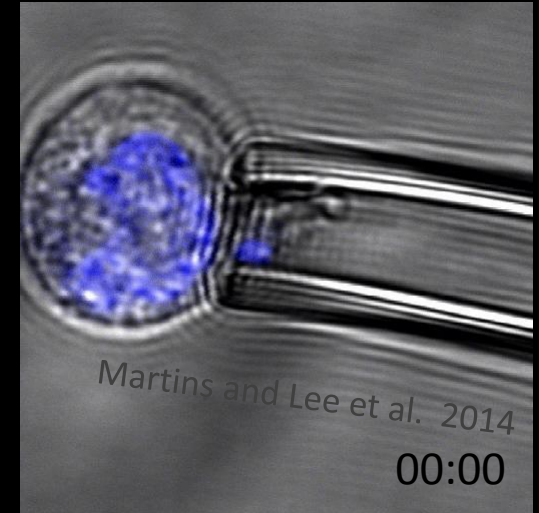
2i-Pluripotent ES cell
medium containing LIF
Bi-allelic Nanog

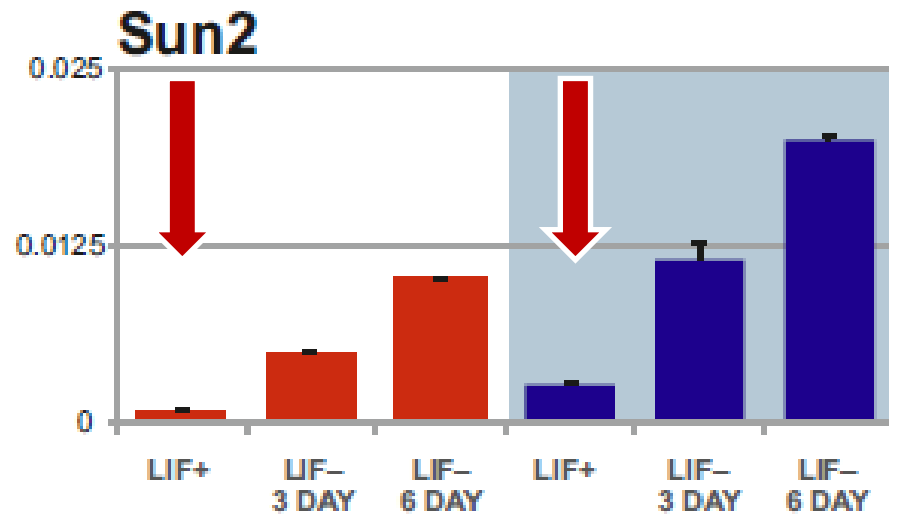
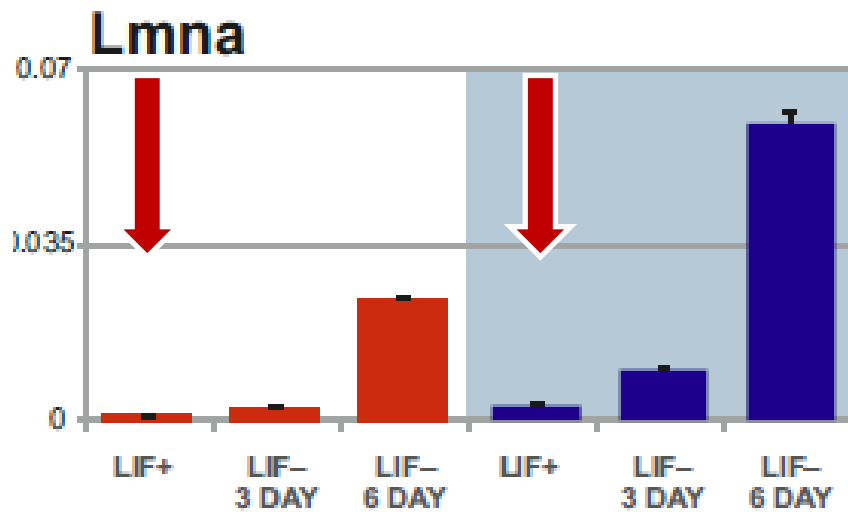


