CHAIRE ÉPIGÉNÉTIQUE ET MÉMOIRE CELLULAIRE

Année 2013-2014 : "Reprogrammations développementales, induites et pathologiques "

<u>Cours III</u>

Reprogrammation experimentale – les cellules pluripotentes induites

24 mars 2014

Seminaire: Dr Claire Rougeulle à 17h30 "Inactivation du chromosome X, pluripotence et reprogrammation, de la souris à l'homme".



E. Heard, March 24th 2014

All cells have the capacity to form a whole organism – through differential gene expression



⇒ Differentiated cells are NOT irreversibly committed to their fate but can be REPROGRAMMED and/or REPURPOSED

Perspectives brought by ES cells and SCNT



Egg

cell

Donor a

n

ES

Therapeutic Cloning?



Derivation of **human ES cells** by Jamie Thomson in 1998 and more recent work by Noggle et al 2011 showing that it is, in principle, possible to reprogram human somatic cells up to the blastocyst stage at least.

=> "**Therapeutic Cloning**" became a real possibility: Patient-specific pluripotent ESCs by somatic cell nuclear transfer into human eggs, which could then be differentiated to the cell type that was defective in the patient *For organ transplant replacement, skin grafts, degenerative diseases (eg Parkinson's), diabetes, spinal cord repair, leukemia...)*

Serious ethical issues in humans!!!

Ideally, one would like to derive human ESCs from patient's own <u>somatic</u> cells? Or to change one cell type into another cell type from the same patient?









In this novel, the Hailsham school children are clones, created to be "donors" that provide vital organs for "normals" through a series of "donations" that eventually lead to the donor's death



E G E

NCE

E. Heard, March 24th 2014

Experiments in the 1970s and 80s demonstrated the reprogramming capacity of one cell another by cell fusion:	over
• EC cells were fused with thymus cells and then injected into mice – forming teratocarcinomas => pluripotency must be dominant (Miller and Ruddle 1976)	
• Subsequently mouse ESCs shown to impose pluripotency onto hybrids generated using various somatic cell fusion partners eg T cells (Tada et al., 2001), splenocytes (Matveeva et al., 1998), bone marrow (Terada et al., 2002), & neural progenitors (Ying et al., 2002).	
Previous isolation of BJ fibroblasts 2n Virus 2n Chemical cell-fusion Co-select	

COLLÈGE DE FRANCE

E. Heard, March 10th 2014

Vakatsuji⁺

. 19

ion i

The inactive X chromosome in a somatic cell can be reactivated after cloning and after fusion with ES cells



Ta

Di 19

19 19

19

19

19

19

Reactivation of the Xi following fusion of Female thymus cells with male ES cells (*Xist* down regulation; shift to synchronous replication timing)

Tada et al, 2002 "Nuclear reprogramming of somatic cells by in vitro hybridization with ES cells", Curr.Biol.

"ES Cell Dominance"
Experiments in the 1970s and 80s demonstrated the reprogramming capacity of one cell over another by cell fusion:
• EC cells were fused with thymus cells and then injected into mice – forming teratocarcinomas => pluripotency must be dominant (Miller and Ruddle 1976)
• Subsequently mouse ESCs found to impose pluripotency onto hybrids generated using various somatic cell fusion partners eg T cells (Tada et al., 2001), splenocytes (Matveeva et al., 1998), bone marrow (Terada et al., 2002), & neural progenitors (Ying et al., 2002).
ES cells: ⇒ can rewire the gene regulatory network of any cell type? ⇒ can reprogram the inactive X chromosome, as well as pluripotency factors (<i>Oct4</i>) ⇒ must contain trans-acting factors capable of reprogramming somatic cell nuclei?
hat could these be?
Overexpression of <i>Nanog</i> , a pluripotency transcription factor, substantially enhanced fusion- used nuclear reprogramming (Silva et al, 2006)
Heterokaryon-based reprogramming of human B lymphocytes for pluripotency requires Oct4 Pereira et al. 2008).



