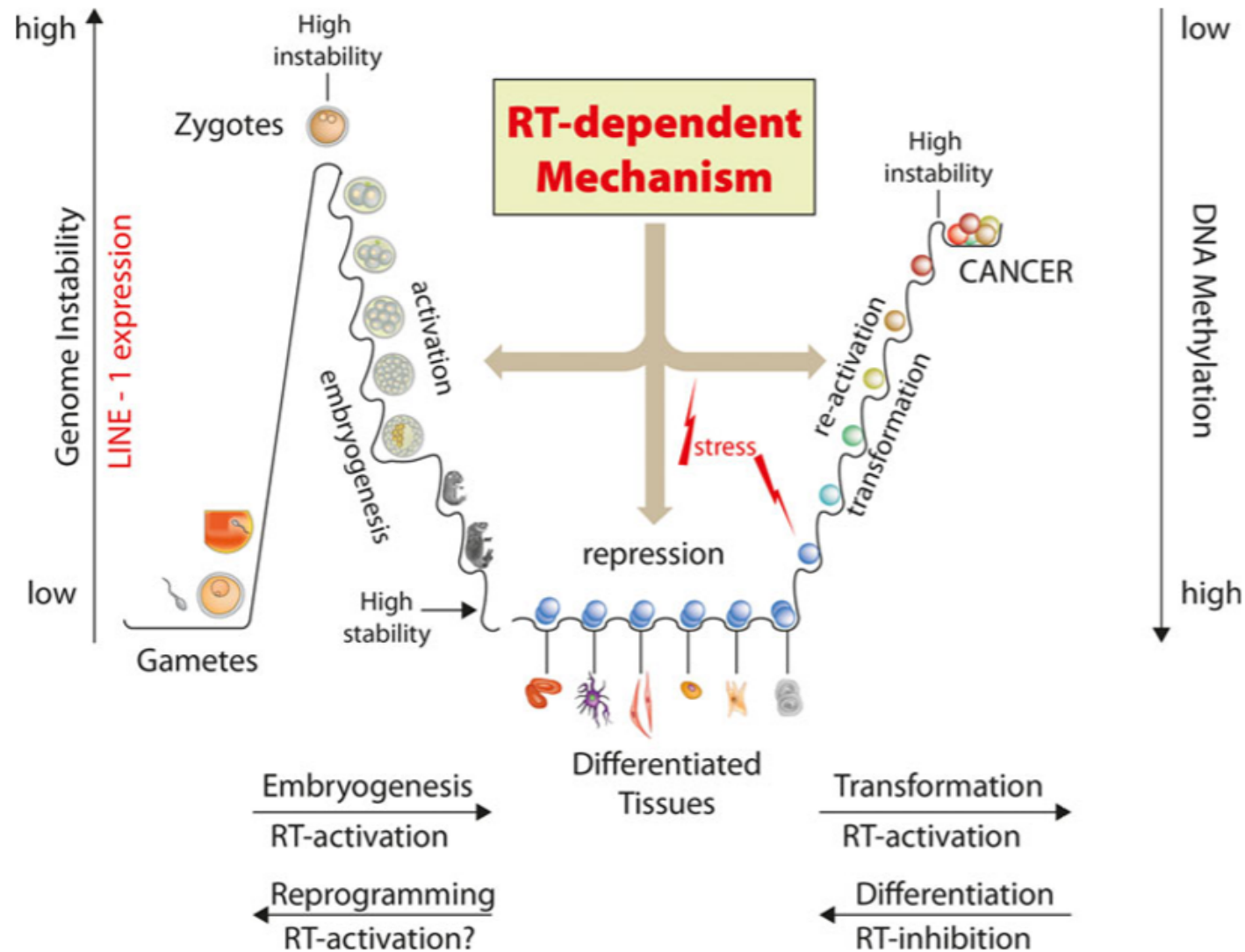


PSEUDOGENE lncRNAs EXAMPLE 1

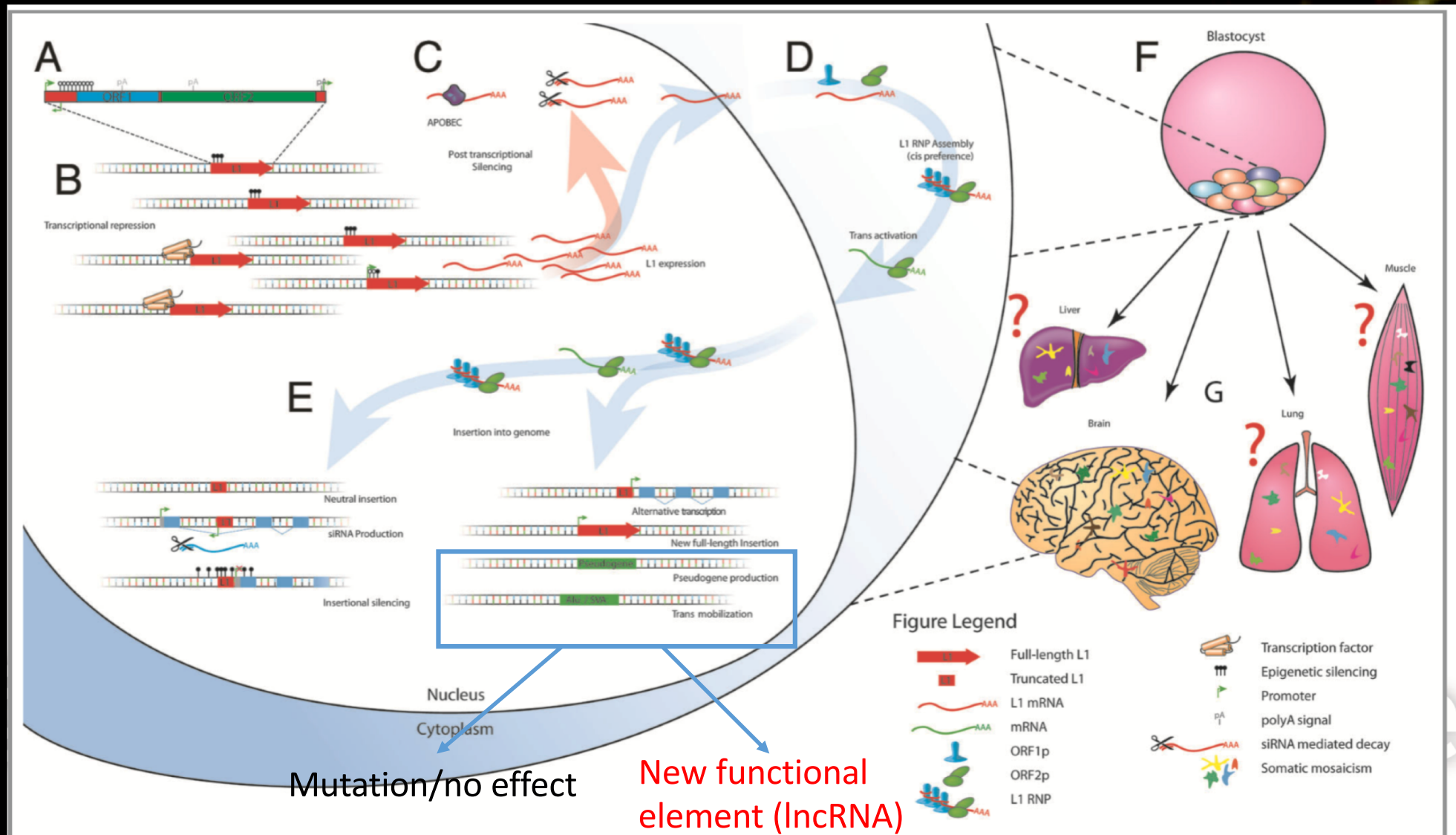
***Pseudogene lncRNA that controls embryonic stem cell
Self-renewal***

***A Oct4P4 pseudogene derived lncRNA silences the
ancestral Oct4 gene in trans***

Retrotransposon activity during development

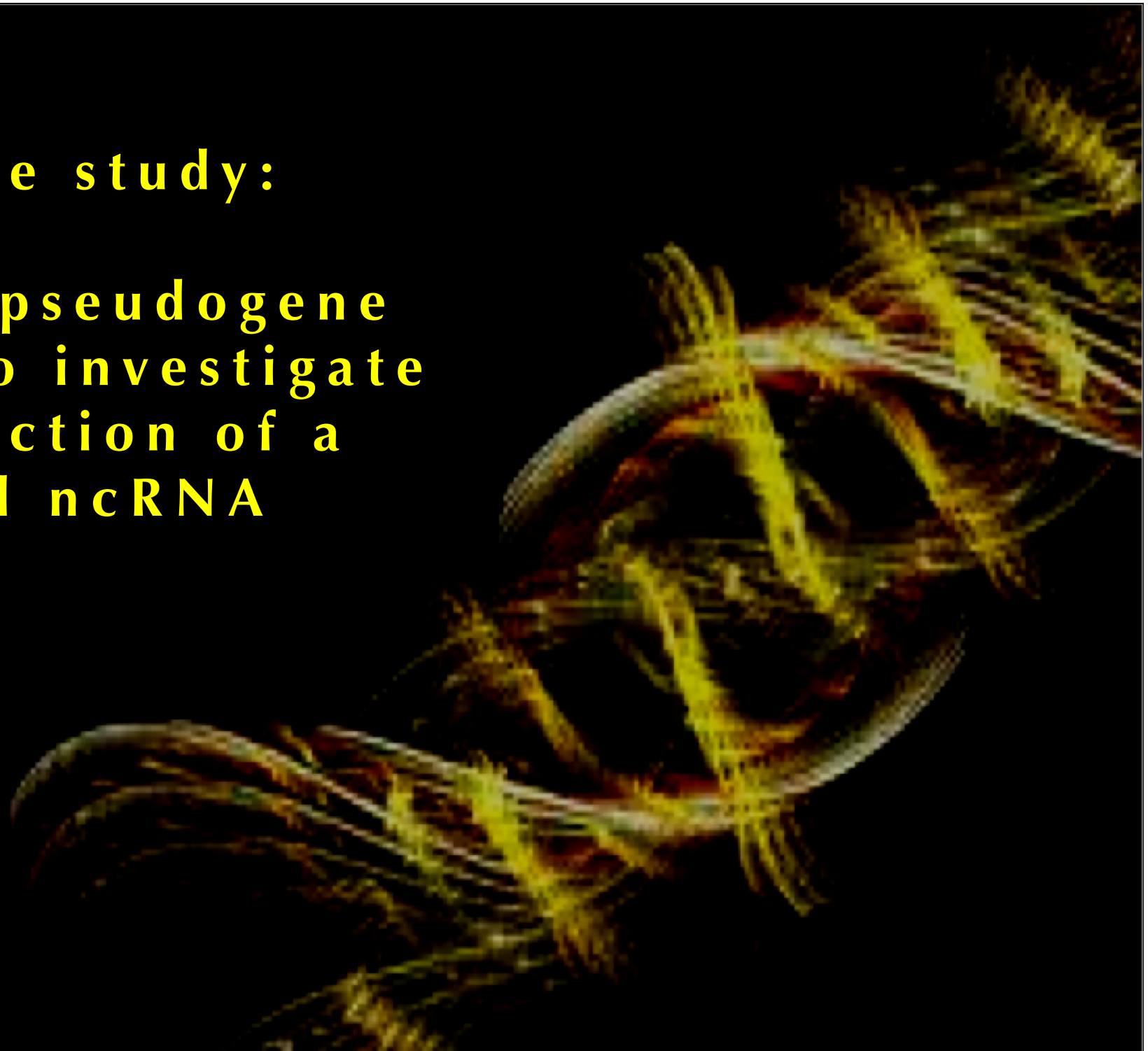


Retrotransposons can change genetic Context → mosaic somatic tissues



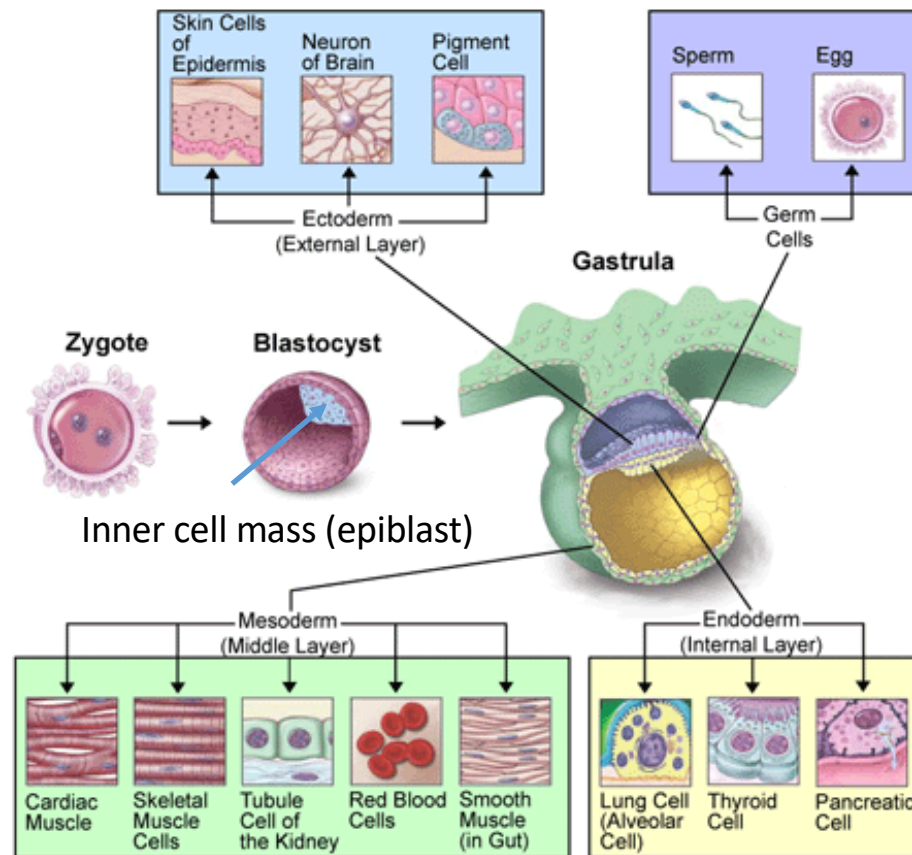
A case study:

**Using a pseudogene
lncRNA to investigate
the function of a
novel ncRNA**

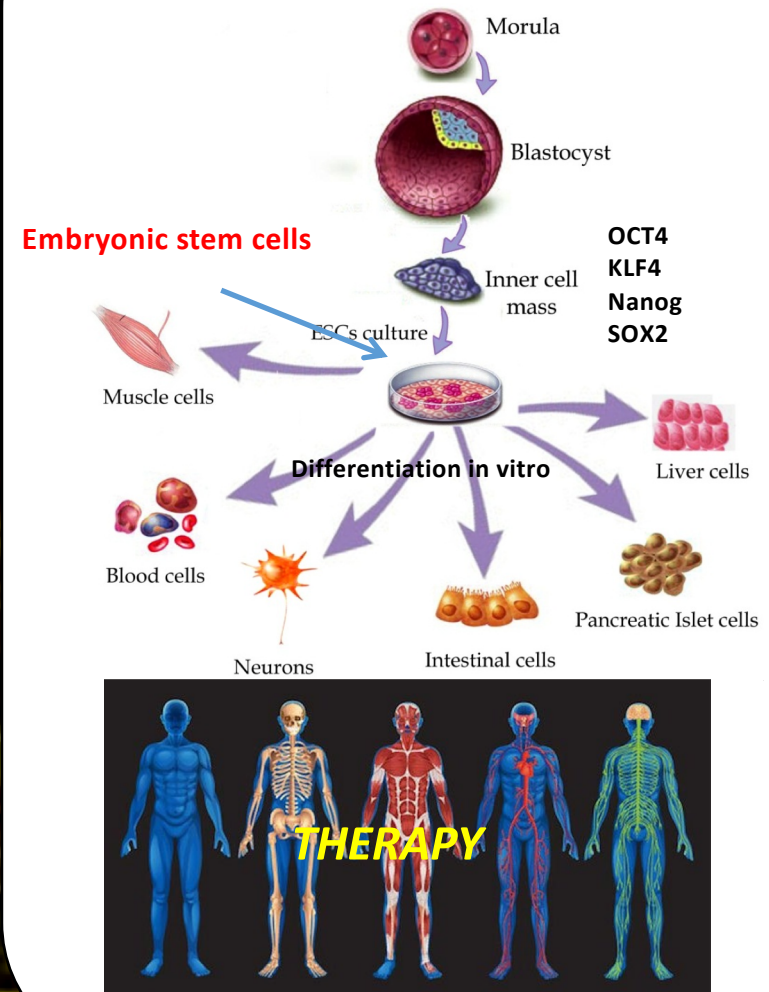


The inner cell mass of the blastocyst are the source of pluripotent embryonic stem cells