Standard ES cell medium
containing serum and LIF
Monoallelic+stochastic



Differentiation conditions
no LIF, 6 Days
Nanog downregulated



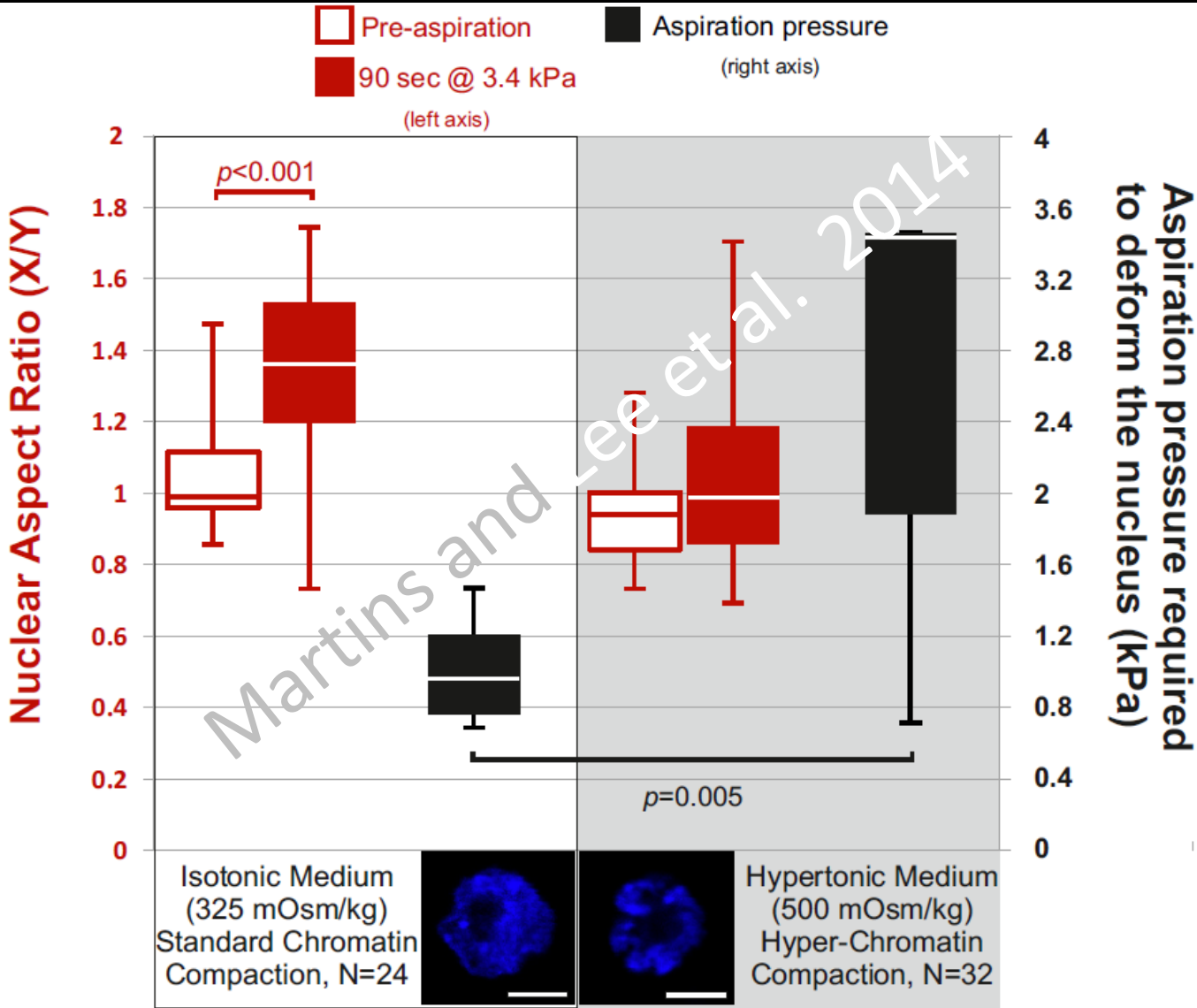


STD ESC CULTURE

 2i: EMB GROUND STATE

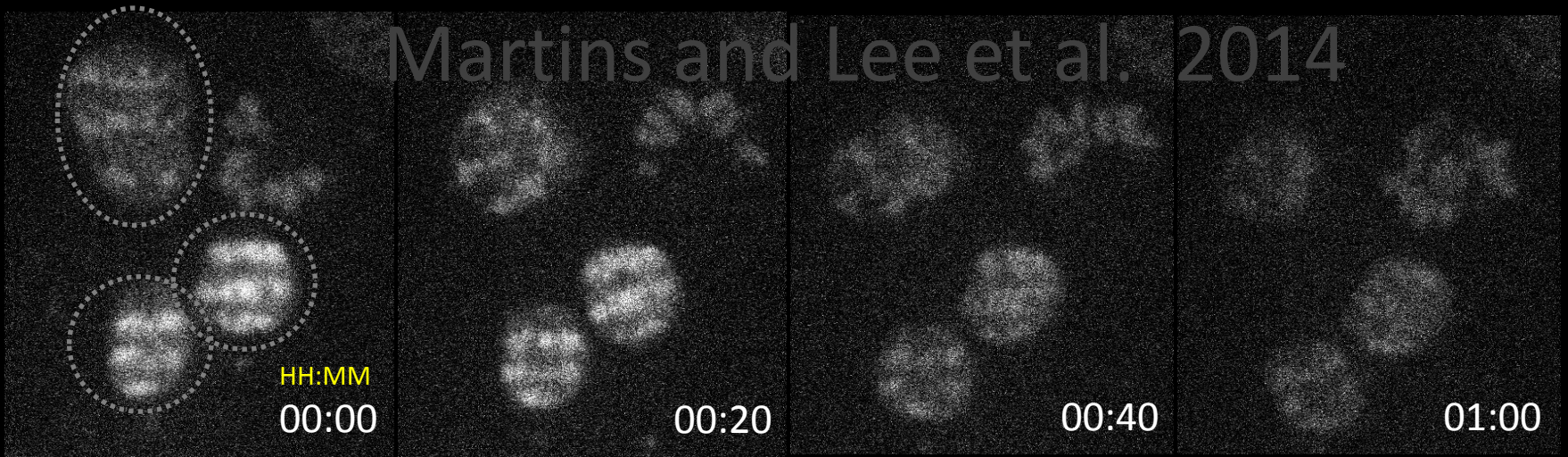


Q: Can chromatin
condensation alone
explain some of the
nuclear stiffness?

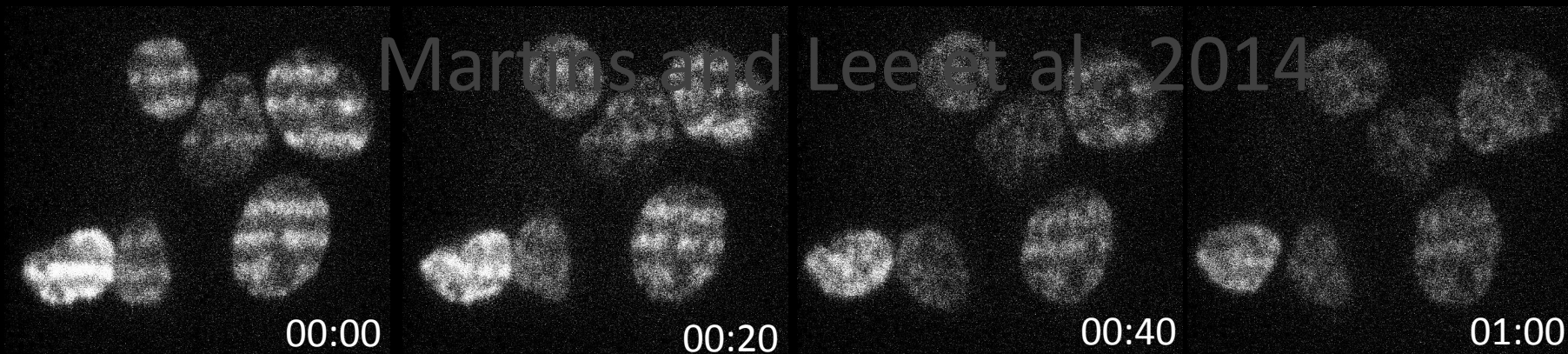


Q: ES cells grown in 2i/LIF which promotes bi-allelic Nanog conditions are stiffer, but does that translate to non-motile chromatin, as is evident in differentiating/ed cells?

21.11.2013 Stack 28



21.11.2013 Stack 40



- Chromatin mobility is much lower in stiffer nuclei- no gross rearrangements visible during interphase.
- Changes in the nuclear envelope composition when ES cells differentiate (Lamin A and Sun 2) provides more putative anchoring points to stabilise chromosomal architecture
- Changes to the overall level of chromatin condensation; with highly condensed chromatin being able to significantly stiffen ES cell nuclei.

So why is it important to maintain a fluid chromatin architecture while in a state primed for differentiation?



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