Vakatsuji⁺

in and a

an

1

Pluripotency TF network and Embryonic Stem (ES) cells

Pluripotency transcriptional network: driven by core transcription factors (TFs) Oct4, Nanog, Sox2, Klf4 – maintain pluripotency and self-renewal.

This network activates genes required for ES cell survival and proliferation & represses target genes that are only activate during differentiation.



E. Heard, March 17th 2014

Transcription factor-mediated repurposing of somatic cells into pluripotent stem cells?

"When I saw the embryo, I suddenly realized there was such a small difference between it and my daughters. I thought, we can't keep destroying embryos for our research. There must be another way." s. Yamanaka - while looking through a microscope at a friend's fertility clinic.



nduced roliferate tem cell

ewal

tors

n at ters



Based on what was known about ES cells, Shinya Yamanaka reasoned that <u>forcing</u> the expression of ES cell-specific genes, particularly transcription factors, in somatic cells might induce them to take on a more embryonic character...

E. Heard, March 24th 2014







The Yamanaka Strategy



Drs. Shinya Yamanaka and Kazutoshi Takahashi

Fbx15 locus



A

Fex15 locus

• Select 24 "ES-cell" gene candidates

Transcription factors involved in self-renewal (Oct3/4, Sox2, and Nanog), or known to be upregulated specifically in ES cells (Mitsui et al., 2003), and some that are associated with transformation but that have also been implicated in the maintenance of ES cell pluripotency (*c-Myc*, *Eras*, and *Klf4*).

Retroviral infection

(Morita et al, 2000) of 24 cDNAs, together, alone, or in combinations

• Mouse embryonic ("young") or adult tail tip fibroblasts ("old")

Harboring selectable marker (b-geo) under control of a promoter active only in ES cells (Fbx15). Fbx15 activation results in Neomycin (G418) resistance (cell survival – colony) and β -galactosidase activity



- Transduction of 24 genes ->NeoR and b-gal+ colonies
- Individual genes alone -> no colonies
- Narrowed down to *Oct4*, *Sox2*, *Klf4 and* c-myc the "magic" cocktail" (OSKM) (NB not Nanog...



Narrowed down factors from 24 to 10 to 4...



E. Heard, March 24^m 2014



The Yamanaka Strategy



Drs. Shinya Yamanaka and Kazutoshi Takahashi



A active Fbx15 locus Bgeo Bgeo Bgeo Bgeo Bgeo G418 selection & G418 selec



Narrowed down factors from 24 to 10 to 4



• Select 24 "ES-cell" gene candidates

Transcription factors involved in self-renewal (*Oct3/4*, *Sox2*, and *Nanog*), or known to be upregulated specifically in ES cells (Mitsui et al., 2003), and some that are associated with transformation but that have also been implicated in the maintenance of ES cell pluripotency (*c-Myc*, *Eras*, and *Klf4*).

Retroviral infection

(Morita et al, 2000) of 24 cDNAs, together, alone, or in combinations

Mouse embryonic ("young") or adult tail tip fibroblasts ("old")

Harboring selectable marker (b-geo) under control of a promoter active only in ES cells (*Fbx15*). *Fbx15* activation results in Neomycin (G418) resistance (cell survival – colony) and β -galactosidase activity

- Transduction of 24 genes ->NeoR and b-gal+ colonies
- Individual genes alone -> no colonies
- Narrowed down to *Oct4*, *Sox2*, *Klf4 and* c-*myc* the "magic" cocktail" (OSKM) (NB not *Nanog*...

Criteria for "stemness"?

- ES cell morphology
- Self renewal (continuous cell division)
- Expression of endogenous pluripotency factors
- Transcriptomes (general gene expression pattern)
- Pluripotency (differentiation into all three germ layers)
- Teratomas
- Germ line transmission

These first iPS cells did NOT pass all the tests in fact



E. Heard, March 24th 2014

Do these "induced" pluripotent stem cells (iPS) fit the criteria for "stemness"?



ed with EcoRI and

Do these "induced" pluripotent stem cells (iPS) fit the criteria for "stemness"?

patterns?