Blastocyst cells give rise to all organs and cell types

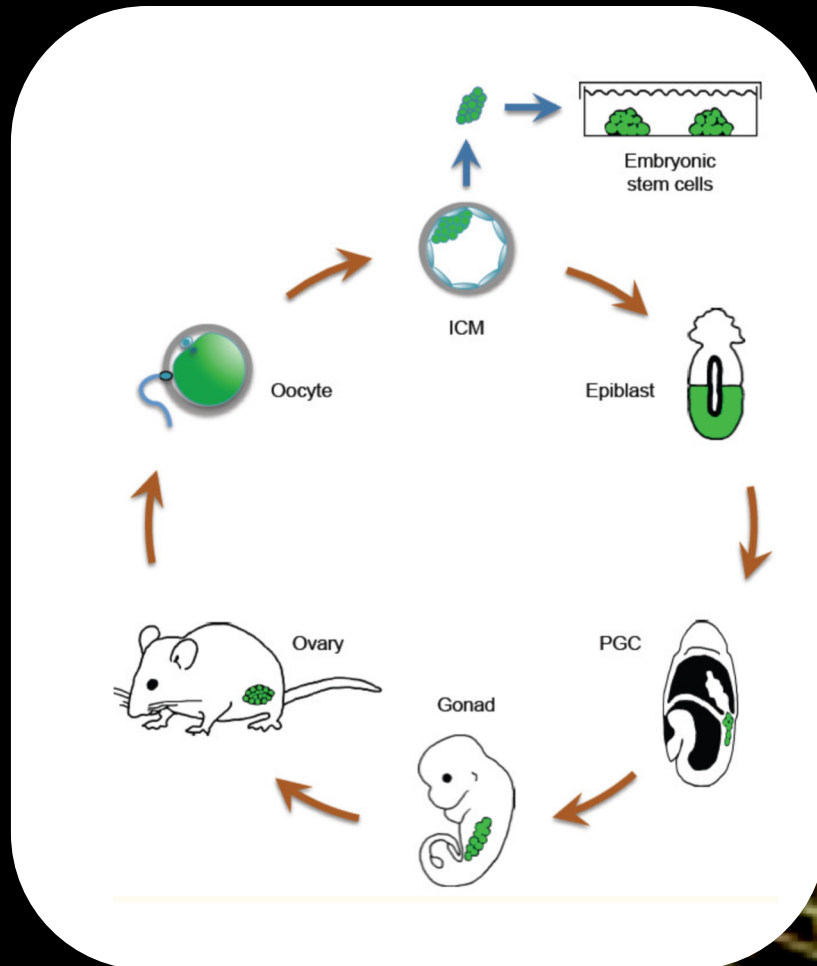


Blastocyst cells can be isolated and cultivated



1. OCT4 EXPRESSION

Oct4 expression during the mouse life cycle

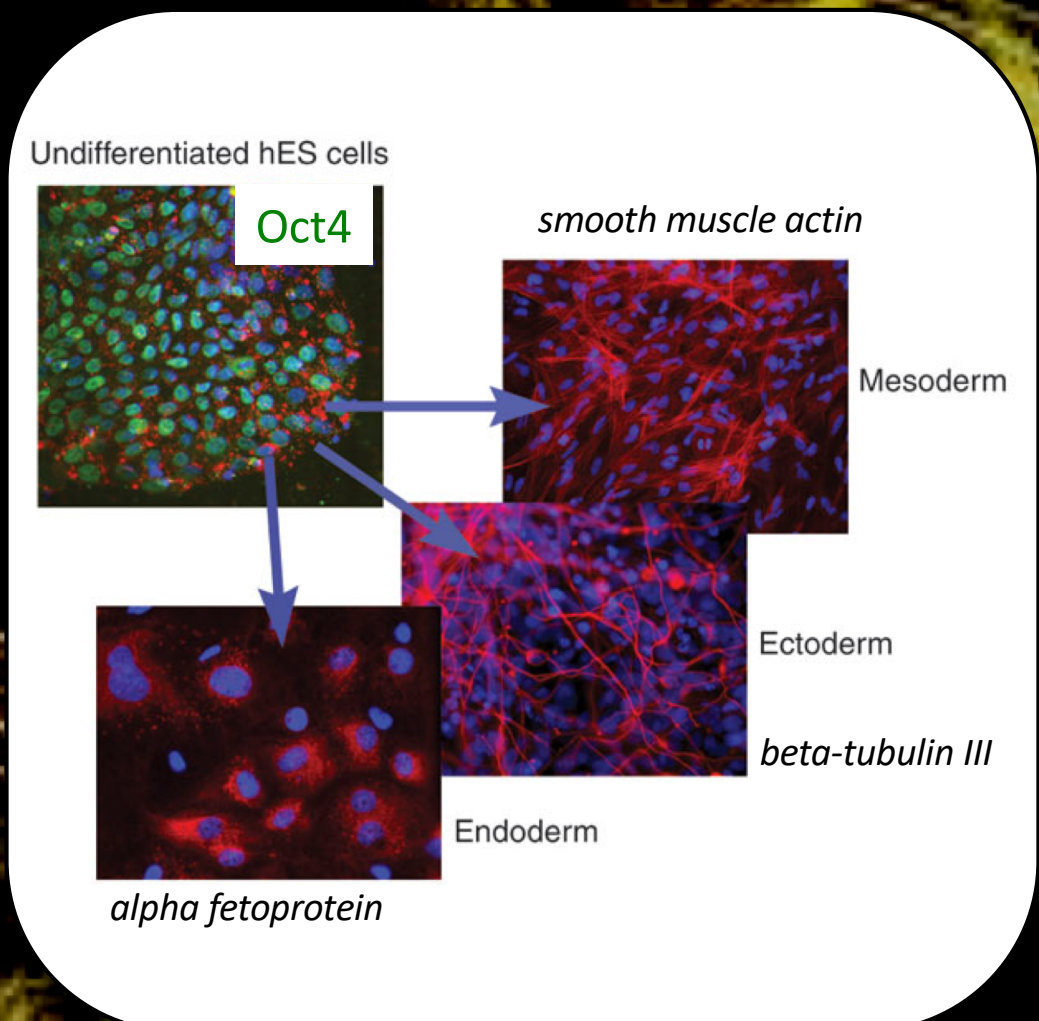
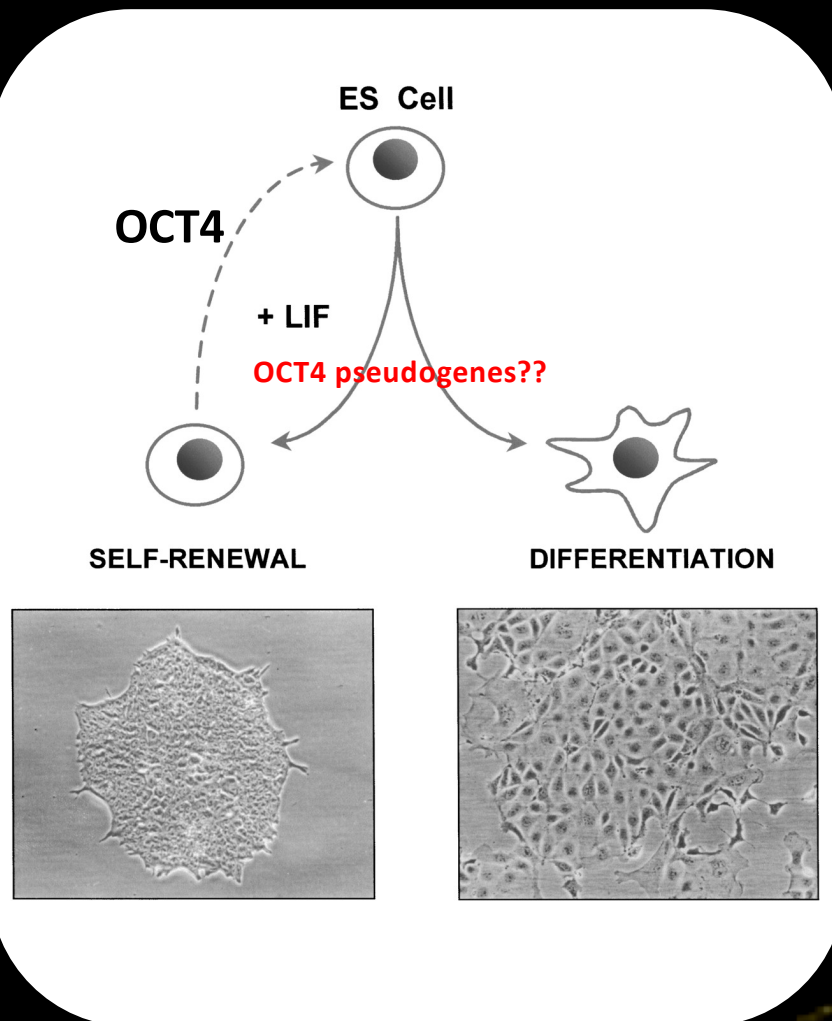


Cells and tissues expressing Oct4 are marked in green.

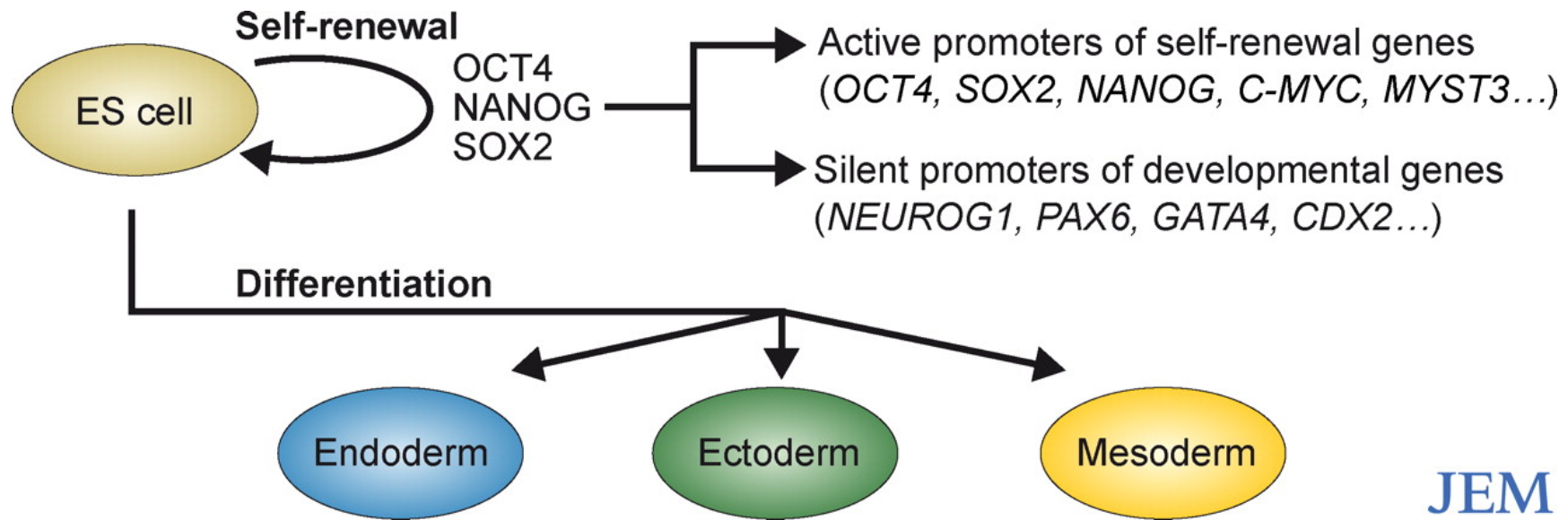
- Oct4 is expressed in mouse oocytes as a maternal transcript and protein.
- Zygotic Oct4 expression is activated prior to the 8-cell stage and is abundant and uniform in all cells of the embryo throughout the morula stage.
- However, as the outer cells of the embryo differentiate into the Trophectoderm, Oct4 expression is restricted to cells of the ICM in the blastocyst.
- After implantation, Oct4 expression is maintained in the epiblast of the blastocyst
- Oct4 expression becomes restricted to primordial germ cells (PGCs), which are first specified in the extraembryonic mesoderm at the base of the allantoic bud during gastrulation. PGCs give rise to gametes, which following fertilization will develop into a new organism of next generation.

2. THE MODEL SYSTEM: ES CELL DIFFERENTIATION

OCT4 expressing ES cells have self-renewing and differentiation potential *in vitro*



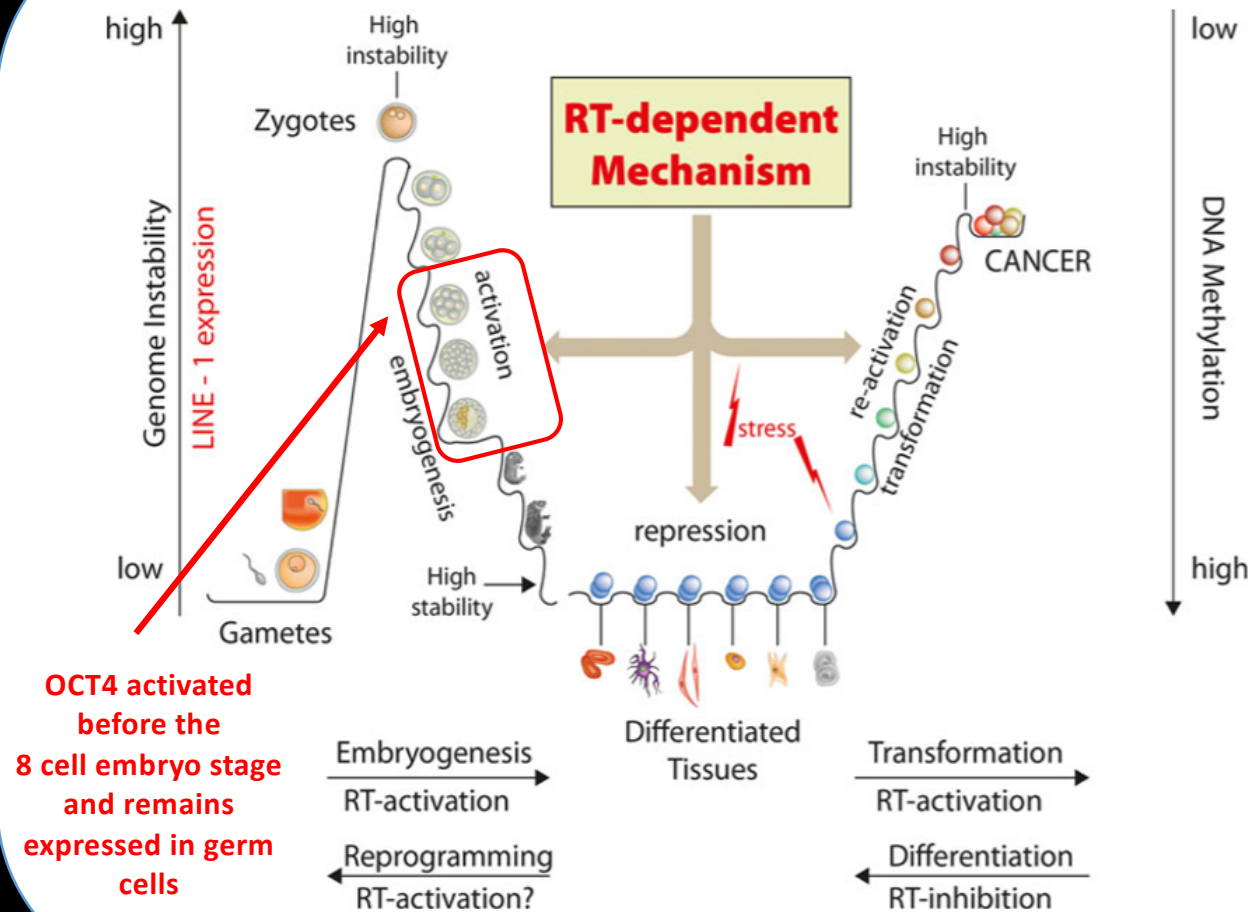
The self-renewal transcription factor Oct4 is essential for embryonic stem cell self-renewal



JEM

Nicolaj Strøyer Christophersen, and Kristian Helin J Exp Med 2010;207:2287-2295

Mouse and human contain several processed OCT4 pseudogenes



Human:
1 ancestral OCT4
7 processed OCT4
pseudogenes

Mouse:
1 ancestral Oct4
5 processed OCT4
pseudogenes

3. Sequence analysis:

Are there other sequences in the mouse genome that are similar to Oct4?

Use Oct4 cDNA and make BlastN

Descriptions

Graphic Summary

Alignments

Taxonomy

Sequences producing significant alignments

Download

New Select columns

Show

100

select all

6 sequences selected

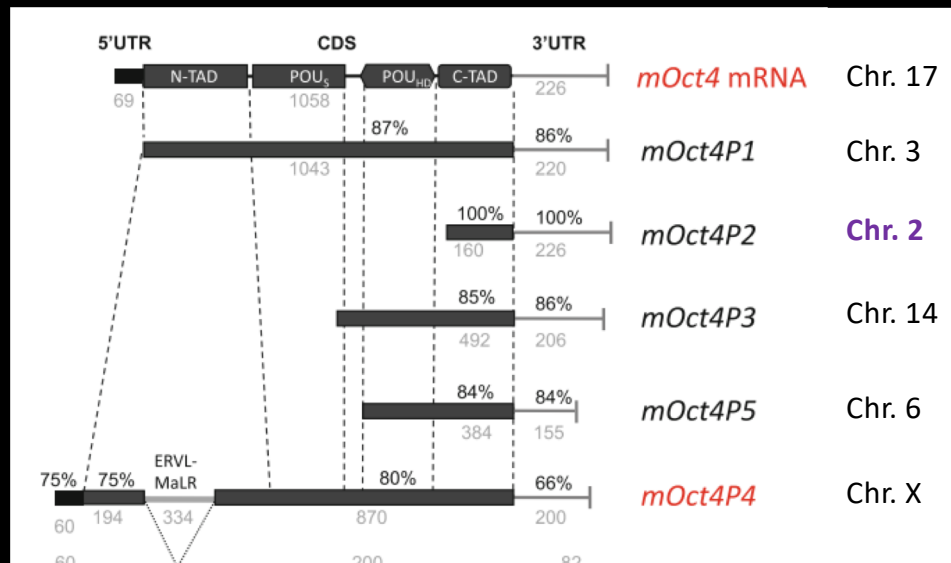
GenBank

Graphics

Distance tree of results

New MSA Viewer

	Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
☒	Mus musculus strain C57BL/6J chromosome 3, GRCh39	Mus musculus	1177	2059	100%	0.0	87.15%	159745316	NC_000069.7
☒	Mus musculus strain C57BL/6J chromosome 17, GRCh39	Mus musculus	710	1977	100%	0.0	100.00%	95294699	NC_000083.7
☒	Mus musculus strain C57BL/6J chromosome X, GRCh39	Mus musculus	638	638	82%	2e-180	80.20%	169476592	NC_000086.8
☒	Mus musculus strain C57BL/6J chromosome 14, GRCh39	Mus musculus	520	520	46%	8e-145	85.71%	125139656	NC_000080.7
☒	Mus musculus strain C57BL/6J chromosome 6, GRCh39	Mus musculus	383	383	36%	1e-103	84.62%	149588044	NC_000072.7
☒	Mus musculus strain C57BL/6J chromosome 1, GRCh39	Mus musculus	303	303	15%	9e-80	100.00%	195154279	NC_000067.7



Unspliced sequence on Chr. 2 with high similarity to Oct4

Mus musculus strain C57BL/6J chromosome 3, GRCh39
Sequence ID: [NC_000069.7](#) Length: 159745316 Number of Matches: 2

Range 1: 129227322 to 129228364
[GenBank](#)
[Graphics](#)

[▼ Next Match](#)
[▲ Previous Match](#)

Score	Expect	Identities	Gaps	Strand
1177 bits(637)	0.0	936/1074(87%)	46/1074(4%)	Plus/Plus

Features:

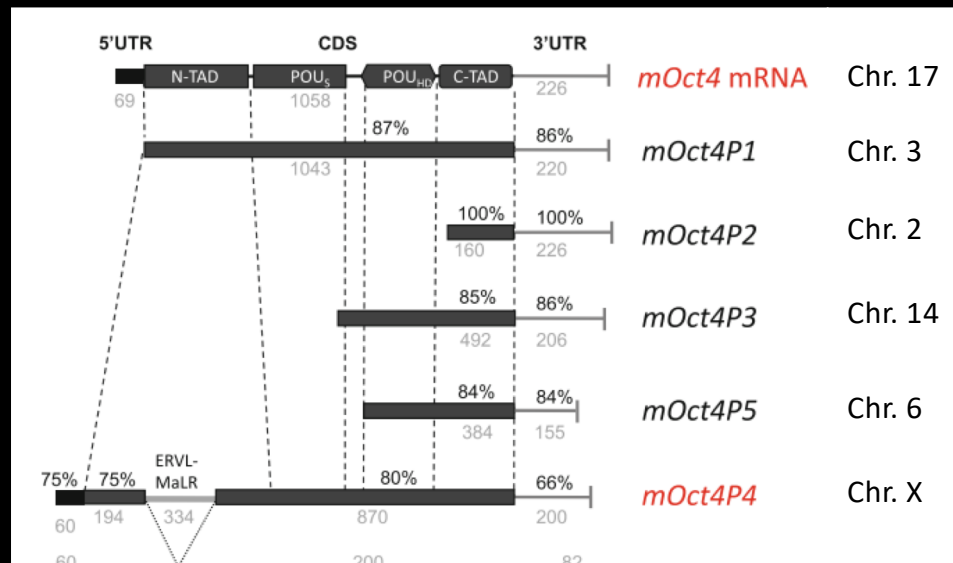
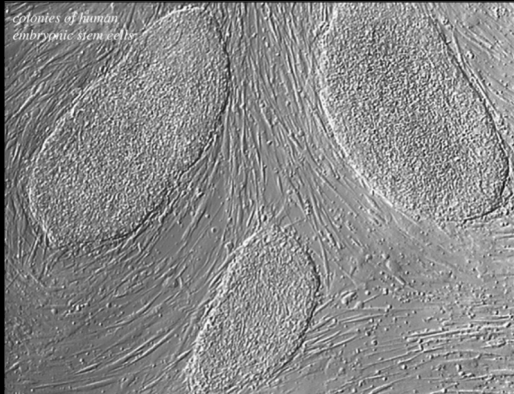
101192 bp at 5' side: [glutamyl aminopeptidase](#)

97965 bp at 3' side: [elongation of very long chain fatty acids protein 6](#)

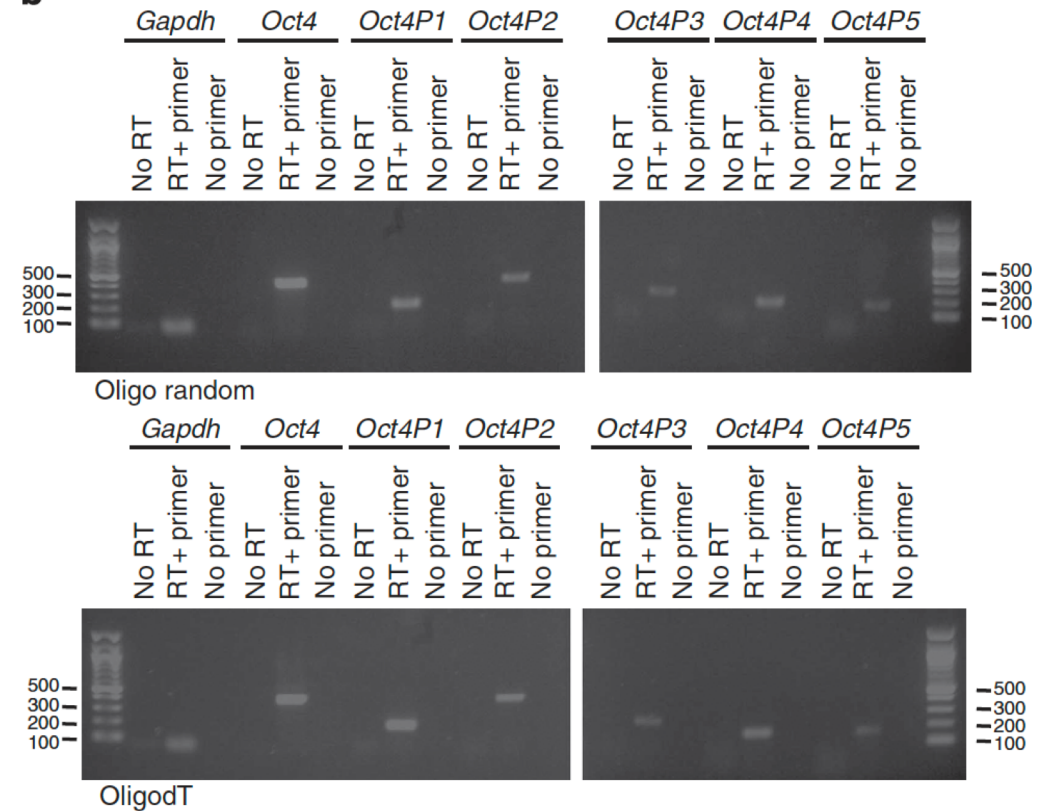
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Query	61	TCAGCAGGGCTGGAGCCGGCTGGGTGGATCCTCGAACCCTGGCTTAAGCTTCCAAGGGCT	120
Sbjct	129227381	TCAGCAGGGCTGGAGCCGGCTGGGTGGACCTCGAACCCTGGCTTAAGCTTCCAAGGGCT	129227440
Query	121	CCAGGTGGGCTGGAATCGGACAGGCTCAGAGGTATTGGGGATCTCCCATGTCGCCGCC	180
Sbjct	129227441	CTGGGTGGGCTGGAATAGGACAGGCTTAGAGGTATT-GGGATCTCCCATGTCACACG	129227499
Query	181	GCATACGAGTTCTGCGGAGGGATGGCATACTGTGGACCTCAGTTGGACTGGGCTAGTC	240
Sbjct	129227500	GTATATGAGTTCTGCGGAGGGATGGCATACTGTGGACCTCAG-----GGGCTGGGCTAGTC	129227541
Query	241	CCCCAAGTTGGCTGGAGACTTTGAGCCCTGAGGGCCAGGAGGACGAGTGGAAAGC	300
Sbjct	129227542	-----GCTGGCTGGAGACTTTGAGCCCTGAGAGCCAGGCAAGAGCAGGTGGAGAGC	129227595
Query	301	AACTCAGAGGGAACCTCTCTGAGCCCTGTGCCGACCGCCCAATGCGCTGAAGTTGAG	360
Sbjct	129227596	AACTCAGAGGGAACCTCTCTGGGCTTGTGCCGACCAACCCAGTGCCTGAAGTTGAG	129227655
Query	361	AAGTTGGAACCAACTCCCGAGGAGTCCCGAGGACATGAAAGCCCTGCAGAGGAGCTAGAA	420
Sbjct	129227656	ATGAGGGAACCAACTCCCGAGGAGTCCCGAGGACATGAAAGCTTCGAGAGGAGCTAGAG	129227715
Query	421	CAGTTTGGCAAGCTGCTGAAGCAGAAGAGGATCACCTTGGGTTACACCCAGGCCGACGTT	480
Sbjct	129227716	AAGTTTGGCAAGCTGCTGAAGCAGAAGAGGATCACCTTGGGTTACACCCAGGCCGACGTT	129227775
Query	481	GGGCTCACCTGGGCGTTCTCTTGGAAAGGTGTTGAGCCAGACCACTCTGTCGCTTC	540
Sbjct	129227776	GGGCTCACCTGGGCGTTCTCTTCAGAAAGTG-TGAGCCAGACCACTCTACTGCTTT	129227834
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Sbjct	129227835	GAGGCCCCGAGCTCAGCCTTATGACCATGCATAGCTGGGGCCCTGCTGGAGAGTGG	129227894
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Sbjct	129227895	GTAGAGGAAGCCGGAAGACAATGAGAACCTTCAGGAGATATGCAAAATCGGAGACCTGGTG	129227954
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Sbjct	129227955	CAGGCCCGGAAGAGAAGCGGAGCATTGAGAACCTGTGAGGCTGAGGCTGGAGAAAT	129228014
Query	721	ATGTTTCTGAAAGTCCCGAAGCCCTTCCCTACAGCAGATCACTCACA-TGGC--CA--AT-	774
Sbjct	129228015	GAGTTTCTGAAAGTCCCGAAGCCCTTCCCGCAGCAGATCACTAGATATAGCATCACCATG	129228074
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Sbjct	129228075	CAGCTTGGGATGGAAGACATGTTTCGAGGATGGTTCTGTAAACGGGCGCAGAGGGC	129228134
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Sbjct	129228135	AAAAGATCAAGTATTGAGTATCCCAACGAGAAAAGTATGAGGCTTTCGGGGATGCTCT	129228194
Query	893	TCC---CA-ggg-ggggCTGTATCCTTTCTCTGCCCCAGGTCCTCCACTTTGGCACCC	946
Sbjct	129228195	TCGGGCGTGGGGTGGGCTGTATCCTTTCTCTGCCCCAGG-CCCCACTTTGGCACCC	129228253
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Sbjct	129228254	CAAGCTATGGGAGCCCCACTTCAACCACT--ACTCGATACCTG-TCTTGAGGGCGAGG	129228310
Query	1007	CCTTTCCCTCTGTGCC--GTCACTGCTTGGGCTCTCCCATGCAATTCAAACCTGA	1059
Sbjct	129228311	CCTTTCCCTCTGTGCCCTGTCACTGCTTGGGCTCTCCCATGCAATTCAAACCTGA	129228364