MEF

with EcoBl and





iPS cells fit many of the criteria of ES cells – BUT:

- Incomplete demethylation of the Oct4 promoter.
- Low level of endogenous Oct4 and Sox2 expression.
- Expression profiles similar but NOT identical to ES cells
- No chimeras, no germ line transmission

No contribution from iPS-derived cells to postnatal animals \Rightarrow Incomplete Reprogramming?

E. Heard, March 24th 2014

Germ-line competent murine iPS cells





Dot3

Sou

ю

Germ-line competent murine iPS cells

Yamanaka group (Okita et al, 2007) produced germ-line competent murine iPS cells

Selection for *Nanog* expression (rather than *Fbx15*) resulted in germ line-competent iPS cells, with increased ES-cell-like gene expression and DNA methylation patterns

- Longer time allowed for reprogramming before selection
- Selection for *Nanog* expression = better readout than Fbx15 for pluripotent state
- Efficiency of iPS clones still low: 0.1%, but clone quality less variable than Fbx15 iPS



Dot3/4

K014

"Out of 121 F1 mice (aged 8-41 weeks) derived from the Nanog-iPS-20D17 cell line, 24 died or were killed because of weakness, wheezing or paralysis. Necropsy of 17 mice identified neck tumours in 13 mice and other tumours in five mice, including two mice with neck tumours....In these tumours, retroviral expression of c-myc, but not Oct3/4, Sox2, or Klf4, is reactivated"



Generation of iPS cells without the Myc oncogene

Yamanaka group (Nakagawa et al, 2008) produced **iPS cells without exogenous** *Myc* to overcome the tumorigenicity problems – less efficient, but clearly safer.





"Our study does not argue that Myc is dispensable for iPS cell generation. We found that MEFs expressed c-Myc from the endogenous gene at $\sim 20\%$ of the levels observed in mouse ES cells. This expression continues in iPS cells. Thus, Oct4, Sox2

Yamanaka, Hochedlinger and others also produced iPS cells without retroviruses to limit the mutagenic effects of retrovrial integration (eg Okita et al "Generation of mouse induced pluripotent stem cells without viral vectors". Science 322, 949–953 (2008) and Nature
 E. I Protocols (2009).

Gener Myc fr

Masato Naka Takashi Aoi¹





ìΕ

CE

How general is the Yamanaka strategy? Can other mammalian cells be reprogrammed?

Yamanaka group (Takahashi et al, 2007) and Thomson group (Yu et al, 2007) produced **iPS** cells from Adult Human Fibroblasts







E. Heard, March 24th 2014

n of

ma

vork

How general is the Yamanaka strategy? Can other mammalian cells be reprogrammed?

Yamanaka group (Takahashi et al, 2007) and Thomson group (Yu et al, 2007) produced **iPS** cells from Adult Human Fibroblasts

Yamanaka : OKSM (OCT4, SOX2, KLF4, c-MYC) • Thomson: OSLN (OCT4, SOX2, LIN28, NANOG) •



- (B) Morphology of human dermal fibroblasts
- (C) Typical image of <u>non-ES</u> cell-like colony.
- (D) Typical image of <u>hES cell-like</u> colony.
- (E) Morphology of established iPS cell line
- (F) Image of iPS cells with high magnification.
- (G) Spontaneously differentiated cells in the

- Human iPS morphology similar to hESC
- Self renewal
- Similar Transcriptomes in iPS cells and hESCs
- ES cell marker expression, OCT4 hypomethylation
- Teratoma induction in SCID mice



pre-let-

Human iPS cells resemble human ES cells – however they are rather different to mouse ES and iPS cells -- SEMINAR, CLAIRE ROUGEULLE

vork

How general is the Yamanaka strategy? Can other mammalian cells be reprogrammed to pluripotency?

Rats, Cats, and Elephants, but Still No Unicorn: Induced Pluripotent Stem Cells from New Species

Li et al, 2009 and Liao et al 2009: Rat primary cells reprogrammed using lentiviral vectors that expressed OSKM.