4. EXPERIENTAL EVALATION OF lncRNA EXPRESSION IN MODEL CELLS

Self-renewing mESCs



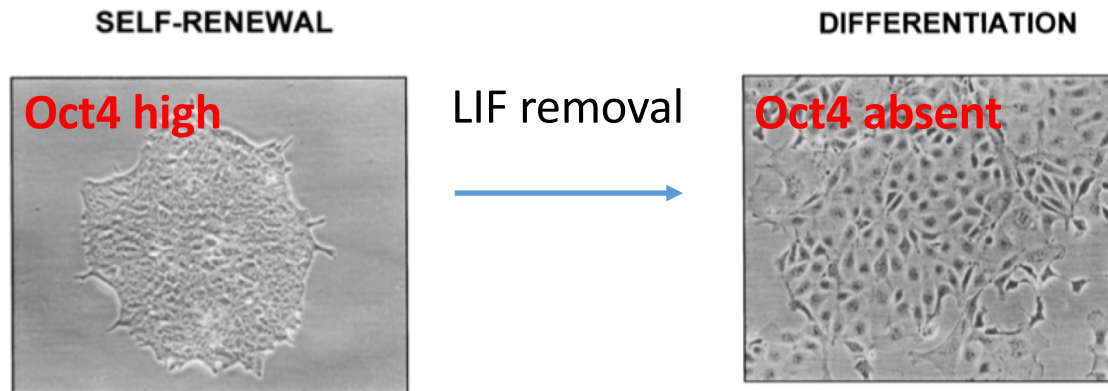
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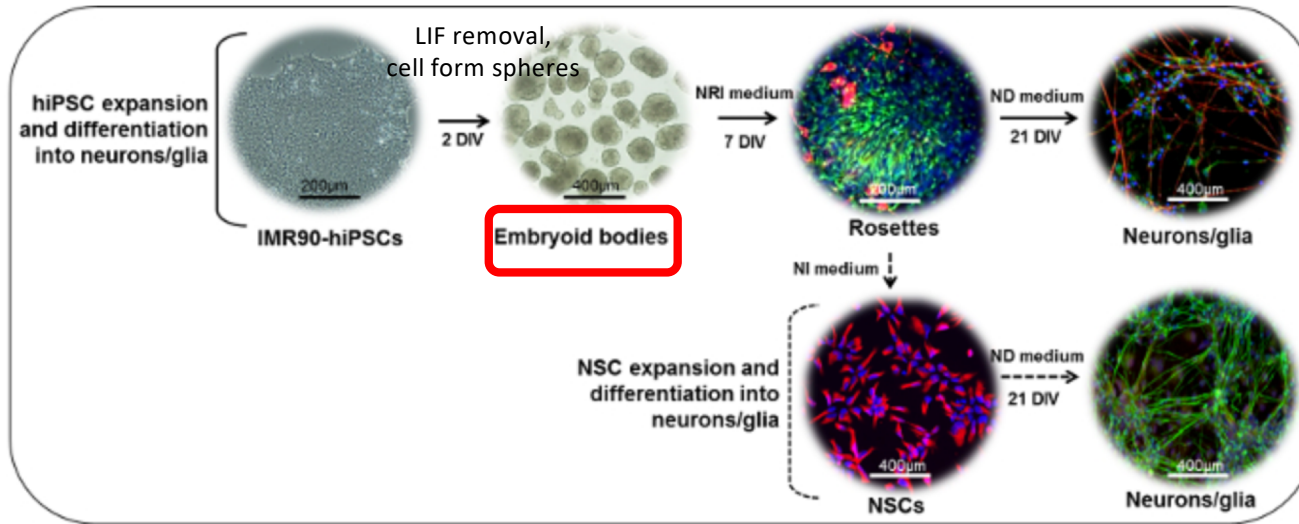
Ancestral OCT4 gave rise to 5 processed pseudogenes that are expressed in mESCs

4. EXPRESSION DURING IN VITRO DIFFERENTIATION

In vitro differentiation; 2D cell culture, attached on petri dishes

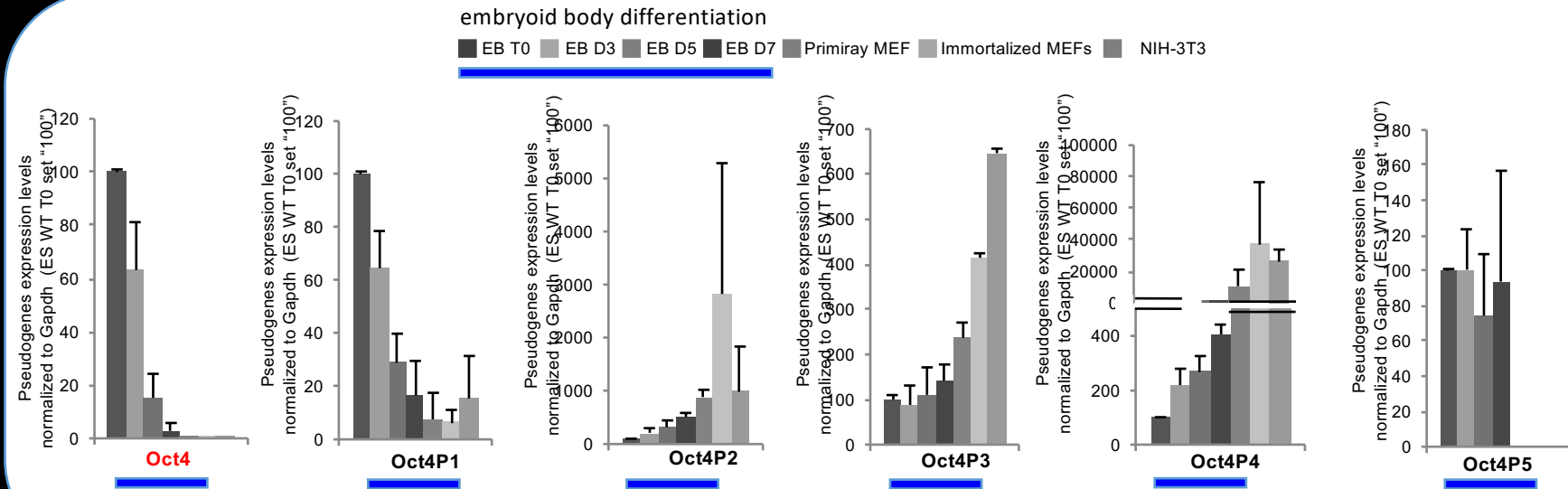


In vitro differentiation; 3D cell culture, embryoid body differentiation

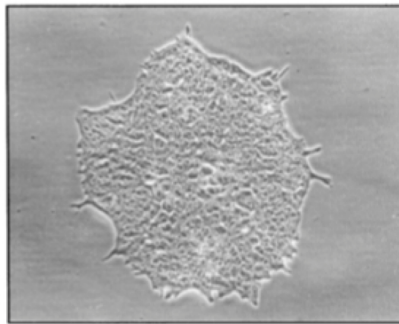


EB: embryoid body differentiation (a model for ES differentiation)

Oct4 pseudogenes are tightly controlled during the differentiation of mESCs

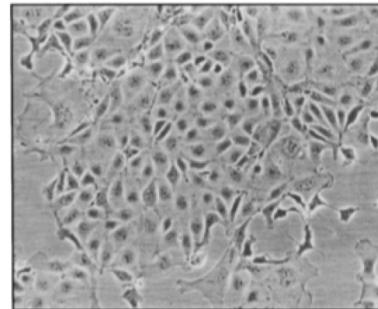


SELF-RENEWAL



Oct4
Oct4P1 (-10X)

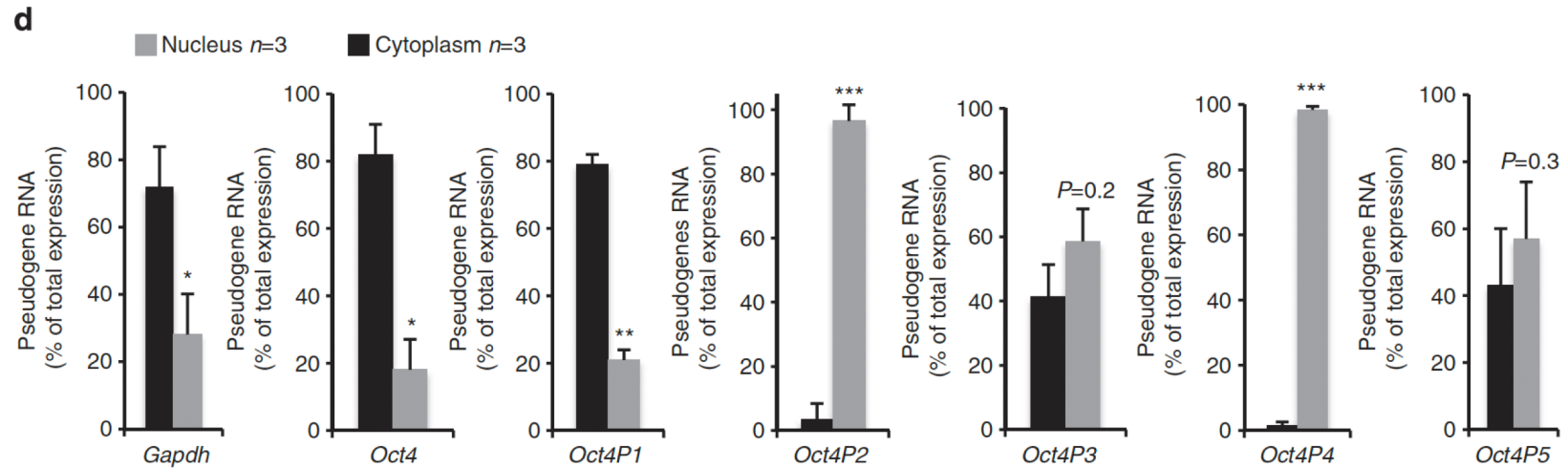
DIFFERENTIATION



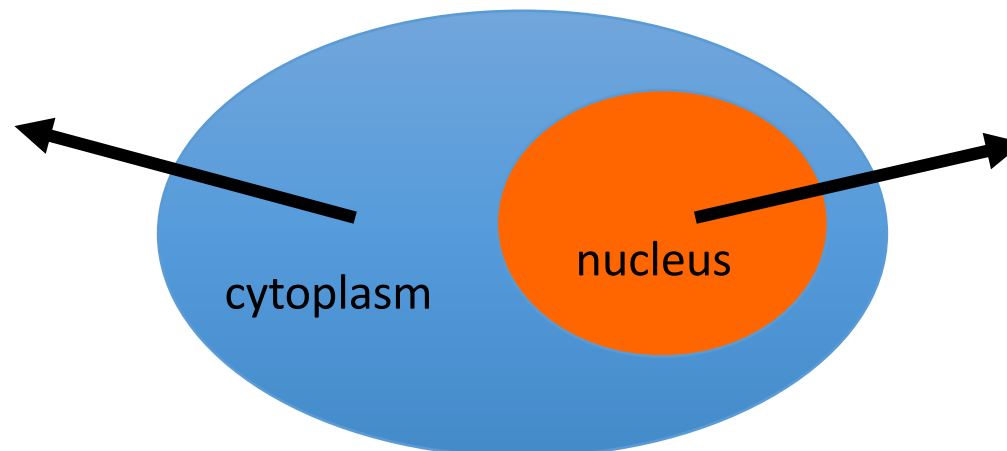
Oct4P2 (+9x)
Oct4P3 (+2x)
Oct4P4 (+4x; Fibrobl. +200x)

5. LOCALIZATION OF PSEUDOGENE lncRNA

OCT4 pseudogenes are localized to nucleoplasm or cytoplasm

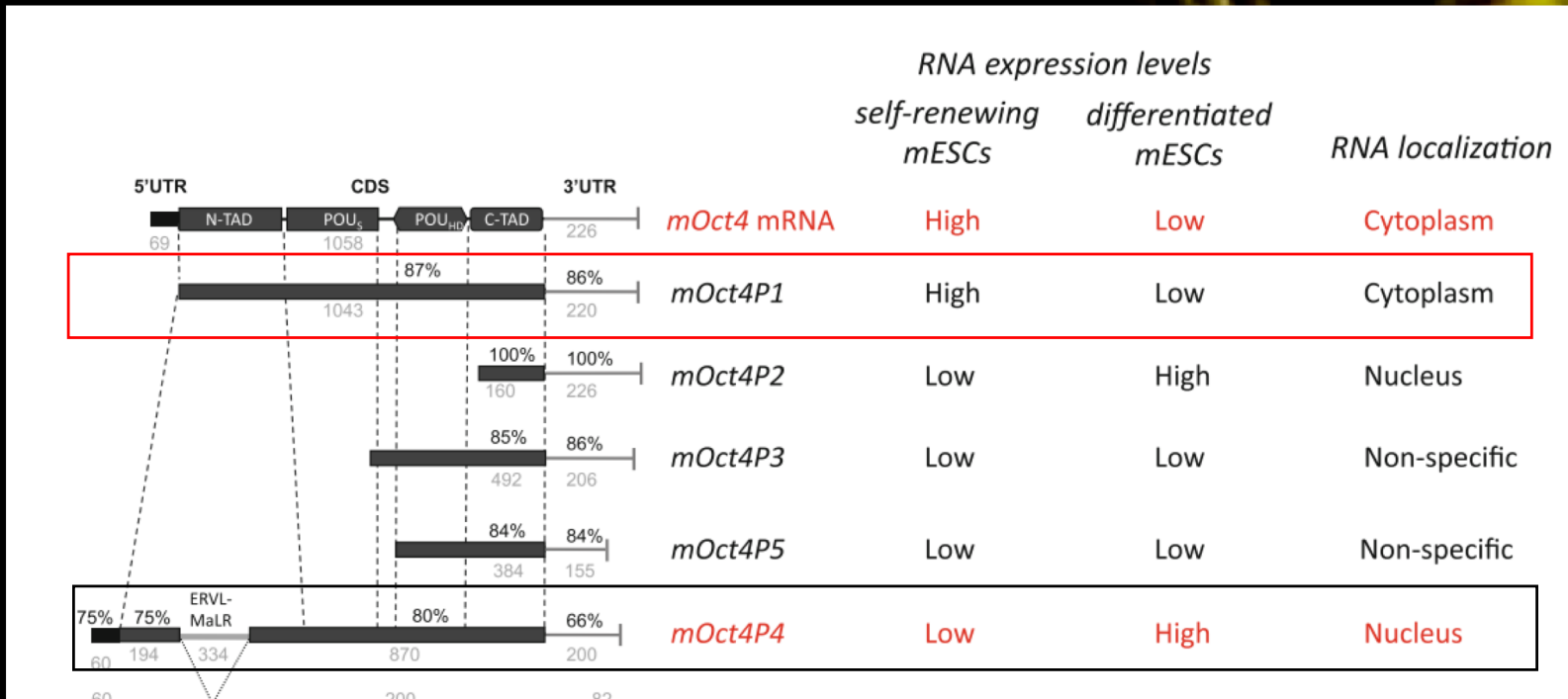


Oct4
Oct4P1

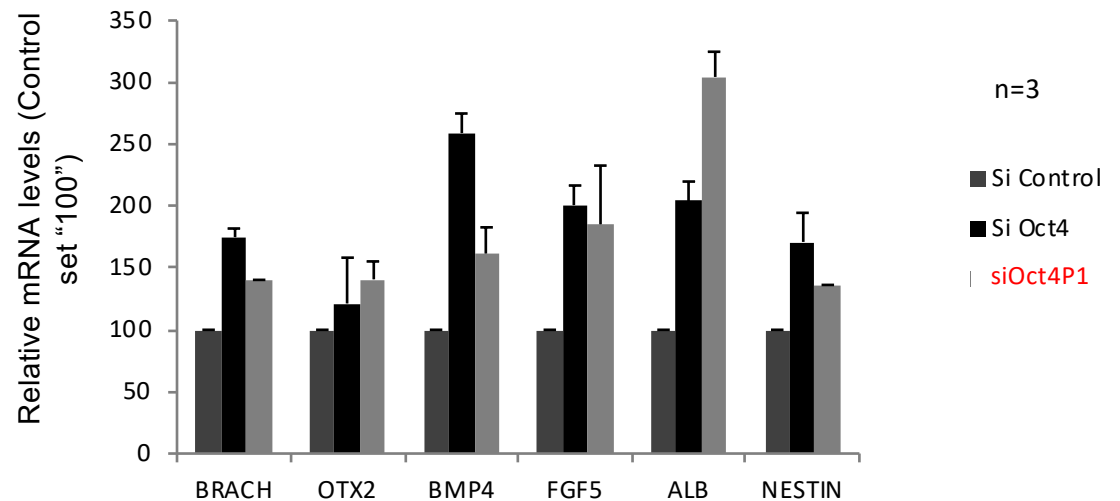
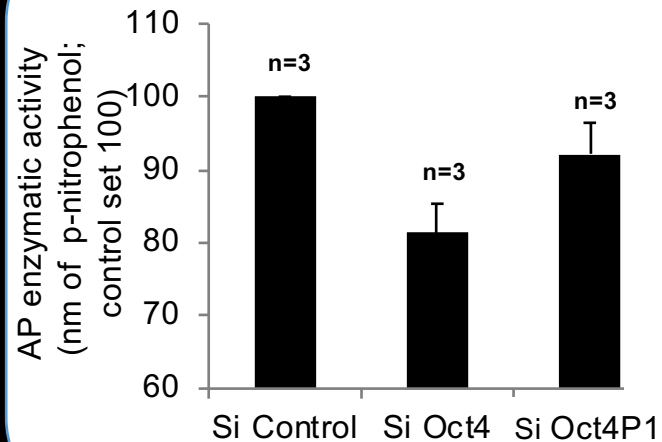
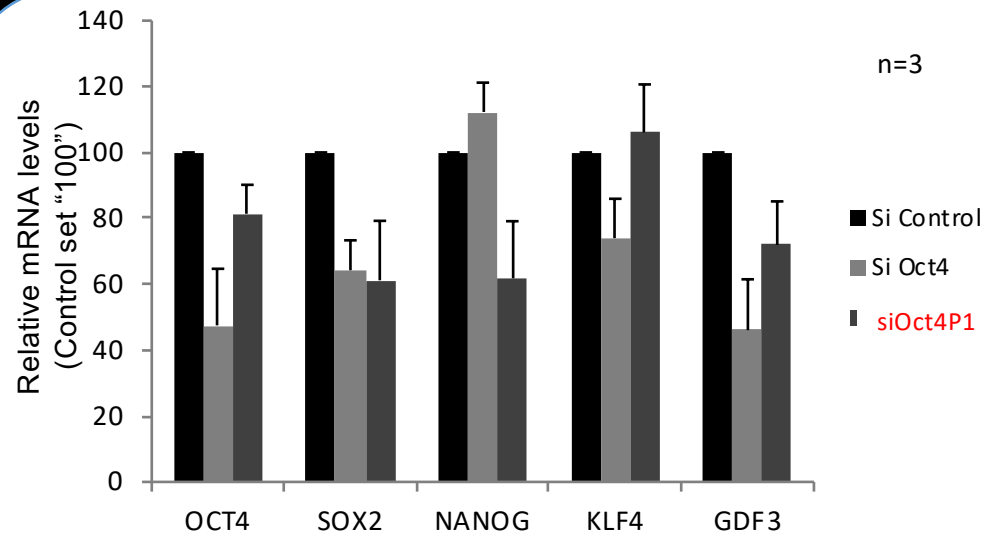
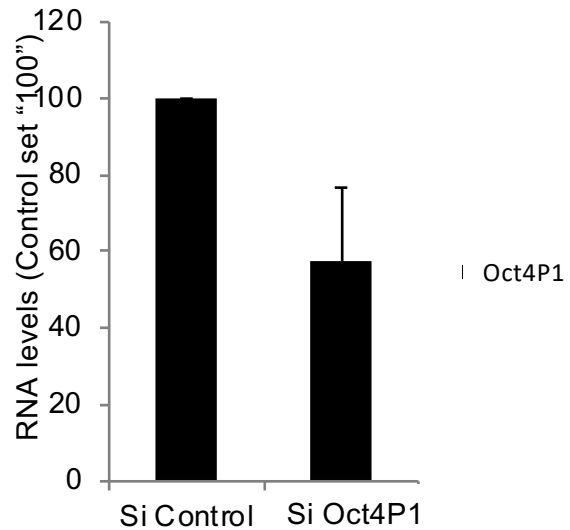