Generation of Induced Pluripotent Stem Cells from Adult Rhesus Monkey Fibroblasts

Haisong Liu,^{1,2,6} Fangfang Zhu,^{1,2,6} Jun Yong,^{1,2,6} Pengbo Zhang,¹ Pingping Hou,¹ Honggang Li,¹ Wei Jiang,¹ Jun Cai,¹ Meng Liu,^{1,2} Kai Cui,¹ Xiuxia Qu,¹ Tingting Xiang,¹ Danyu Lu,³ Xiaochun Chi,³ Ge Gao,⁴ Weizhi Ji,⁵ Mingxiao Ding,¹ and Hongkui Deng^{1,2,*}



Retrovirus-mediated transduction of OSKM into monkey fibroblasts:

- Monkey iPS morphology similar to hESC
- Self renewal
- Similar Transcriptomes in iPS cells and hESCs
- ES cell marker expression, OCT4 hypomethylation
- Differentiation into 3 germ layers and teratoma induction in SCID mice.



E. Heard, March 24th 2014

· · · · ·

Rats, 0

Alan Trounson ¹California Institu ¹Correspondenc DOI 10.1016/j.st

Two indeper pluripotent s phenotypic t

How general is the Yamanaka strategy? Can other mammalian cells be reprogrammed to pluripotency?

Rats, Cats, and Elephants, but Still No Unicorn: Induced Pluripotent Stem Cells from New Species

Self-renewing and pluripotent iPSCs

- Powerful alternatives for animal research
- Hope for endangered species ?
- Therapeutic applications for captive animals.
- For nearly extinct species, iPSCs may be a means to rescue species from extinction?
- Preserving the genomes of individual animals as pluripotent stem cells opens the possibility of producing iPSC-derived germ cells (COURS V), which could be used in conjunction with advanced assisted reproduction efforts to increase the size and diversity of the population.







How general is the Yamanaka strategy? Can other mammalian cells be reprogrammed to pluripotency?



Ben-Nun et al, Nature Methods, 2011





factors ve. The factors (Boyer). They le supmouse heir tarrinhibihistone hodifies bind to teracts is gene (Evans

of hES). They i differis great S cells , transnen the role of ster of enance

Summary and open questions from the first iPS papers

- Four transcription factors (Oct4, Sox2, Klf4 and Myc) are sufficient for nuclear reprogramming of somatic cells into "induced pluripotent stem" iPS cells.
- **Inefficient** (0.1-2%) **but highly reproducible** : three subsequent studies from different labs obtained reprogramming to pluripotency within one year!
- **iPS resemble ESCs** (self renewing, pluripotent -> give rise to all three germ layers *in vitro*) but are they truly equivalent? ? Do they have any "memory" of their somatic history?



nto

Table 1 - Data 5 The differential iPSCr-ESCrind L: Lin28, Mr e-1

Donie cella

NPC MSC fibroblast Foreskin Foreskin Foreskin Fibroblast Fibroblast Fibroblast dH1f fibrobla dillent BJ fibroblast dH1F fibrobla dRTF fibroblas BJI fibroblast R1 fibroblast MRC5 fibrob1 UII fibroblast CB CD133+ CB CD133+ BJ sample

factors (Boyer (Boyer). They le supmouse heir tarrinhibihistone hodifies bind to iteracts is gene (Evans

of hES They differis great S cells transten the role of enance

Summary and open questions from the first iPS papers

- Four transcription factors (Oct4, Sox2, Klf4 and Myc) are sufficient for nuclear reprogramming of a somatic cell into an "indued pluripotent stem" iPS cells.
- **Inefficient** (0.1-2%) **but highly reproducible**: three subsequent studies from different labs obtained reprogramming to pluripotency within one year!
- **iPS resemble ESCs** (self renewing, pluripotent -> give rise to all three germ layers *in vitro*) but are they truly equivalent? ? Do they have any "memory" of their somatic history?
- Why do these <u>particular</u> factors achieve reprogramming?
- Oct4, Sox2, Klf4 TFs cooperatively suppress lineage specific genes and activated ES-related genes leading to self-sustaining pluripotency network essential TFs for pluripotency (Boye et al, 2005; Loh et al, 2006; Wang et al 2006; Orkin and Hochedlinger, 2011 for review)
 c-*Myc* = Facilitator of reprogramming? It increases expression of many genes important for proliferation and self-renewal (eg Telomerase Cartwright et al., 2005).
- Why is induced pluripotency so slow and inefficient?



Dot3/4

Sox2

c-myc

Figure

Role(s) of the Murine Reprogramming Factors in the Induction of Pluripotency?



stone

hylases , Jmjd2b/2c, /1b, Utx, Uty,

e variants

13.3, H2A.Z, croH2A

> Each (and all) of the OKSM factors can be functionally replaced by <u>other TFs, by</u> <u>miRNAs, small compounds...(=> COURS IV</u>)

> > Figure 7. Ro A) c-Myc is a n activates many

в

Are iPS cells truly equivalent to ES cells?



Why is induced pluripotency so inefficient?



Although I in the char

How simil

(such as tel markers, ar several stud differences individual I alterations teome and iPSCs were about the s However, o genetic abn from ESCs² extent of va similar to v within diffe

Recently malities se stress indu A substant H2A.X - (double-stra exposed to homologou for error-fr ming proce way in mai the availabl tions seen i process per genetic an parental fib Much et

Much et ties, such a vidual ESC unpredicta potentially research. I disease-spe be relevant tem-immai towards de ESC deriva of the cells

E. Heard, March 24th 2014

Adapted from Loh and Lim, 2010

Induced Pluripotency is slow and inefficient

WHY?

Passaging improves iPS cell pluripotency Dividing and amplifying the cells allows epigenetic memory to be erased



MET gener

H3K4me2

Must have similar steps to normal reprogramming (eg in the germ line):

Silencing of somatic cell program and activation of self-renewing/pluripotency program

Need to override epigenetic barriers:

Interfering with epigenetic processes increases frequencies up to 10% or more! (COURS IV)

Figure 1 process of shorter, h After a fil assume of reprogra modifical processin delays th In this ph regulator cases, the embryon

Histone Functi merk H3K4me2 Marks p H3K4me3 Marks a H3K27me3 Marks f H3K4me3 Marks f H3K4me3 Marks f H3K9me3 Marks f H3K36me2 Marks f Soch a

EMT, epithelial-to-mes lysine 4: H3K27ac, histo

Table 2 | Roles of ex Chromatin modifier factor

KDM2A and KDM2B

EHMT1 and SETDB1 BMI1, RING1, EZH2, EED and SU[212

T genes

Induced Pluripotency is slow and inefficient

Epigenetic barriers, that were imposed on the genome during differentiation, to stabilize cell identity and prevent aberrant cell fate changes, must be overcome during reprogramming



Nature of this Epigenetic Memory? Ways to overcome it? (COURS IV)

X-chromosome reactivation as a "gold standard" for efficient reprogramming?



reality -

Figure 1 process of shorter, h After a fil assume of reprogra modifical processis delays th In this phy regulator cases, th embryon

nerk 13K4me2 Marks 13K4me3 Marks 13K27me3 Marks 13K4me1 Marks 13K4me1 Marks

> Ame3 Marks F S6me2 Marks p (such a P9me2 Marks t 27ac Marks c enhanc

Table 2 | Roles of exa Chromatin modifier factor UTX

KDM2A and KDM2B

EHMT1 and SETDB1 BMI1, RING1, EZH2,

E. Heard, March 24th 2014

iPS and epigenetic reprogramming of mouse cells Xi Reactivation?



Mouse female iPS cells showed reactivation of a somatically silenced X chromosome and undergo random X inactivation upon differentiation.



COLL

iPS and epigenetic reprogramming of human cells Xi Reactivation?