Oct4P2
Oct4P4

6. FUNCTIONAL ASSAY – lncRNA and self renewal

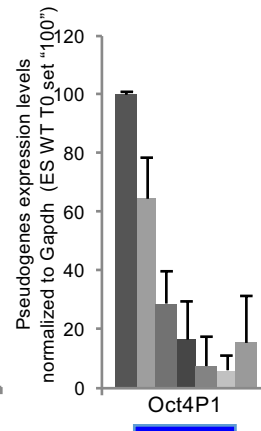
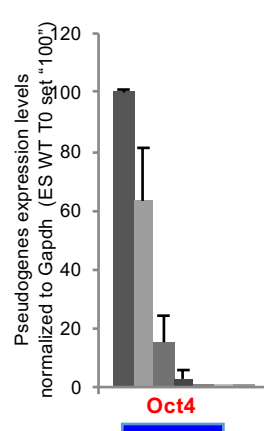
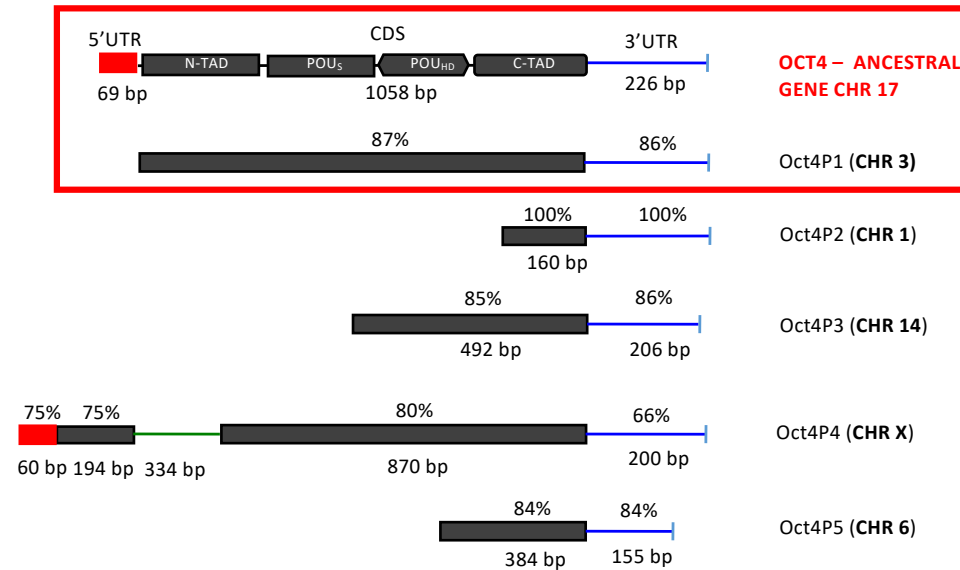


6. FUNCTIONAL ASSAY – lncRNA and self renewal



Cytoplasmatic OCT4P1 promotes mESC self-renewal

Ancestral OCT4 gave rise to 5 processed pseudogenes that are expressed in mESCs



siOct4: high in ESC

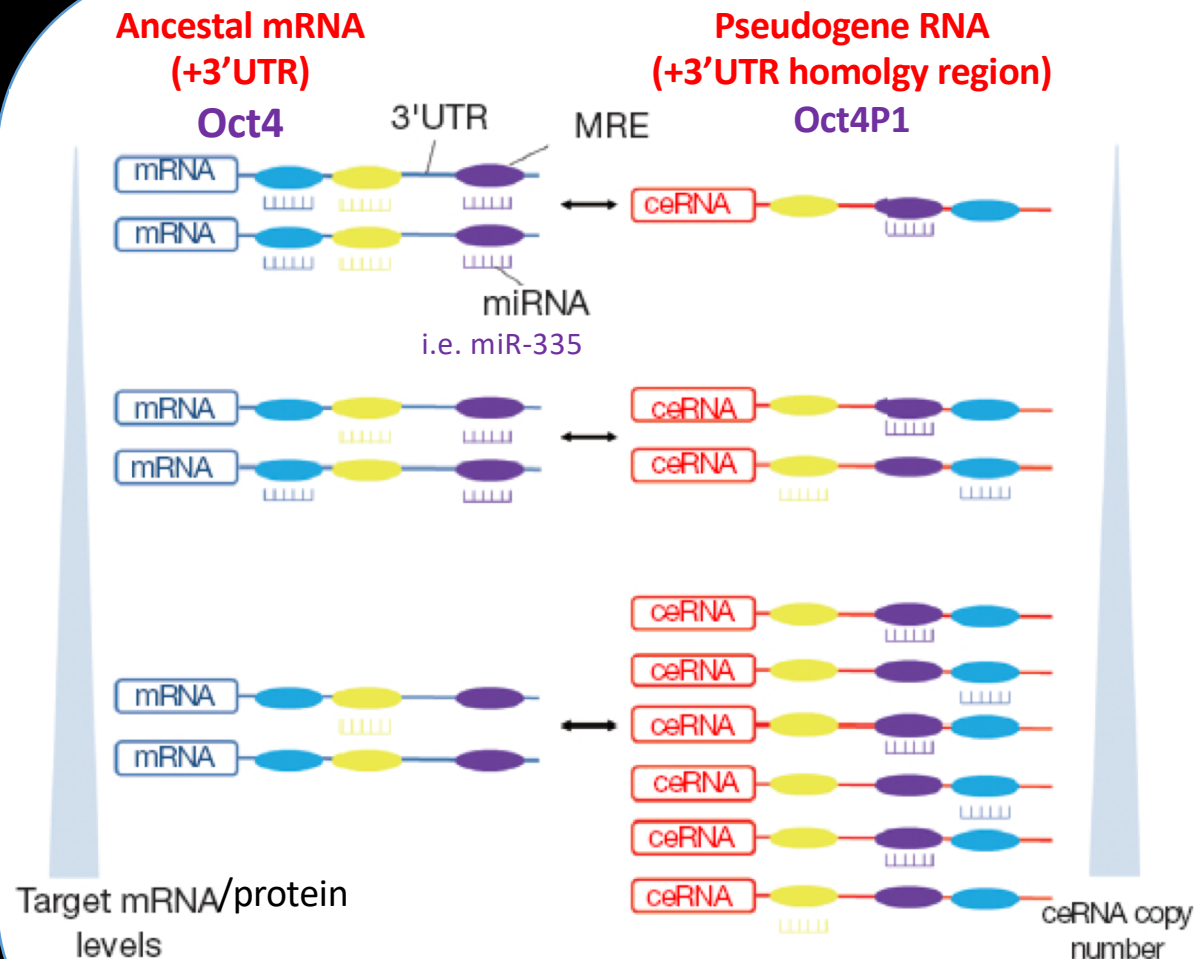
siOct4P1: high in ESC, low in differentiated cells

siOct4: differentiation

siOct4P1: differentiation

Oct4P1 needed to keep high Oct4 expression in self renewing ESC??

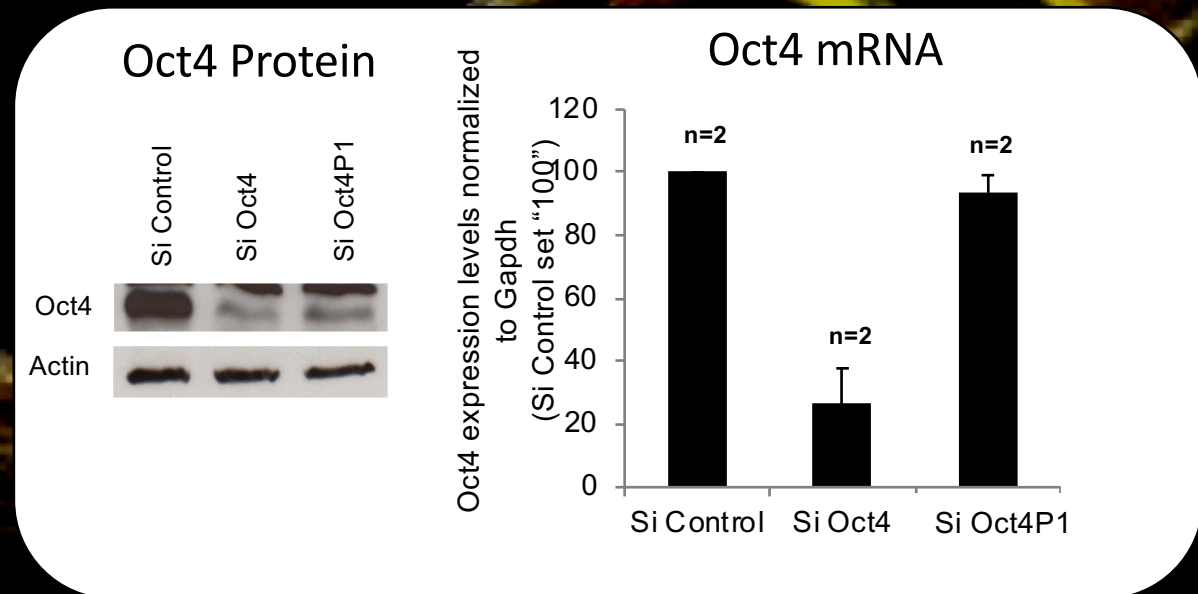
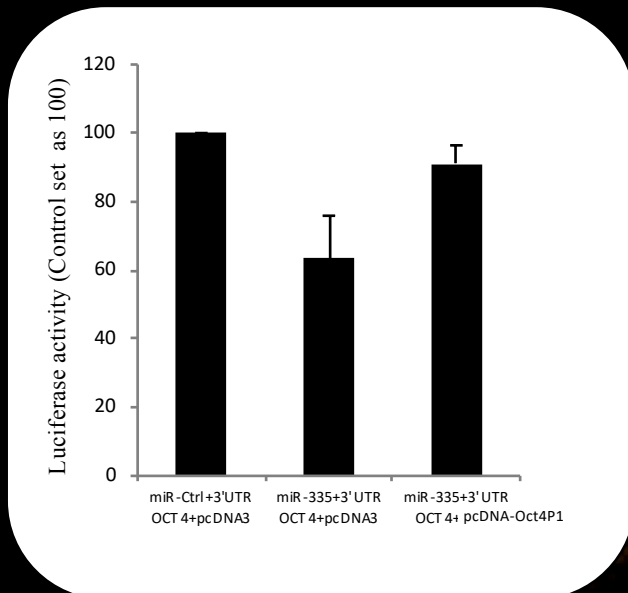
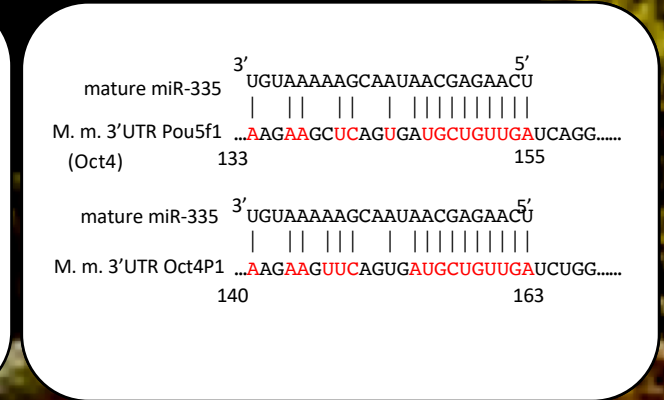
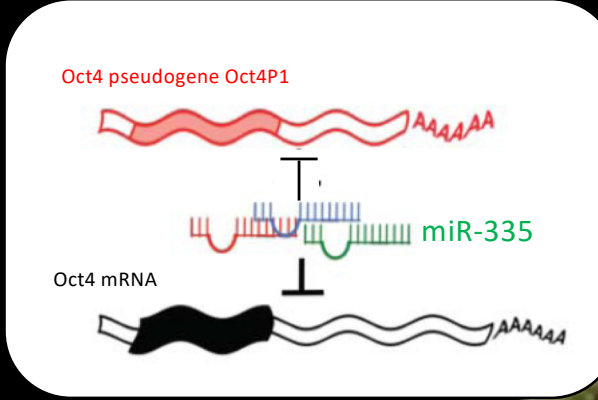
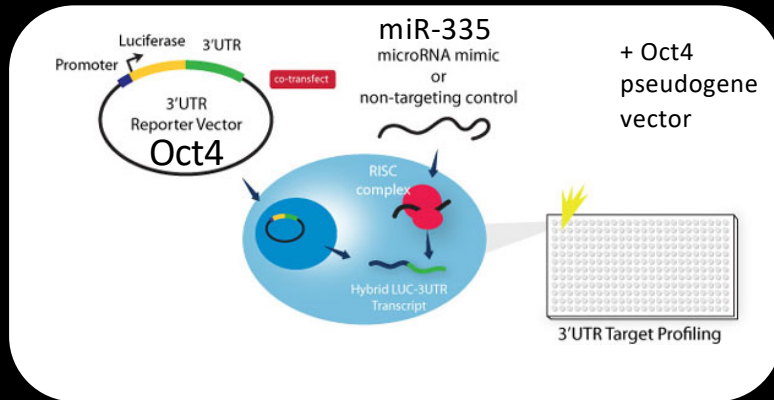
Pseudogene sponge miRNAs that target the ancestral gene



miRNA that targets Oct4: i.e. miR-335
MRE: miRNA response element

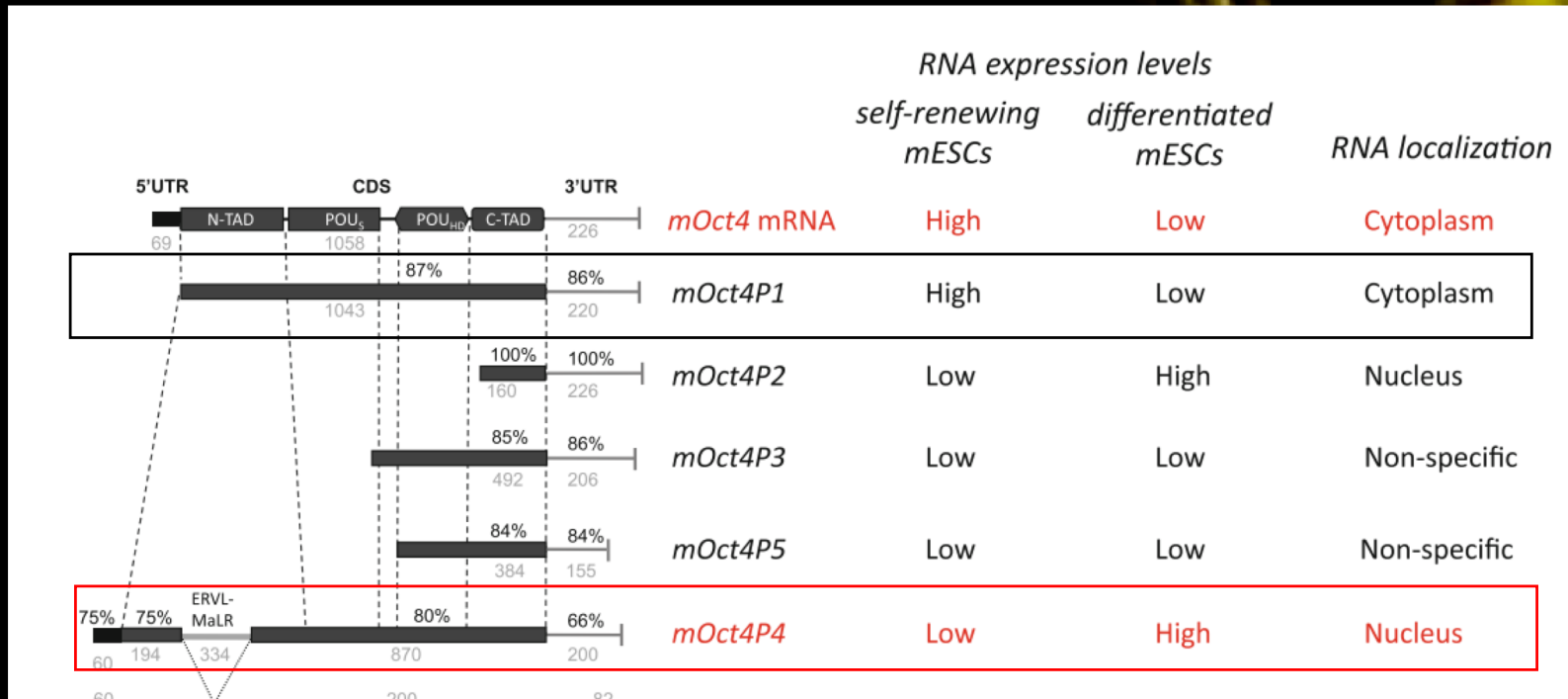
Evolution of
pseudogenes
to fine-tune the
expression
of ancestral genes

Cytoplasmatic OCT4P1 acts as Oct4/Rb1 ceRNA



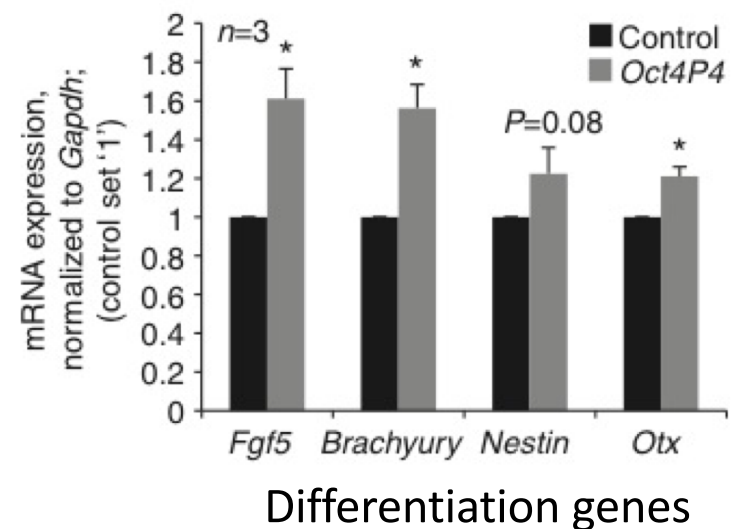
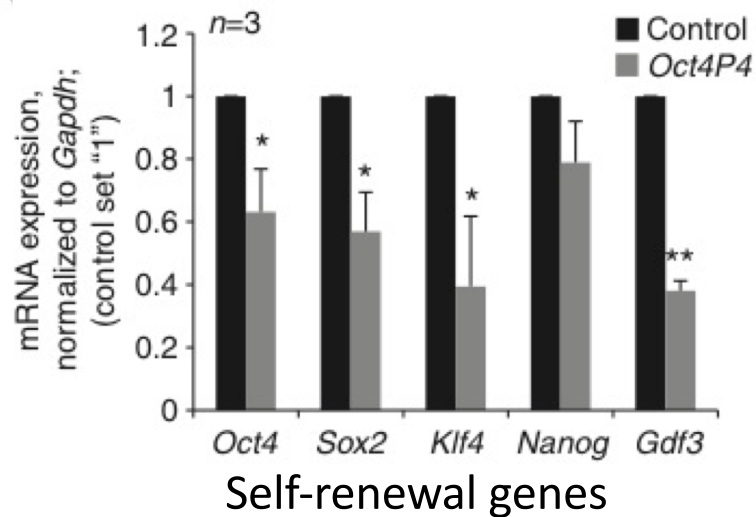
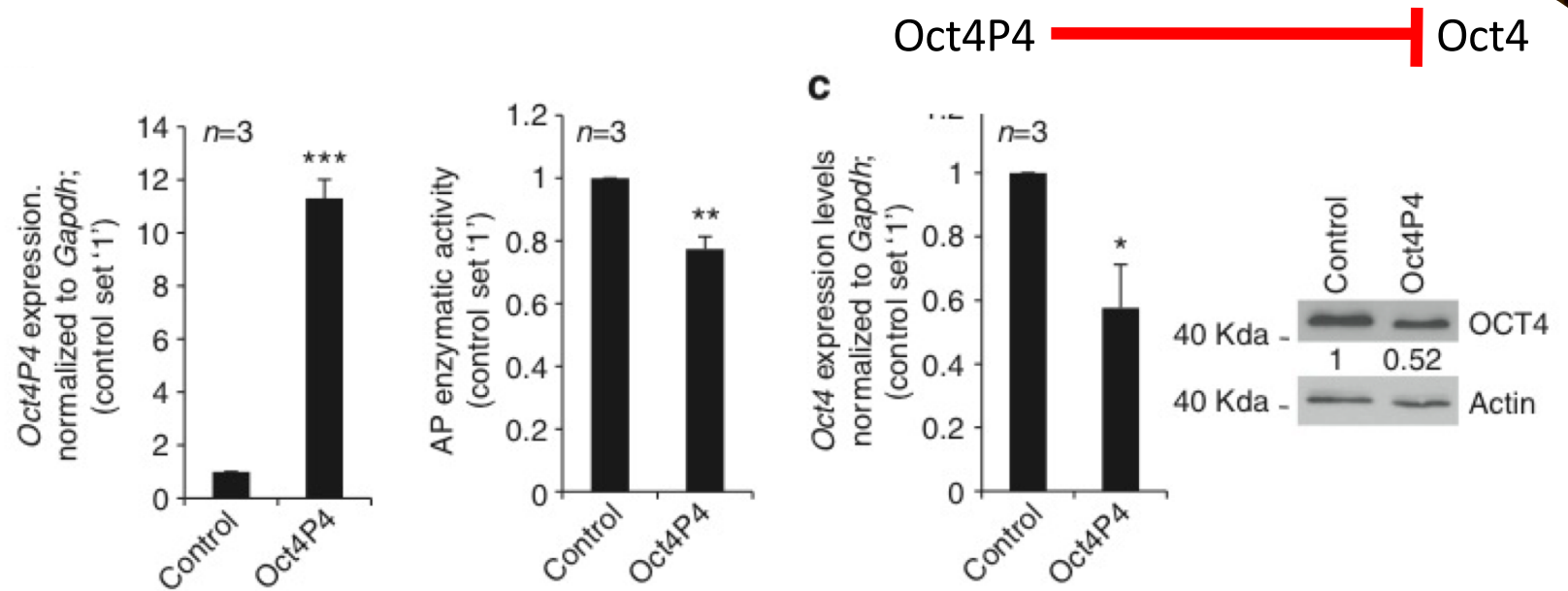
OCT4P1 promotes self-renewal by sponging miRNAs that target Oct4

6. FUNCTIONAL ASSAY – lncRNA and self renewal



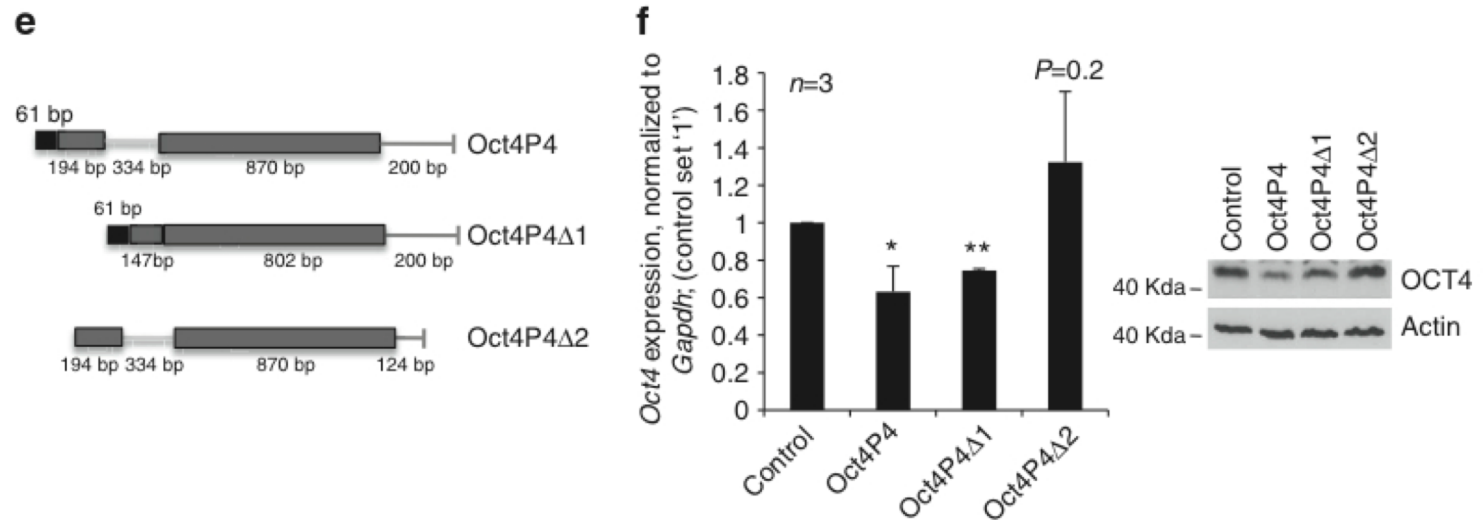
Overexpression of OCT4P4 down regulates Oct4 to promote mESC differentiation

mESCs with stable *mOCT4P4* expression construct

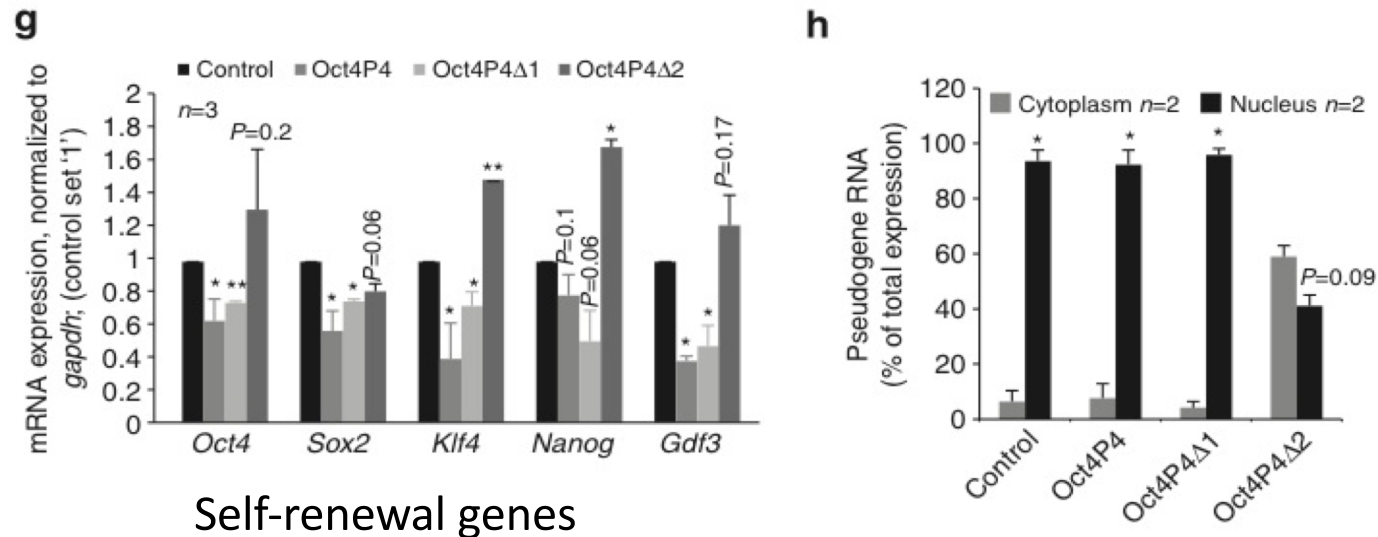


7. MOLECULAR ANALYSIS: WHAT ARE THE RELEVANT SEQUENCES??

Express Oct4P4 deletion constructs and follow parental gene expression



Express Oct4P4 deletion constructs and follow self-renewal programs



5' and 3' UTR homology regions are required to repress self-renewal marker genes

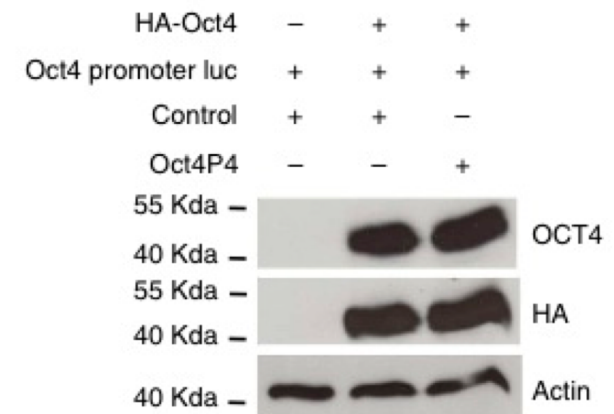
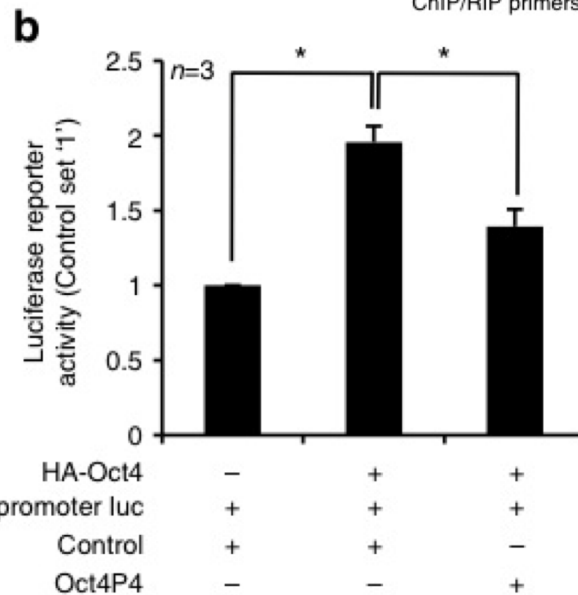
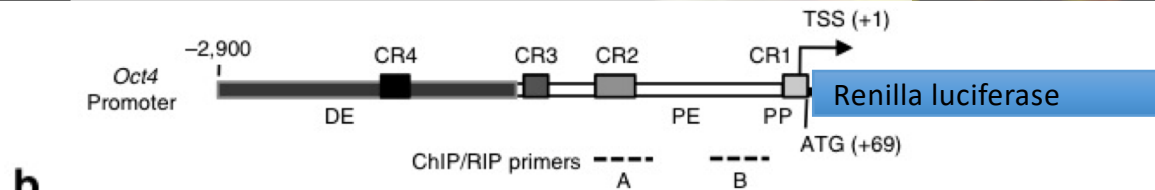
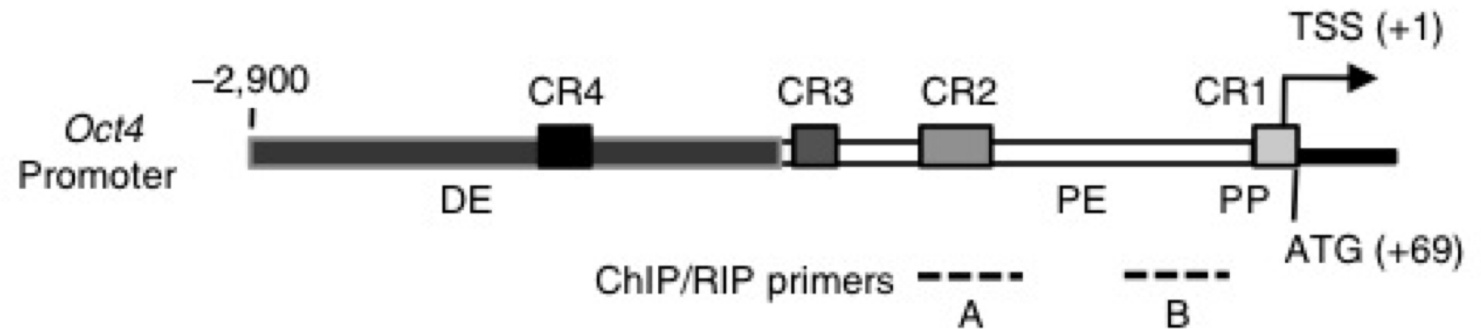
8. MOLECULAR ANALYSIS: USE SIMPLIFIED MODEL TO GET INDICATION ON MOLECULAR MECHANISM??

Oct4P4 is a nuclear RNA that suppresses Oct4 transcription

Hypothesis:
transcriptional suppression of Oct4 promoter?

Use simplified models to support hypothesis:

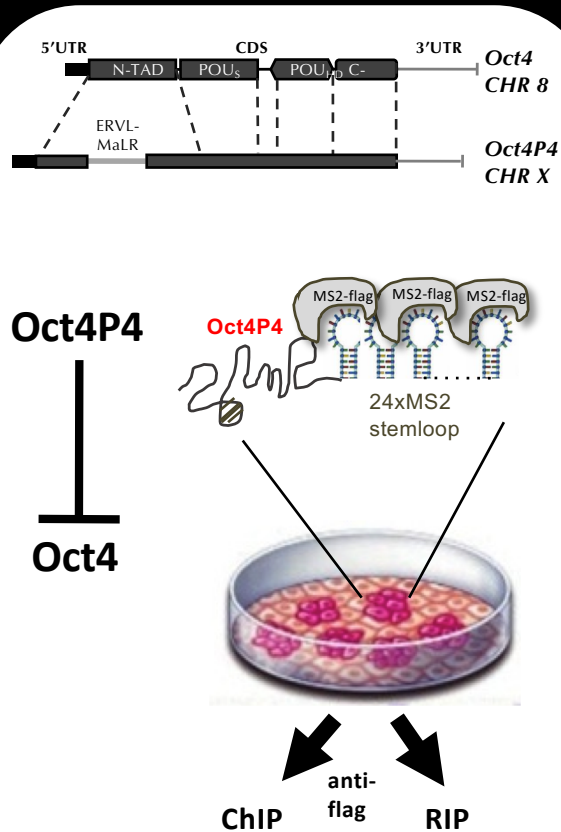
- Clone Oct4 promoter
- Clone luc reporter downstream
- Test whether Oct4P4 lncRNA suppresses reporter



Oct4P4 may interfere with the ancestral Oct4 promoter

9. MOLECULAR ANALYSIS: DEMONSTRATE MECHANISM AT ENDOGENOUS Oct4 PROMOTER??

- Generate model system to:
1. study Oct4P4 lncRNA localization
 2. study chromatin at OCT4 promoter

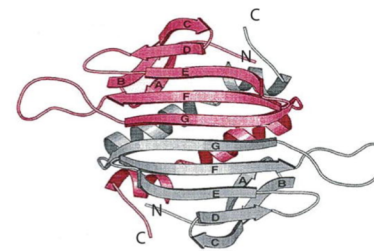


Step 1: stably transfect mESCs with MS2-flag

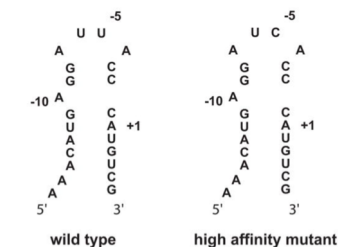
Step 2: stably transfect mESCs with an expression vector encoding mOct4P4 lncRNA fused to 24 MS loop stem loops



Diagram of MS2 coat protein dimer



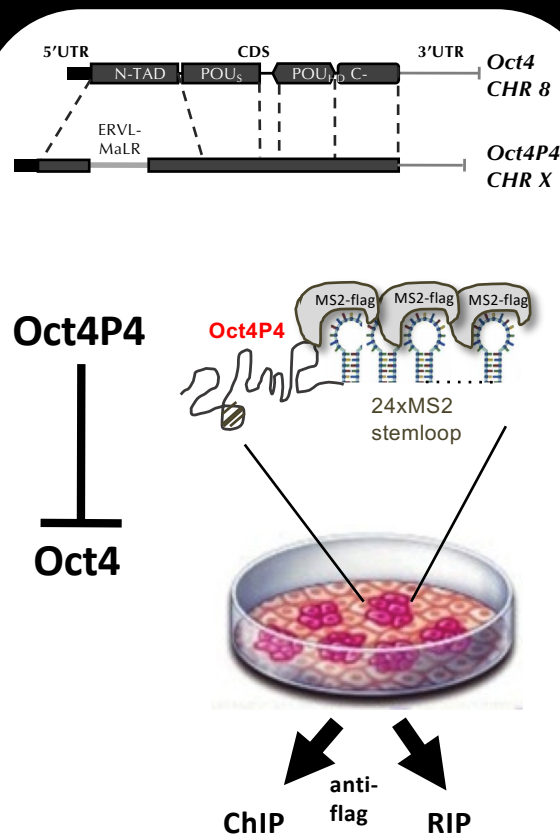
MS2 stem loop sequence



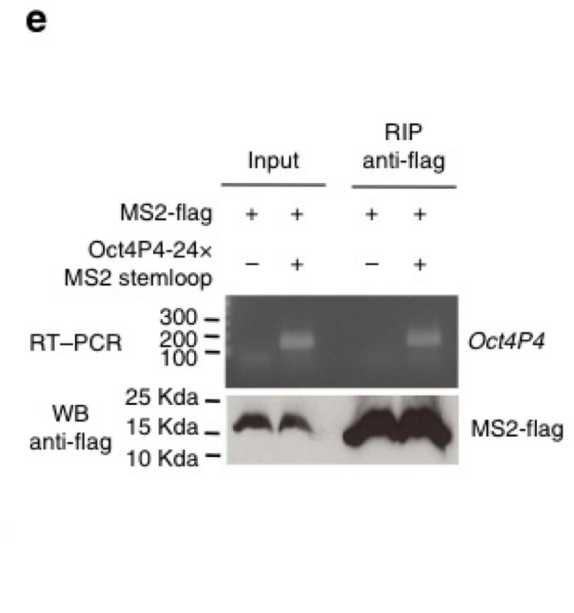
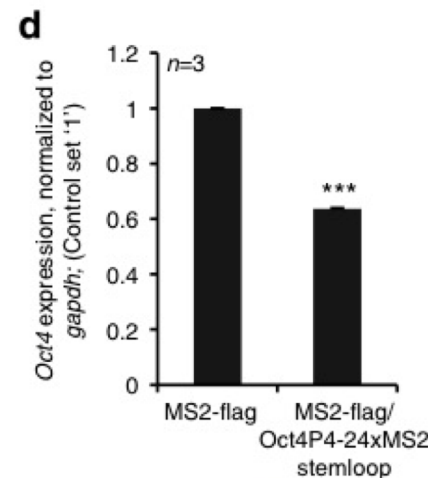
MS2 stem loop tagged Oct4P4 co-expressed with flag-MS2 in mESCs

9. MOLECULAR ANALYSIS: DEMONSTRATE MECHANISM AT ENDOGENOUS Oct4 PROMOTER??

- Generate model system to:
1. study Oct4P4 lncRNA localization
 2. study chromatin at OCT4 promoter

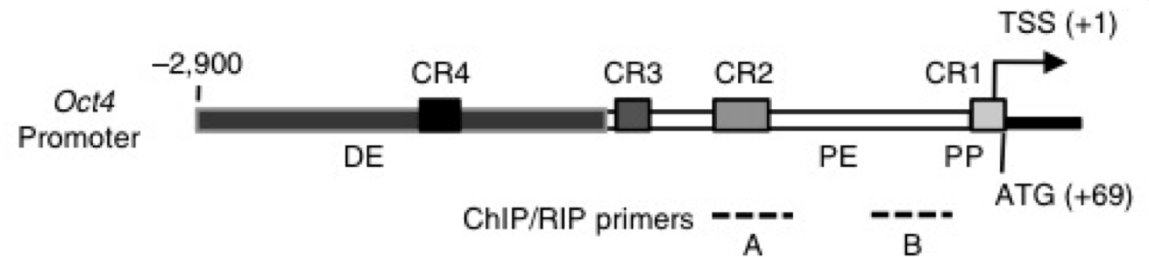
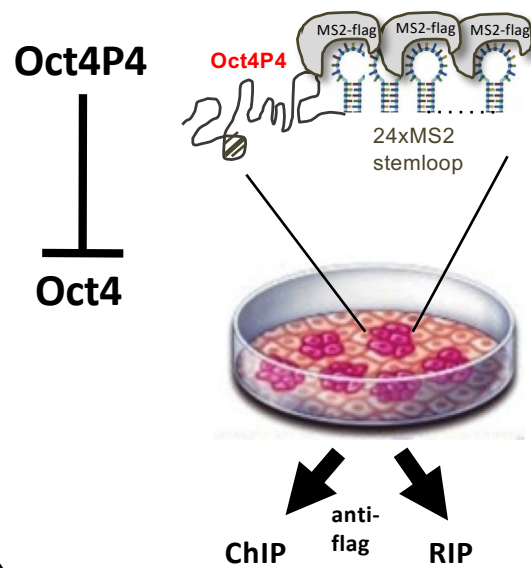
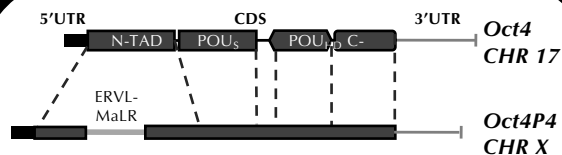