Female Human iPSCs Retain an Inactive X Chromosome

The inactive X is not reactivated during human iPS cell induction Culture conditions, Xi status in hESCs, Xi status in human embryos (ICM) SEMINAR, CLAIRE ROUGEULLE





Tchieu et al, 2010

0/6419

rnold,

Perspectives brought by iPS cells

- Cell and Tissue therapy without use of human embryos and reduces problems associated with compatibility (COURS V)
- > Enables study of differentiation and development *in vitro*
- Powerful tool for drug screening
- Enables genetic engineering for functional investigation of development and disease *in vitro* and *in vivo* (production of chimeric animals)





Issues/problems:

- Obtaining iPS cell lines swifly and then sufficient cell numbers upon *in vitro* differentiation can be a problem (in a therapeutic context)
- iPS still brings a risk of cancer (even avoiding *myc* and integrative vectors...) (*Why does the iPS cell production process create oncogenically transformed cells*)

Can safety and efficiency be improved?



3 G E NCE

IMPROVED DELIVERY OF TFs

OVERCOMING EPIGENETIC BARRIERS

IMPROVED CULTURE CONDITIONS (COURS V)



IMPROVED DELIVERY OF TFs

Induced Pluripotent Stem Cell Generation Using a Single Lentiviral Stem Cell Cassette

CESAR A. SOMMER,^a MATTHIAS STADTFELD,^b GEORGE J. MURPHY,^c KONRAD HOCHEDLINGER,^b DARRELL N. KOTTON,^{d,e} GUSTAVO MOSTOSLAVSKY^a



Human Induced Pluripotent Stem Cells Free of Vector and Transgene Sequences

Junying Yu,^{1,2,3}* Kejin Hu,³ Kim Smuga-Otto,^{1,2,3} Shulan Tian,^{1,2} Ron Stewart,^{1,2} Igor I. Slukvin,^{1,4} James A. Thomson^{1,2,3,5}*

Stem Cell Reports



-OPEN ACCESS

Reprogramming to Pluripotency Using Designer TALE Transcription Factors Targeting Enhancers

Xuefei Gao,¹ Jian Yang,¹ Jason C.H. Tsang,¹ Jolene Ooi,¹ Donghai Wu,² and Pentao Liu^{1,*}

A more efficient method to generate integration-free human iPS cells We report a simple method, using p53 suppression and nontransforming L-Myc, to generate human induced pluripotent stem cells (iPSCs) with episomal plasmid v we generated human iPSCs from multiple donors, inclu

Keisuke Okita¹, Yasuko Matsumura¹, Yoshiko Sato¹, Aki Okada¹, Asuka Morizane^{1,2}, Satoshi Okamoto³, Hyenjong Hong¹, Masato Nakagawa¹, Koij Tanabe¹, Ken-ichi Tezuka⁴, Toshiyuki Shibata⁵, Takahiro Kunisada⁴, Masayo Takahashi^{1,3}, Jun Takahashi^{1,2}, Hiroh Saji⁶ & Shinya Yamanaka^{1,7–9} We report a simple method, using p53 suppression and nontransforming L-Myc, to generate human induced pluripotent stem cells (iPSCs) with episomal plasmid vectors. We generated human iPSCs from multiple donors, including two putative human leukocyte antigen (HLA)-homozygous donors who match ~20% of the Japanese population at major HLA loci; most iPSCs are integrated transgene-free. This method may provide iPSCs suitable for autologous and allologous stem-cell therapy in the future.

OVERCOMING EPIGENETIC BARRIERS



IMPROVED DELIVERY OF TFs

Induced Pluripotent Stem Cell Generation Using a Single Lentiviral Stem Cell Cassette

CESAR A. SOMMER,^a MATTHIAS STADTFELD,^b GEORGE J. MURPHY,^c KONRAD HOCHEDLINGER,^b DARRELL N. KOTTON,^{d,e} GUSTAVO MOSTOSLAVSKY^a



Human Induced Pluripotent Stem Cells Free of Vector and Transgene Sequences

Junying Yu,^{1,2,3}* Kejin Hu,³ Kim Smuga-Otto,^{1,2,3} Shulan Tian,^{1,2} Ron Stewart,^{1,2} Igor I. Slukvin,^{3,4} James A. Thomson^{1,2,3,5}*