RNA immunoprecipitation anti-flag then RT-PCR for Oct4P4 lncRNA

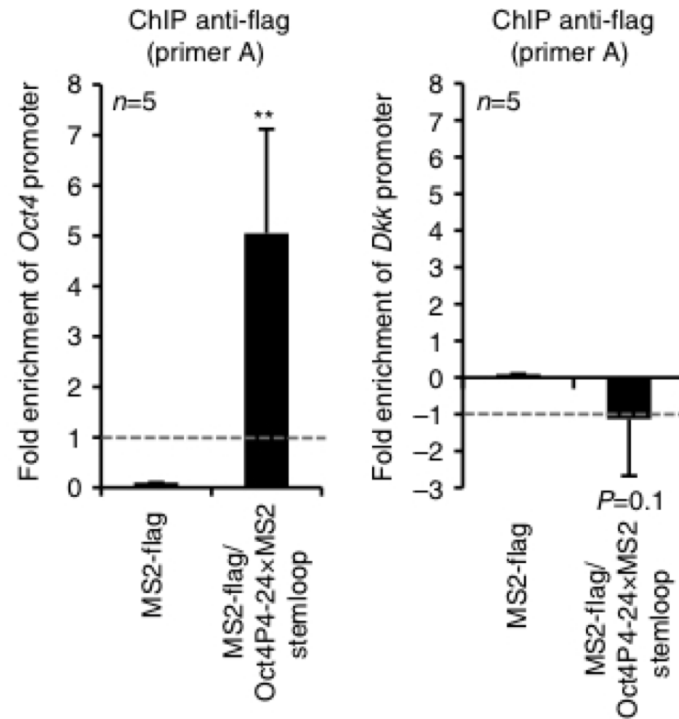


MS2 stem loop tagged Oct4P4 co-expressed with flag-MS2 in mESCs

A model system to study Oct4P4 lncRNA localization



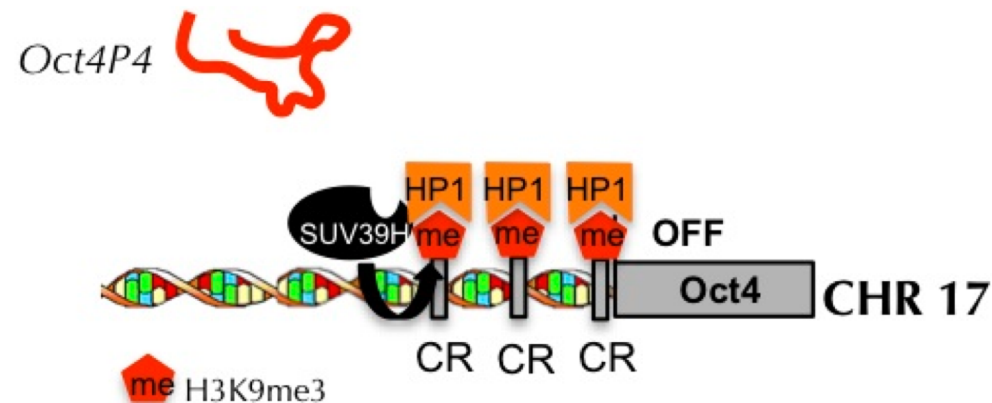
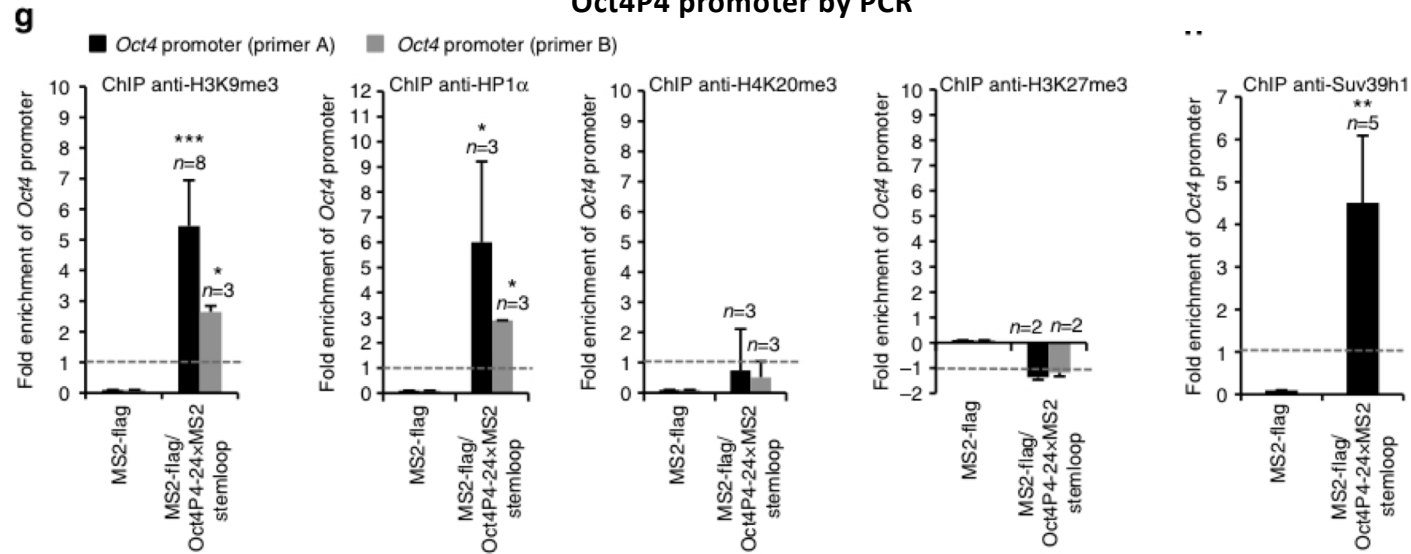
ChIP using anti-flag then use the immuno-precipitate to detect the Oct4P promoter by PCR



Oct4P4-MS2 lncRNA localizes to Oct4 promoter

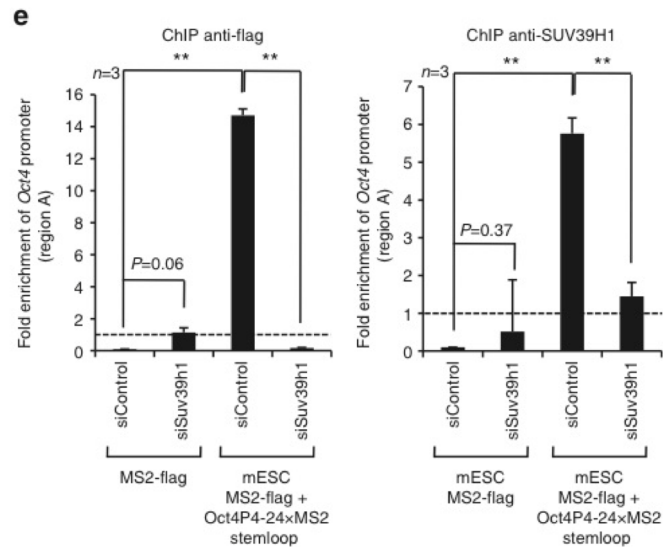
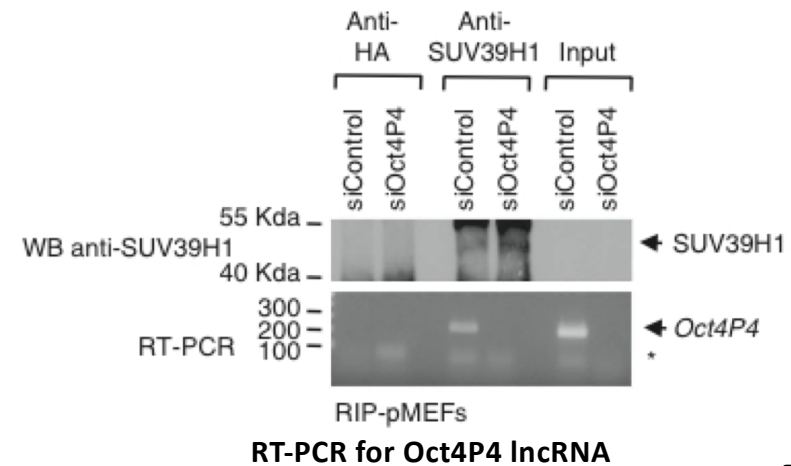
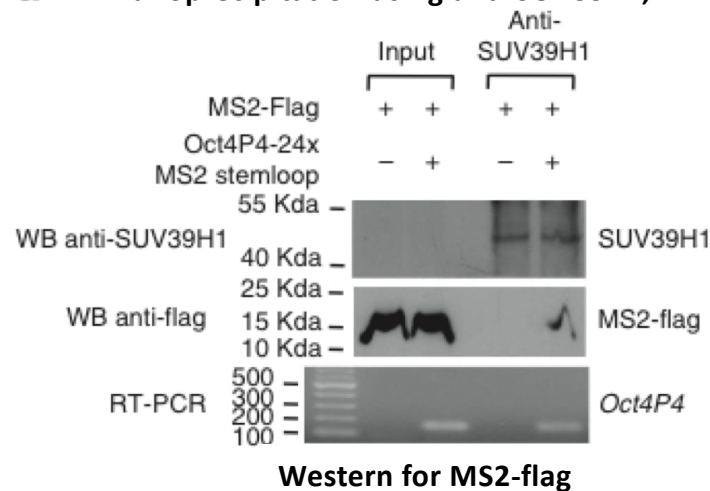
Oct4P4-MS2 directs Suv39h1 to Oct4 promoter

ChIP using specific antibodies then use the immuno-precipitate to detect the Oct4P4 promoter by PCR

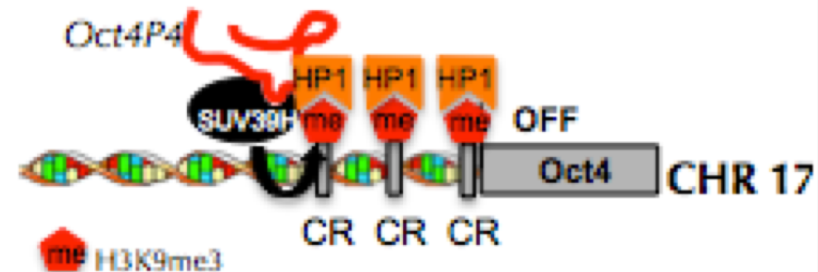


Oct4P4-MS2 directly interacts with Suv39h1

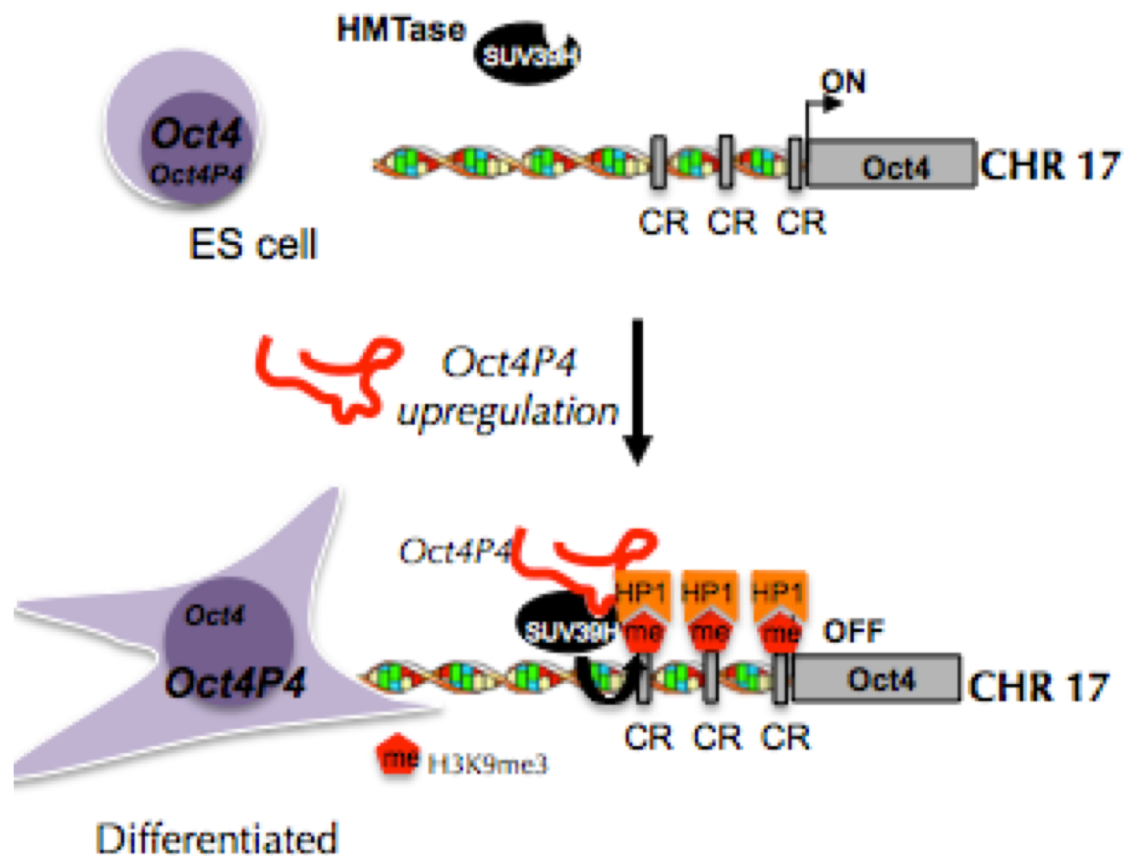
RNA immunoprecipitation using anti-SUV39h1;



ChIP using specific antibodies then use the immuno-precipitate to detect the Oct4P4 promoter by PCR



Oct4P4-MS2 recruits Suv39h1 To direct silencing of the Oct4 promoter

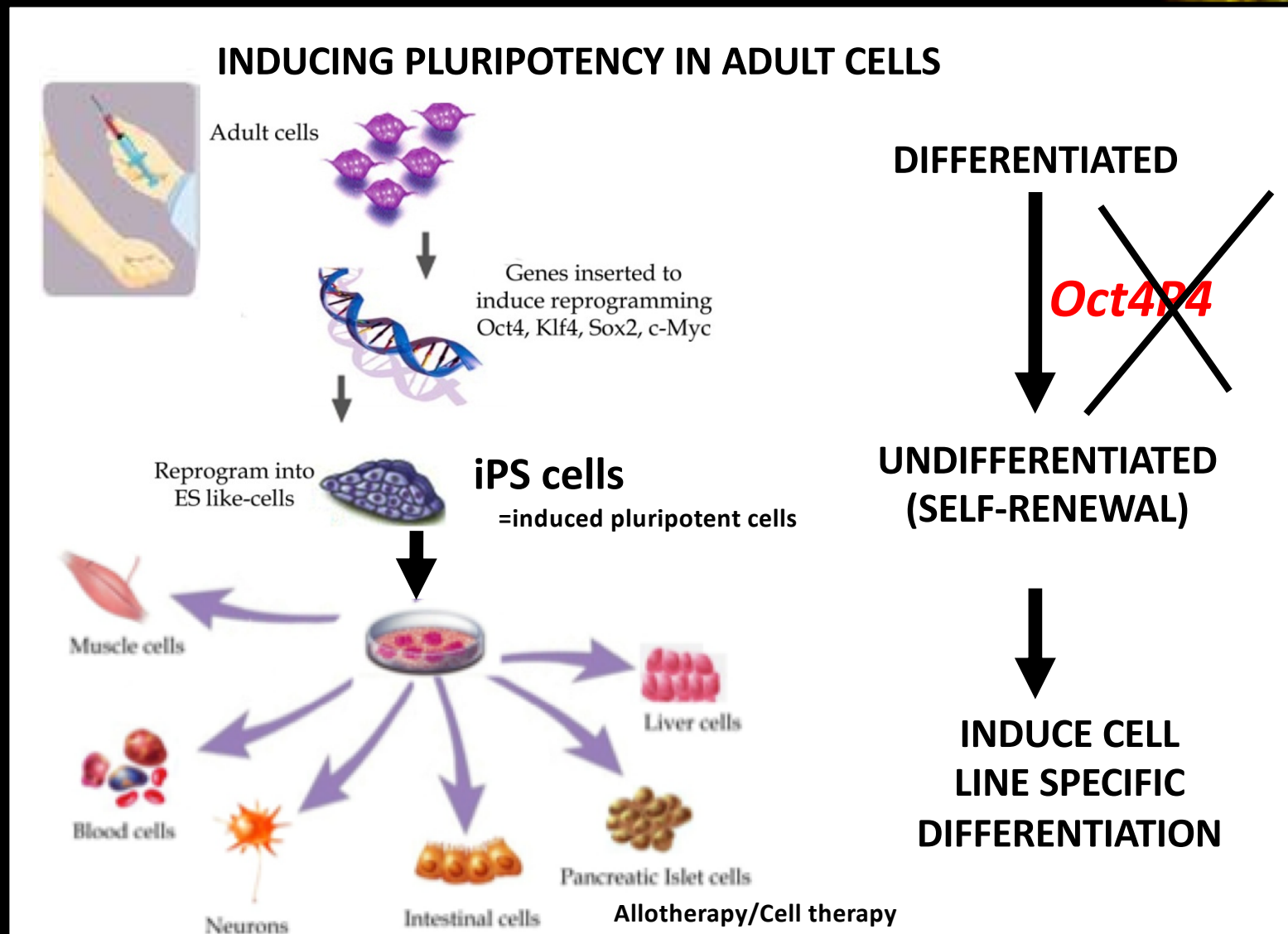


FUNCTIONAL
RELEVANCE????

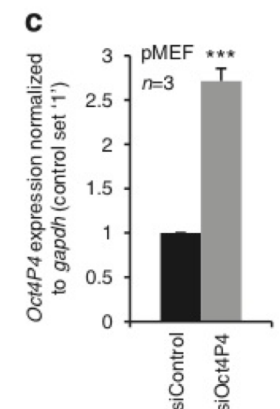
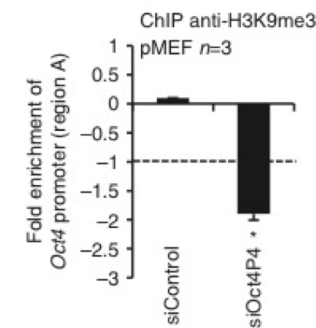
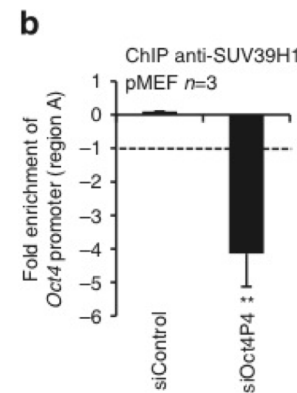
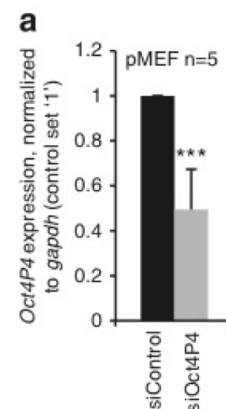
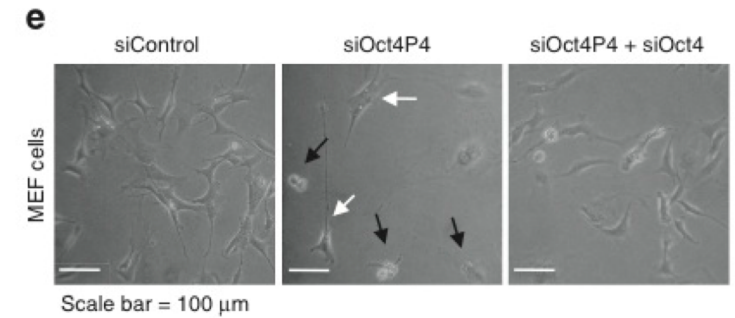
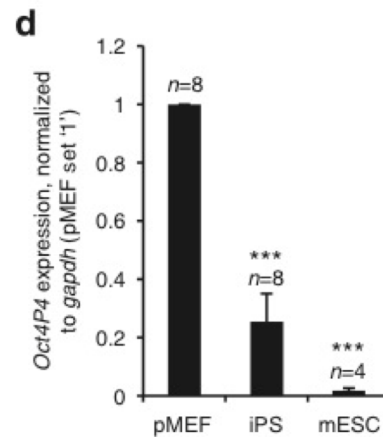
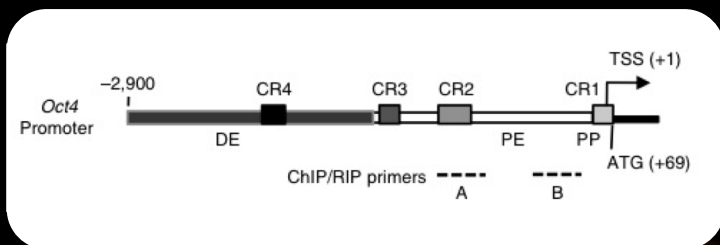
REVERSIBILITY???