Stem Cell Reports



OPEN ACCESS

Reprogramming to Pluripotency Using Designer TALE Transcription Factors Targeting Enhancers

Xuefei Gao,¹ Jian Yang,¹ Jason C.H. Tsang,¹ Jolene Ooi,¹ Donghai Wu,² and Pentao Liu^{1,*}

A more efficient method to generate integration-free human iPS cells

Keisuke Okita¹, Yasuko Matsumura¹, Yoshiko Sato¹, Aki Okada¹, Asuka Morizane^{1,2}, Satoshi Okamoto³, Hyenjong Hong¹, Masato Nakagawa¹, Koij Tanabe¹, Ken-ichi Tezuka⁴, Toshiyuki Shibata⁵, Takahiro Kunisada⁴, Masayo Takahashi^{1,3}, Jun Takahashi^{1,2}, Hiroh Saji⁶ & Shinya Yamanaka^{1,7–9} We report a simple method, using p53 suppression and nontransforming L-Myc, to generate human induced pluripotent stem cells (iPSCs) with episomal plasmid vectors. We generated human iPSCs from multiple donors, including two putative human leukocyte antigen (HLA)-homozygous donors who match -20% of the Japanese population at major HLA loci; most iPSCs are integrated transgene-free. This method may provide iPSCs suitable for autologous and allologous stem-cell therapy in the future.

OVERCOMING EPIGENETIC BARRIERS

Vitamin C modulates TET1 function during somatic cell reprogramming

Jiekai Chen^{1,2,6}, Lin Guo^{1,2,6}, Lei Zhang^{3,6}, Haoyu Wu^{1,4}, Jiaqi Yang^{1,2}, He Liu^{1,2}, Xiaoshan Wang^{1,2}, Xiao Hu³, Tianpeng Gu³, Zhiwei Zhou^{1,2}, Jing Liu^{1,2}, Jiadong Liu^{1,5}, Hongling Wu^{1,2}, Shi-Qing Mao³, Kunlun Mo^{1,2}, Yingying Li^{1,2}, Keyu Lai^{1,2}, Jing Qi^{1,2}, Hongjie Yao^{1,2}, Guangjin Pan^{1,2}, Guo-Liang Xu³ & Duanqing Pei^{1,2}

How microRNAs facilitate reprogramming to pluripotency

Frederick Anokye-Danso^{1,*} Melinda Snitow^{2,*} and Edward F. Morrisev^{1,2,3,4,‡}

The use of small molecules in somatic-cell reprogramming

Alexander J. Federation^{1,2,3}, James E. Bradner^{1,2,4}, and Alexander Meissner^{2,5,6}

Derivation of novel human ground state naive pluripotent stem cells

Ohad Gafni¹*, Leehee Weinberger¹*, Abed AlFatah Mansour¹*, Yair S. Manor¹*, Elad Chomsky^{1,2,3}*, Dalit Ben-Yosef^{4,5}, Yael Kalma³, Sergey Viukov¹, Itay Maza¹, Asaf Zviran¹, Yoach Rais², Zohar Shipony^{2,3}, Zohar Mukamel^{2,3}, Vladklav Krupalnik¹, Mirie Zerbib¹, Shay Geula¹, Inbal Caspi¹, Dan Schneir¹, Tamar Shwartz⁴, Shlomit Gilad⁶, Daniela Amann-Zalcenstein⁶, Sima Benjamin⁶, Ido Amit², Amos Tanay^{2,3}, Rada Massarwa¹, Noa Novershtem¹ & Jacob H. Hanna¹

Removing Reprogramming Roadblocks: Mbd3 Depletion Allows Deterministic iPSC Generation

Justin Brumbaugh^{1,2} and Konrad Hochedlinger^{1,2,3,*}



IMPROVED DELIVERY OF TFs

OVERCOMING EPIGENETIC BARRIERS

NO EXOGENOUS FACTORS

Pluripotent Stem Cells Induced from Mouse Somatic Cells by Small-Molecule Compounds