10. Functional relevance:

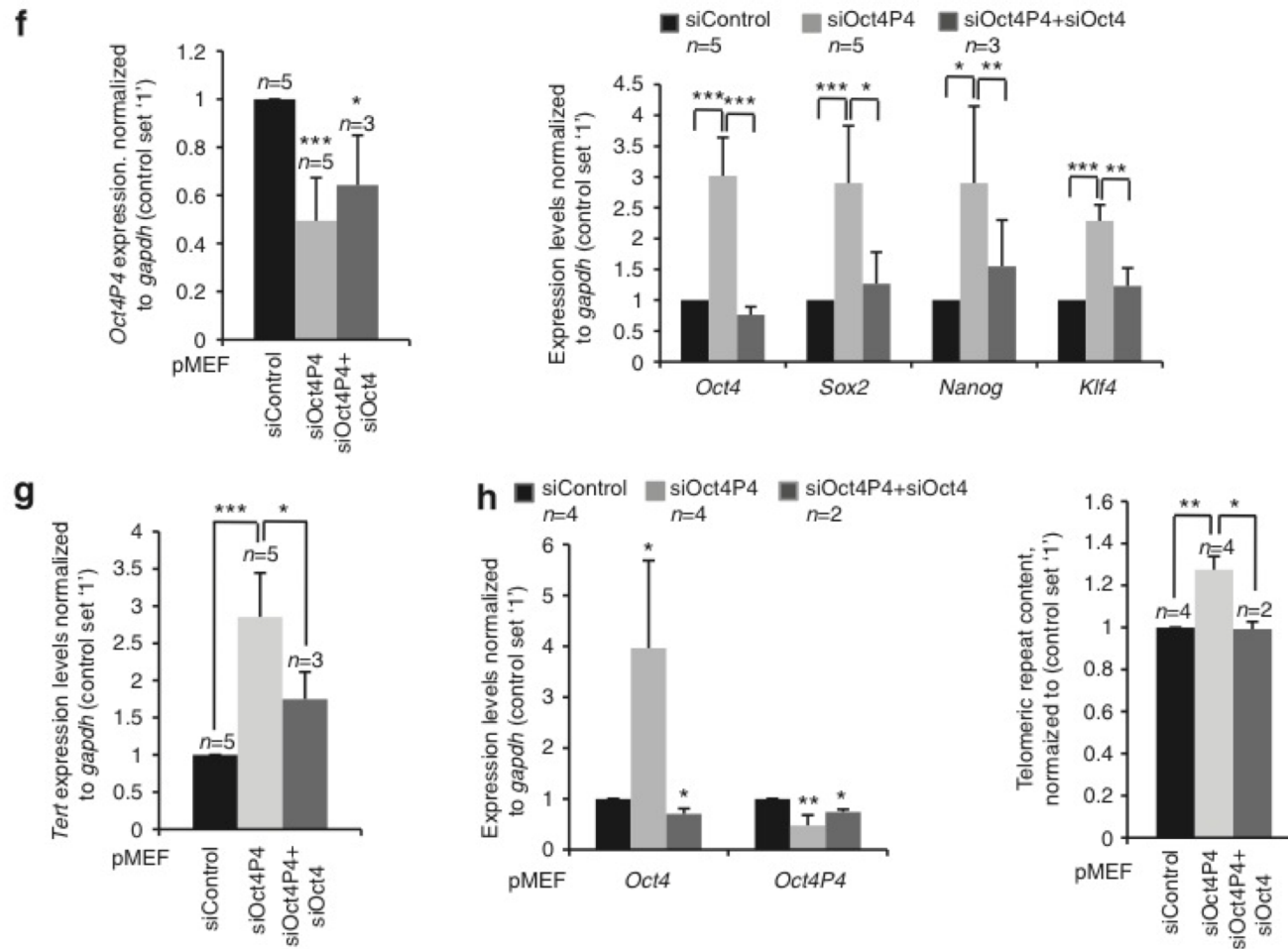
Oct4P4 depletion in pMEFs causes the re-acquisition of self-renewal features



Oct4P4 depletion in pMEFs causes the re-acquisition of self-renewal features

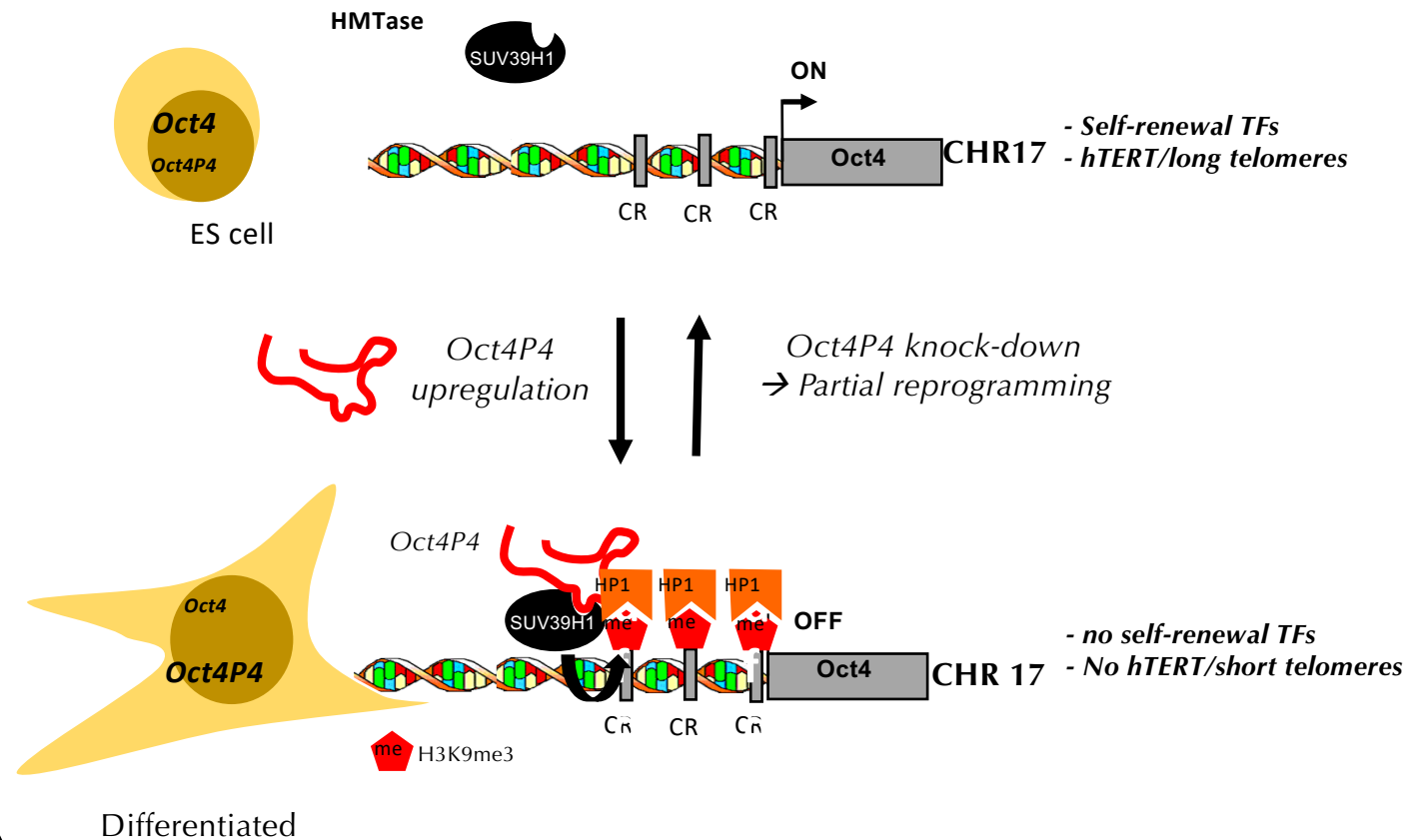


Oct4P4 depletion in pMEFs causes the re-acquisition of self-renewal features



Pseudogenes control the epigenetic status of ancestral genes

Oct4 pseudogene lncRNA silences ancestral gene

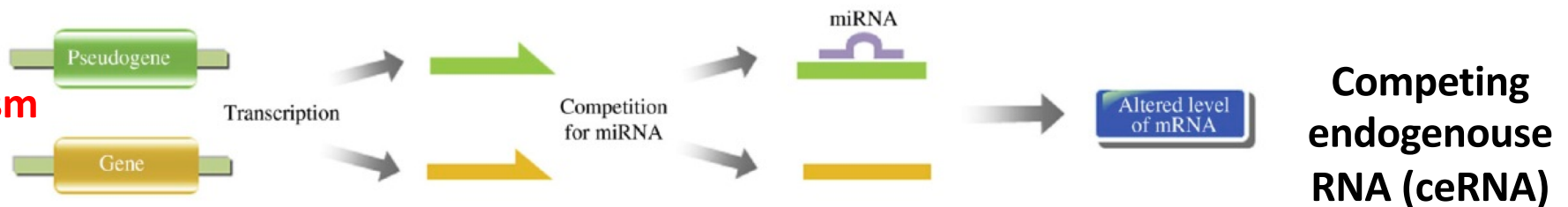


Pseudogenes are powerful regulators of gene expression

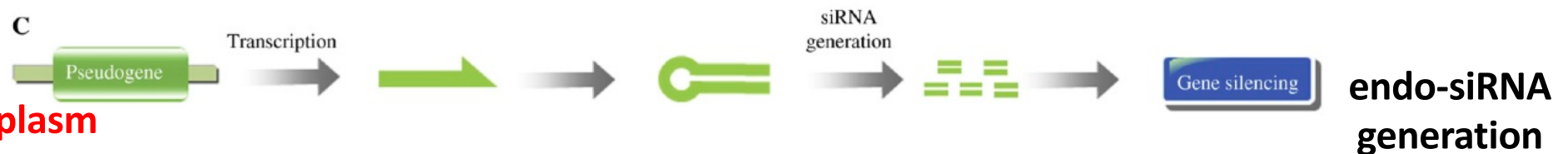
A
Nuclear



B
Cytoplasm



C
Cytoplasm



D
Cytoplasm

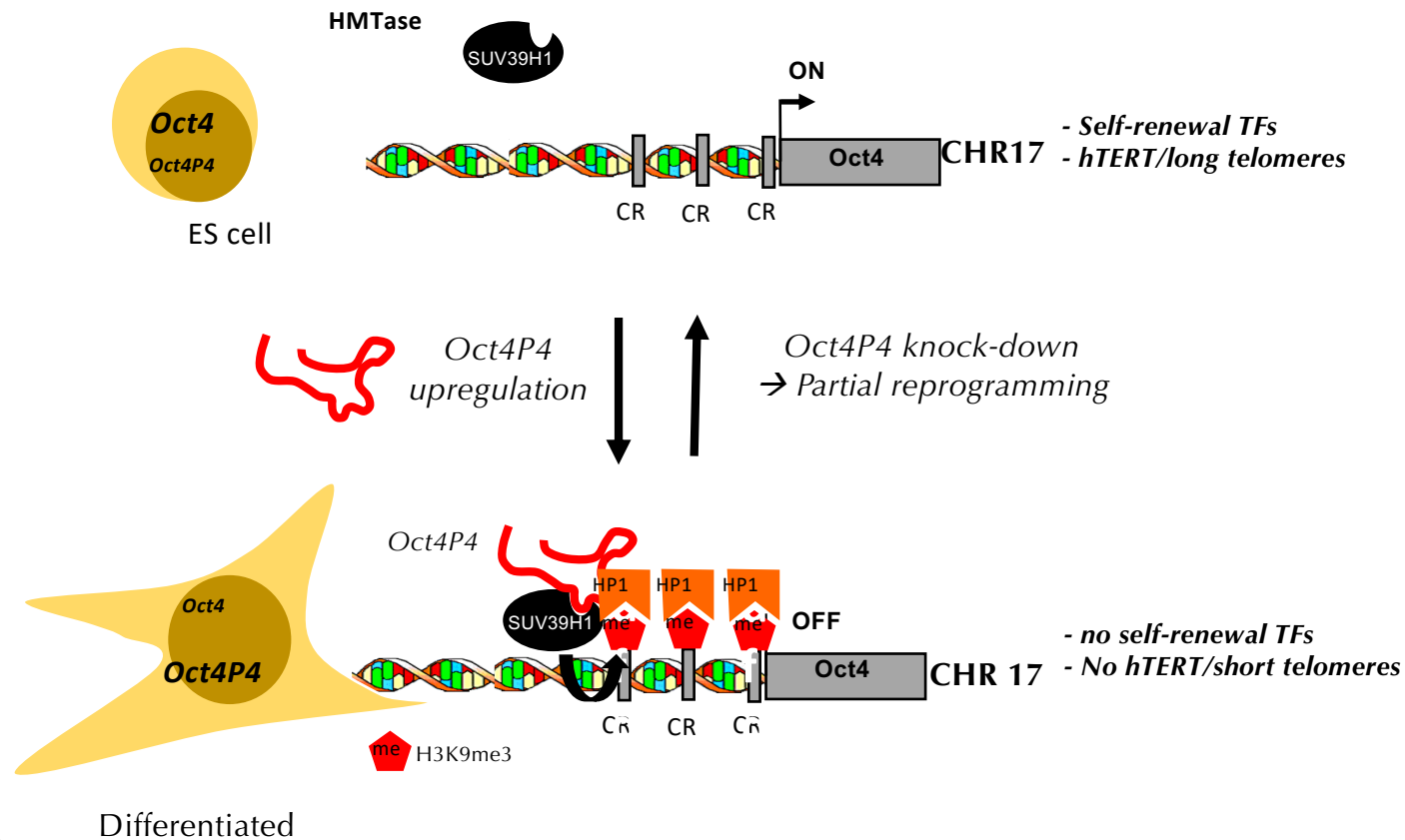


Nuclear



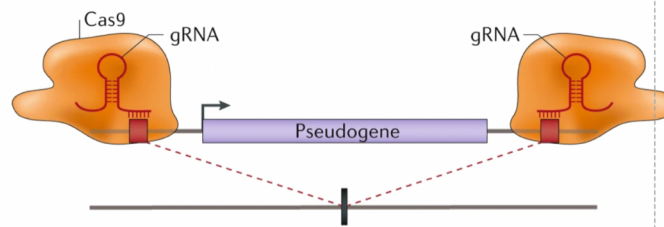
WHAT WOULD YOU DO NEXT TO UNDERSTAND OCT4P4 FUNCTION?

Oct4 pseudogene lncRNA silences ancestral gene

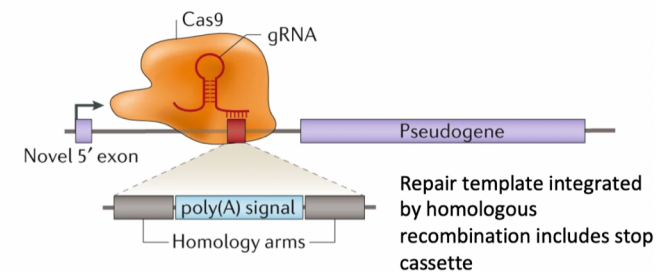


- GENETIC MODEL SYSTEM
- FUNCTIONAL RELEVANT REGIONS
- INTERACTING PROTEINS

c CRISPR-mediated deletion

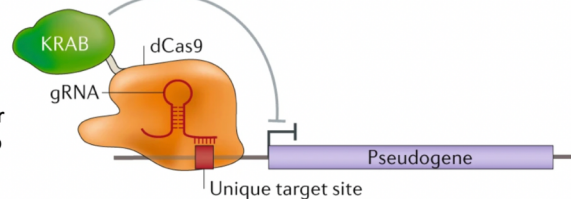


d Integration of transcriptional terminator



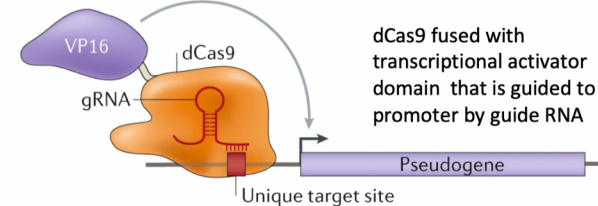
e CRISPR interference

dCas9 fused with transcriptional repressor domain that is guided to promoter by guide RNA



f CRISPR activation

dCas9 fused with transcriptional activator domain that is guided to promoter by guide RNA



mESCs

	WT	dCas9 empty	dCas9 <i>sgOct4P4</i>	
170 kDa —				WB
130 kDa —				HA-dCAS9
40 kDa —				ACTIN
100bp —				RT-PCR
				<i>sgOct4P4</i>
100bp —				<i>Gapdh</i>

Stable expression of dCAS9 and *sgOct4P4* promoter

e CRISPR interference

dCas9 fused with transcriptional repressor domain that is guided to promoter by guide RNA

Pseudogene

Unique target site

Bar graph showing mOct4P4 expression, normalized to Gapdh; (dCas9 empty EB T0 set 4.1^{th}) for dCas9 empty (white bars) and dCas9 sgOct4P4 (black bars) at T0, D3, and D10. The y-axis represents mOct4P4 expression, normalized to Gapdh; (dCas9 empty EB T0 set 4.1^{th}), ranging from 0 to 50. The x-axis shows time points T0, D3, and D10. Sample sizes (n) are indicated above each bar. P-values for comparisons between dCas9 empty and dCas9 sgOct4P4 are shown: $p=0.2821$ at T0 and $p=0.0013$ at D3.

Time Point	dCas9 empty (n)	dCas9 sgOct4P4 (n)	Approx. Expression
T0	3	3	~1.0
D3	3	3	~25.0 (empty), ~4.0 (sgOct4P4)
D10	2	2	~38.0 (empty), ~7.0 (sgOct4P4)

Figure 2: Expression levels of pluripotency markers in EB D6 cells.

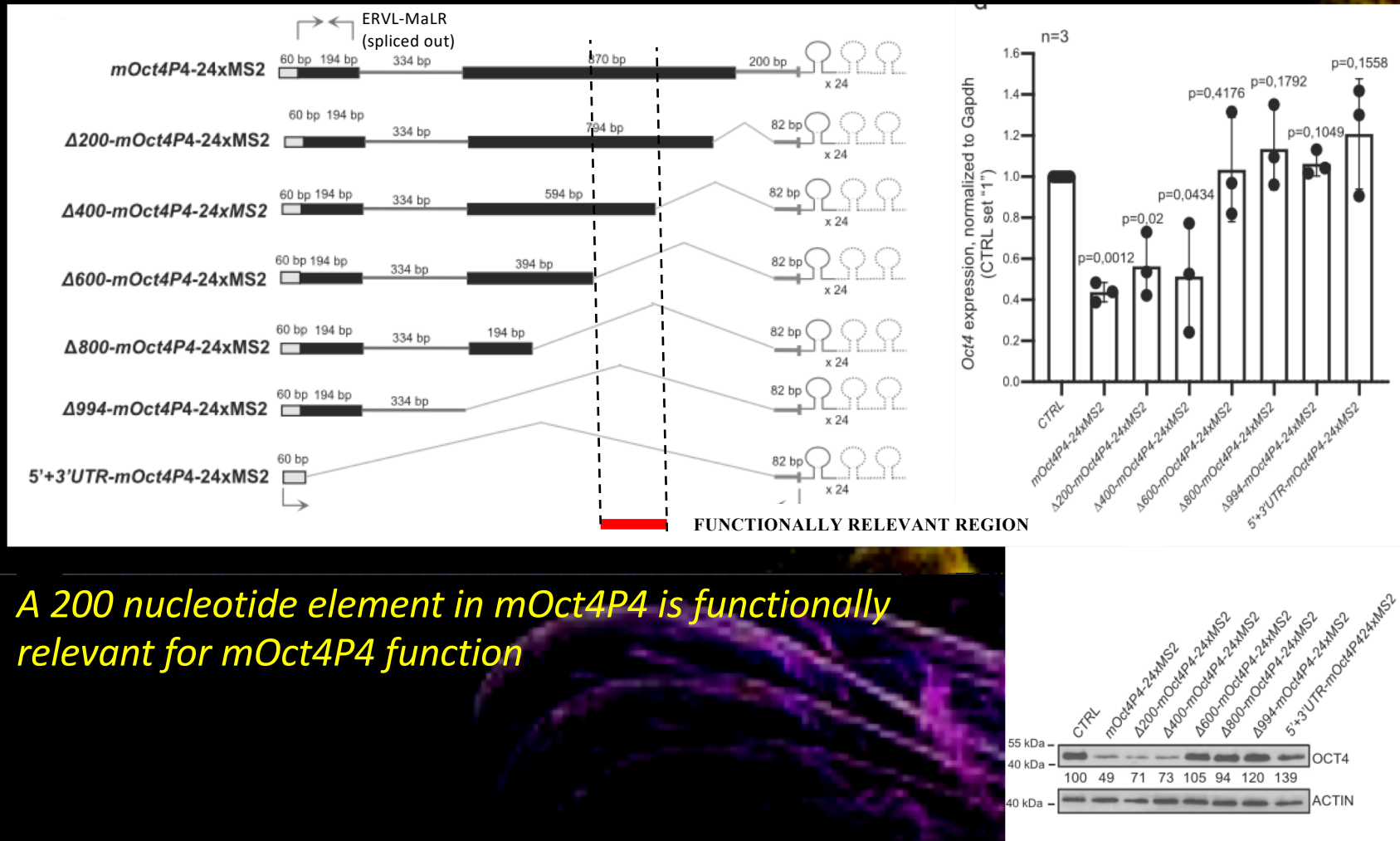
Left Graph: Expression levels of Sox2, Nanog, and Gdf3, normalized to Gapdh.

- Sox2:** dCas9 empty EB D6 (grey bar) is approximately 1.0; dCas9 sgOct4P4 EB D6 (black bar) is approximately 8.0. $p=0,01137$, $n=3$.
- Nanog:** dCas9 empty EB D6 (grey bar) is approximately 1.0; dCas9 sgOct4P4 EB D6 (black bar) is approximately 7.0. $p=0,01498$, $n=3$.
- Gdf3:** dCas9 empty EB D6 (grey bar) is approximately 1.0; dCas9 sgOct4P4 EB D6 (black bar) is approximately 19.0. $p=0,04038$, $n=3$.

Right Graph: Expression levels of Fgf5 and Nestin, normalized to Gapdh.

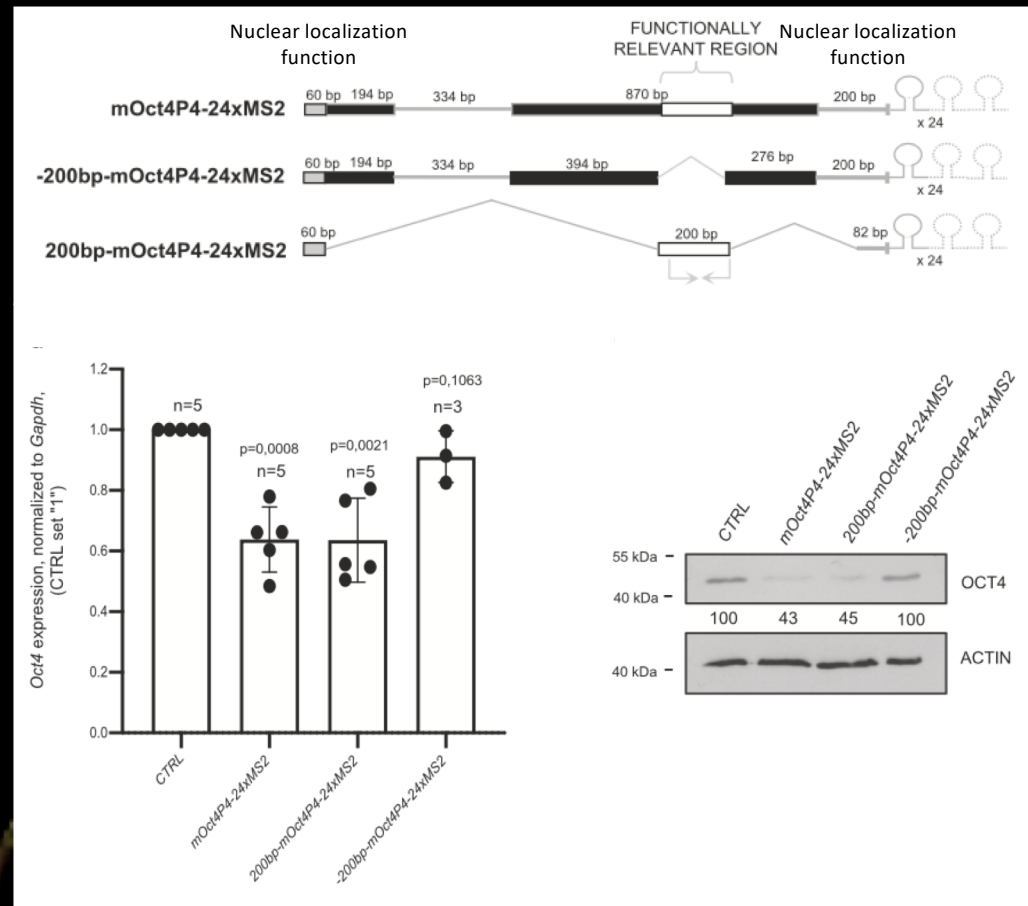
- Fgf5:** dCas9 empty EB D6 (grey bar) is approximately 1.0; dCas9 sgOct4P4 EB D6 (black bar) is approximately 0.55. $p=0,03373$, $n=3$.
- Nestin:** dCas9 empty EB D6 (grey bar) is approximately 1.0; dCas9 sgOct4P4 EB D6 (black bar) is approximately 0.35. $p=0,00195$, $n=3$.

A detailed deletion analysis to identify functional relevant RNA sequences

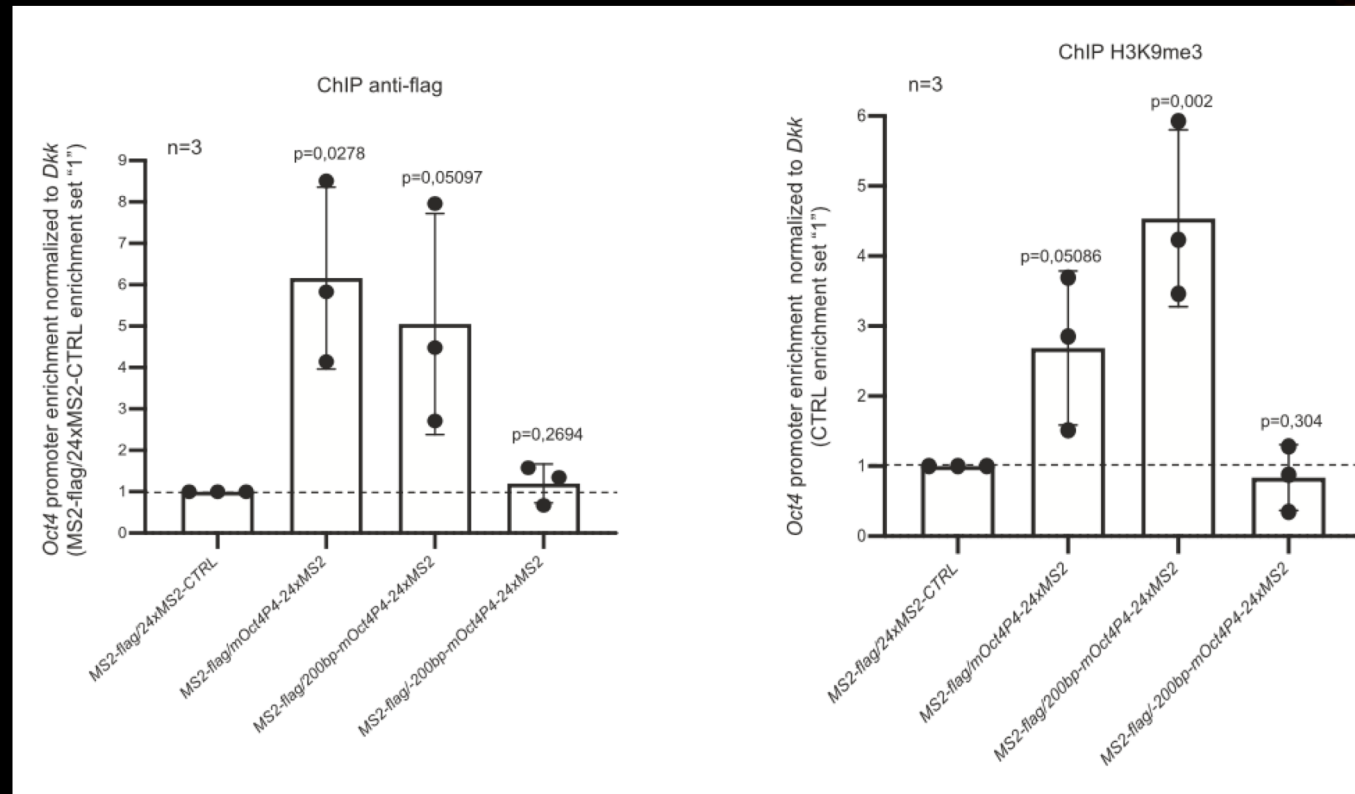


A 200 nucleotide element in mOct4P4 is functionally relevant for mOct4P4 function

A 200nt RNA sequence recapitulates mOct4P4 function

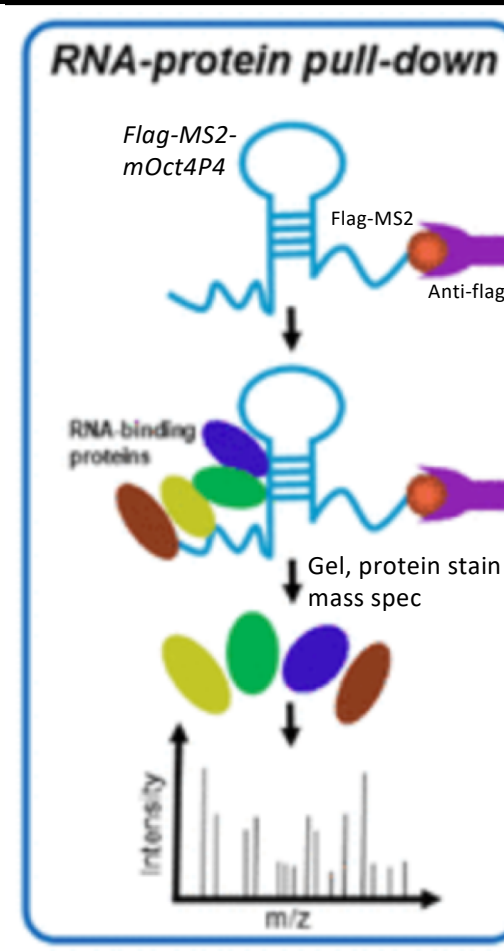
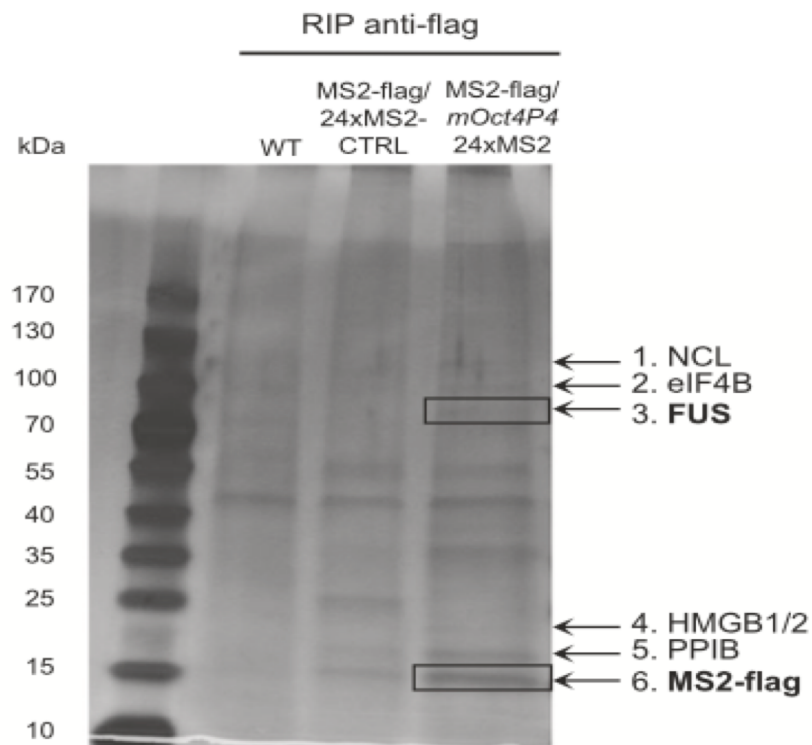


A 200nt RNA sequence recapitulates mOct4P4 function



Are there proteins that bind Oct4P4 lncRNA to mediate silencing function?

RNA Immunoprecipitation (flag-MS2 tagged RNA)



d

MS2-flag + +

Input

RIP anti-flag + +

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-flag 55 kDa

RT-PCR 200bp

100bp

24xMS2-CTRL siCTRL

24xMS2-CTRL siFus

mOct4P4-24xMS2 siCTRL

mOct4P4-24xMS2 siFus

70 kDa

55 kDa

40 kDa

100 112 17 88

40 kDa

MS2-flag

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-FLAG 55 kDa

RT-PCR 200 bp

100 bp

mOct4P4

e

MS2-flag + +

Input

RIP anti-flag + +

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-flag 55 kDa

RT-PCR 200bp

100bp

24xMS2-CTRL siCTRL

24xMS2-CTRL siFus

mOct4P4-24xMS2 siCTRL

mOct4P4-24xMS2 siFus

70 kDa

55 kDa

40 kDa

100 112 17 88

40 kDa

MS2-flag

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-FLAG 55 kDa

RT-PCR 200 bp

100 bp

mOct4P4

f

MS2-flag + +

Input

RIP anti-flag + +

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-flag 55 kDa

RT-PCR 200bp

100bp

24xMS2-CTRL siCTRL

24xMS2-CTRL siFus

mOct4P4-24xMS2 siCTRL

mOct4P4-24xMS2 siFus

70 kDa

55 kDa

40 kDa

100 112 17 88

40 kDa

MS2-flag

mOct4P4-24xMS2

WB anti-FUS 70 kDa

WB anti-FLAG 55 kDa

RT-PCR 200 bp

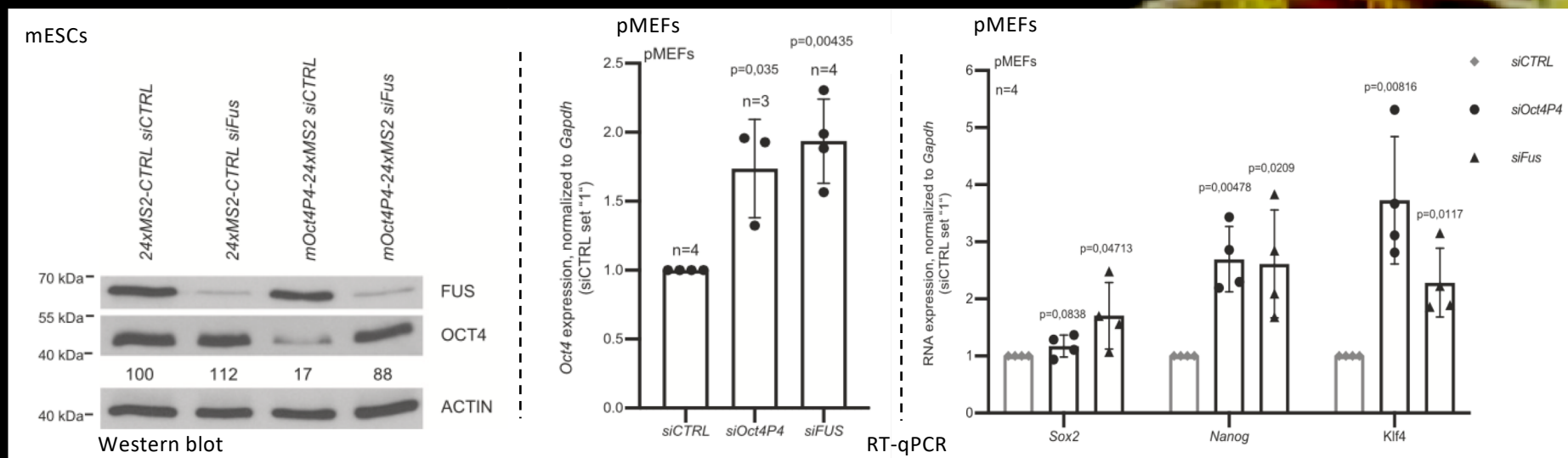
100 bp

mOct4P4

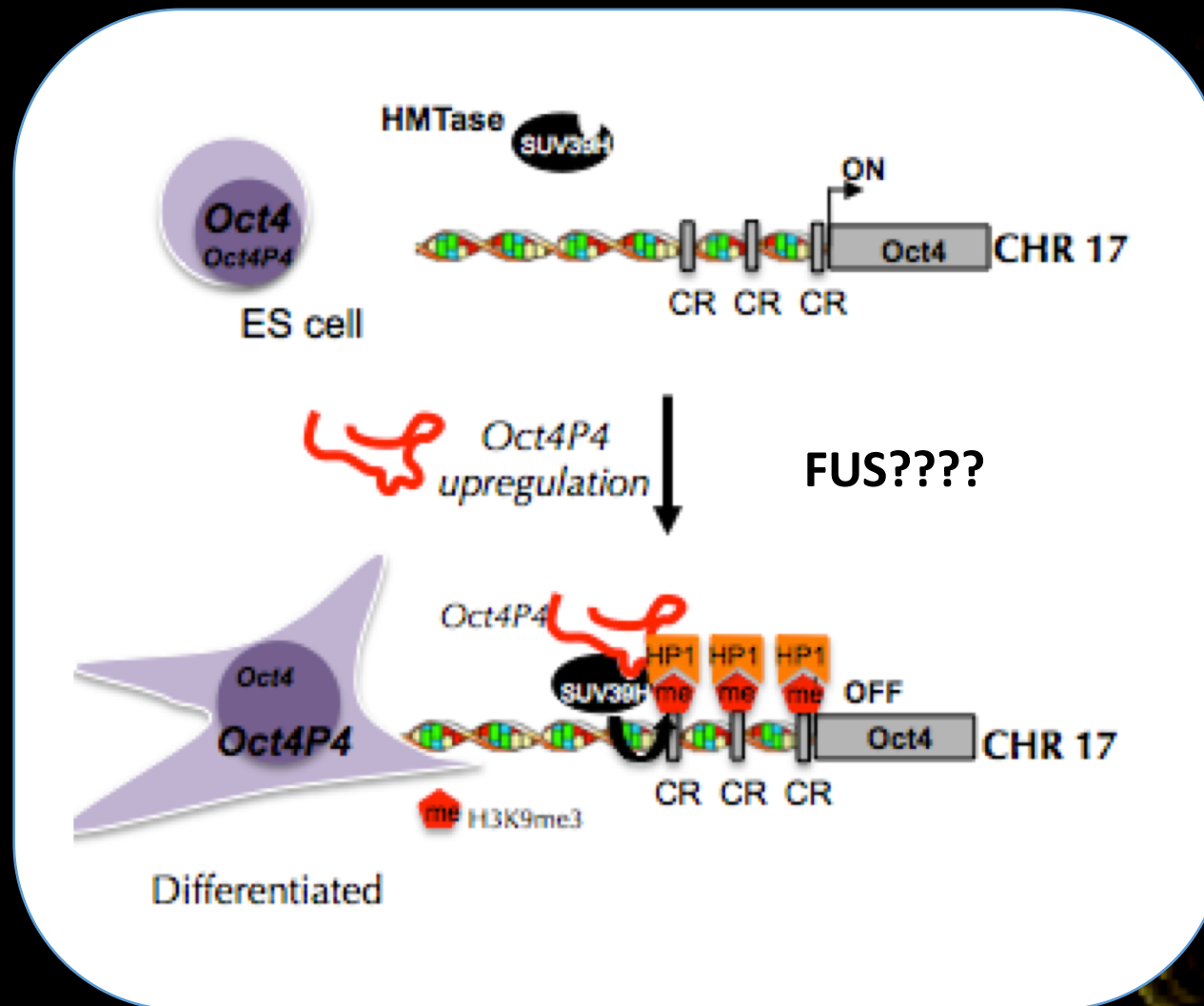
FUS is required for mOct4P4 lncRNA mediated silencing of Oct4

Question 1: Does FUS loss of function break mOct4P4 function in mESCs?

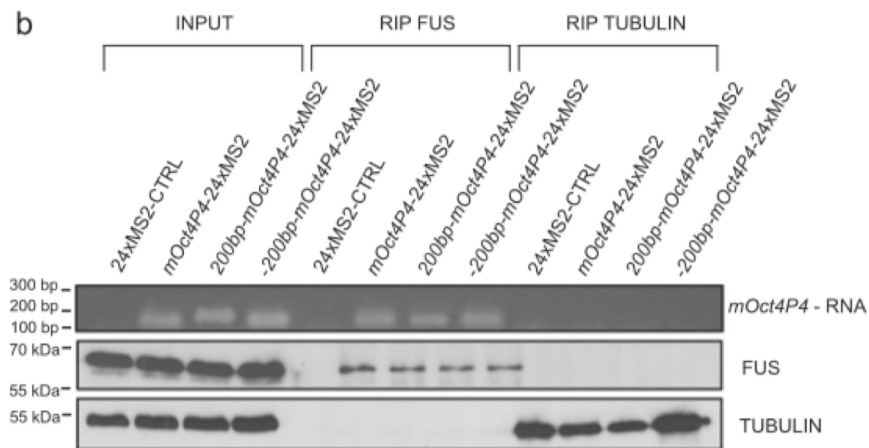
Question 2: Does loss of function of FUS break silencing of Oct4 in differentiated cells?



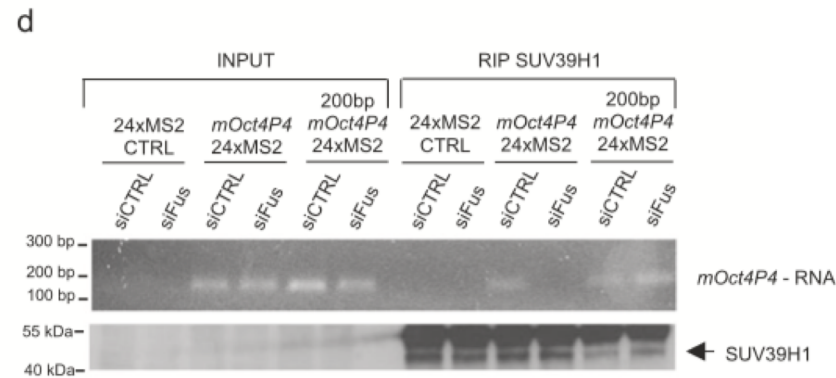
HOW DOES FUS RELATE TO SUV39H1???



FUS binds to Oct4P4 to allow Suv39h1 recruitment

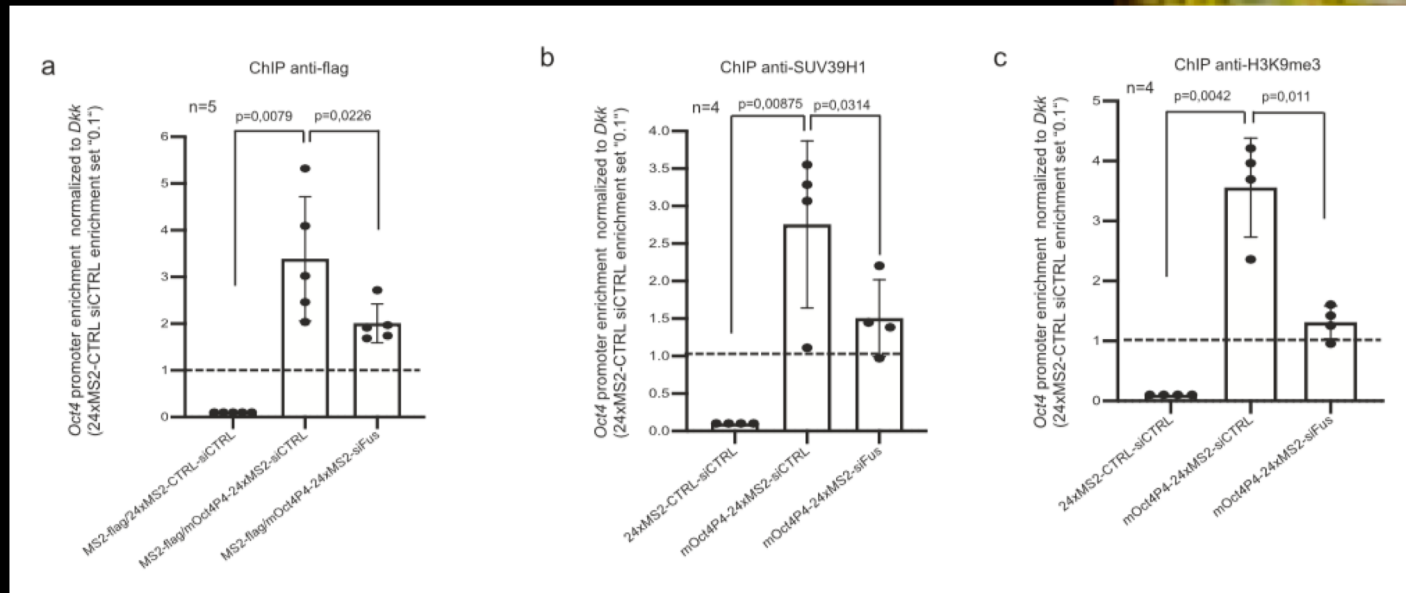


FUS binds mOCT4P4 with or without 200bp region



Suv39h1 binds mOCT4P4 only when FUS binds mOct4P4 lncRNA

FUS binds to Oct4P4 to allow Suv39h1 recruitment and targeting of the Oct4 locus



FUS binds to Oct4P4 to allow Suv39h1 recruitment and targeting of the Oct4 locus