Pingping Hou,¹* Yanqin Li,¹* Xu Zhang,^{1,2}* Chun Liu,^{1,2}* Jingyang Guan,¹* Honggang Li,¹* Ting Zhao,¹† Junqing Ye,^{1,2}† Weifeng Yang,³† Kang Liu,¹† Jian Ge,^{1,2}† Jun Xu,¹† Qiang Zhang,^{1,2}† Yang Zhao,¹‡ Hongkui Deng^{1,2}‡

Stimulus-triggered fate conversion of somatic cells into pluripotency

Haruko Obokata^{1,2,3}, Teruhiko Wakayama³†, Yoshiki Sasal⁴, Koji Kojima¹, Martin P. Vacanti^{1,5}, Hitoshi Niwa⁶, Masayuki Yamato⁷ & Charles A. Vacanti¹



ransgenes and chemical modulation, thus signifying a milestone in advancing our understanding ncy induction.

Beyond the Yamanaka concept: Induced Pluripotency by Lineage Specifiers



Using TFs to induce *direct* lineage conversion (Trans-differentiation)



Phenotypes Induced in 1 OT: Cultures after Treatment with 5-aza-CR (a) Adipocytes (4 weeks after treatment); (b) myotubes (2 weeks after treatment); (c) chondrocytes (5 weeks after treatment). *Taylor and Jones, 1979.*

Observations by Peter Jones and colleagues in the 1970's

5-aza-C treated fibroblasts sometimes gave rise to cells with "new phenotypes"(adipocytes, myotubes, chondrocytes...)

"possibly by inducing a reversion to a more pluripotential state from which the new phenotypes subsequently differentiate..?"

Using TFs to induce *direct* lineage conversion (Trans-differentiation)



Identify key TFs that induce lineage conversion

A computational biology approach to find regulators of cell fate (Heinäniemi et al, 2013): Bioinformatics approach that searches public gene-expression data sets for candidate transcriptional regulators for many human cell types. \Rightarrow a new resource for experiments aimed at direct lineage conversion



Pairwise comparison of TF expression levels extracted from many gene array data sets yields candidate cell type– specific master regulators. Solid-colored ovals represent the dominant TF in a given pair for a given cell type. (b) Quantification of pairwise relationships between TFs can be used to visualize lineage relationships in a topographical manner. (c) Identified TFs may be candidate factors to reprogram one cell type into another.



Direct lineage conversion for generation of specific cell types



- Identification of specific transcription factors controlling differentiated cell identity allows for the forced conversion of one lineage into another in the absence of cell proliferation.
- Direct conversion of numerous cell types now shown including neurons, cardiomyoctes and hepatocytes. Direct lineage conversion is sufficient for the generation of distantly related cell types crossing germ layer boundaries.

From Sancho-Martinez et al, Nat. Cell Biol. 2012



Bm4 Pou3t4 Sox2 KH4 c-Myc E47 Tot3

E. Heard, March 24th 2014

Indirect lineage conversion: for generation of specific cell types in absence of specific TFS



Ignacio



- Indirect lineage conversion: a more general approach in the absence of specific transcription factors.
- ➤ Relies on the use of pluripotency TFs (OSKM) initially.
- ➢ On forced OSKM expression, this first leads to removal of differentiated marks, creating an unstable state suitable for further differentiation on exposure to appropriate signals.



of neage ktail nsion phase'. tal cues

Different conversion methodologies to generate cells of a given fate



From Sancho-Martinez et al, Nat. Cell Biol. 2012



Multiple Reprogramming Perspectives

- Cell and tissue therapy perspectives using iPS, direct or indirect cell conversions
- Study of development and disease *in vitro*
- > Powerful tools for drug screening
- Genetic engineering for functional investigations







Seminaire

Dr Claire Rougeulle

Inactivation du chromosome X, pluripotence et reprogrammation, de la souris à l'homme