

SUPPLEMENTARY INFORMATION

KDM4B-mediated reduction of H3K9me3 and H3K36me3 levels improves somatic cell reprogramming into pluripotency

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SUPPLEMENTAL MATERIALS AND METHODS

Teratomas

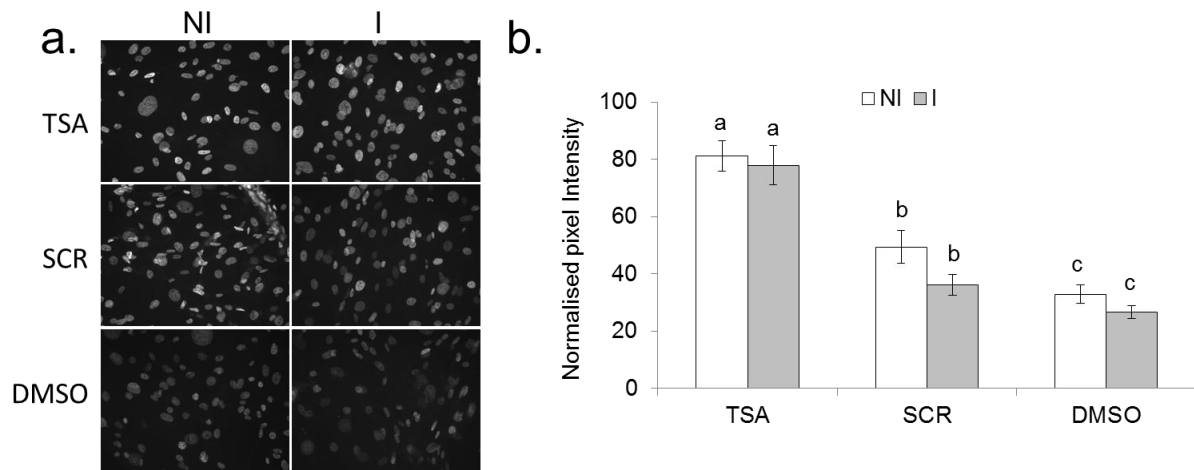
iPSC lines at passage 3-6 were harvested with a cell scraper, centrifuged and re-suspended in PBS + 1% PVA (10-30k). Using a 23G needle, 100 μ l of cell suspension (approx. $1 - 5 \times 10^6$ cells per site) was injected intramuscularly into the quadriceps of adult immune-deficient mice. After 5–7 weeks, tumors were dissected, fixed overnight in Davidson's fixative, embedded in paraffin, sectioned, haematoxylin–eosin stained and analyzed by a pathologist service (NZVP, Hamilton, NZ).

Chimeras

iPSC lines were either aggregated with morulae or injected into host blastocysts (Swiss or FVB/NJArc) and transferred into the oviduct of day 0.5 pseudo-pregnant females. Coat-colour chimeras were bred with Swiss females for germline transmission.

SUPPLEMENTAL FIGURES

Figure S1



SUPPLEMENTAL FIGURE LEGENDS

Figure S1. Effect of HDACi on F-*Kdm4b*-EGFP expressing MEFs (a)

Immunofluorescence of H3K9ac in 48 h non-induced (NI) vs induced (I) F-*Kdm4b* MEFs. **(b)**

Quantification of immunofluorescence analysis in (A). Values represent normalized pixel intensity (ROI/area) \pm SEM. Asterisks indicate significant differences between induced (I) vs non-induced (NI) MEFs and their corresponding HDACi treatments (N=20 nuclei quantified per treatment, n=4 replicates). Within each group (NI vs I), bars with different superscripts differ ^{ab}P<0.005 by two-tailed unpaired t-test.

SUPPLEMENTAL TABLES

Supplemental Table 3. Functional annotation of differently expressed genes upon *Kdm4b*-induction

<i>Top 3 Molecular and Cellular Functions*</i>	<i>P-value</i>
Cellular Compromise	5.60E-03 - 1.65E-04
Cellular Function and Maintenance	5.60E-03 - 1.65E-04
Cell Death and Survival	3.74E-03 - 6.24E-04
<i>Top 3 Canonical Pathways</i>	<i>P-value</i>
Oncostatin M Signaling	2.10E-02
Inhibition of Matrix Metalloproteases	2.41E-02
Transcriptional Regulatory Network in Embryonic Stem Cells	2.47E-02
<i>Top 3 Associated Network Functions</i>	<i>Score</i>
Cell Cycle	27
Immunological Disease, Molecular Transport	3
Carbohydrate Metabolism, Skeletal and Muscular System Development and Function, Small Molecule Biochemistry	3

*Gene Ontology was determined by QIAGEN's Ingenuity® Pathway Analysis.

Supplemental Table 4: In vitro development of F-*Kdm4b* NT reconstructs treated \pm HDACi

Induction ^a	HDACi ^b	n	nNT ^c	Morulae (% \pm SEM)	Blastocysts (% \pm SEM)
NI	-	5	130	1 (1 \pm 0.2%) ^a	1 (1 \pm 1%) ^a
I	-	12	230	16 (7 \pm 1%) ^b	11 (5 \pm 2%)
NI	SCR	2	59	4 (7 \pm 1%)	9 (15 \pm 2%) ^b
I	SCR	4	86	6 (7 \pm 2%)	9 (11 \pm 2%)
NI	TSA	3	81	4 (5 \pm 1%)	3 (4 \pm 1%)
I	TSA	9	155	12 (8 \pm 1%)	7 (5 \pm 1%)

^aMEFs (#1, 3, 9) were either Dox-induced (I) or non-induced (NI) for 48 hours: ^bNT

reconstructs were treated with scriptaid (SCR), trichostatin (TSA) or DMSO; ^cnumber of NT reconstructs that extruded a single polar body and were cultured until day 4. Significance was

determined via Fisher 2x2 test; ^{ab}Cells with different superscripts within a column differ with

P<0.01

Supplemental Table 5. Teratoma formation from *Kdm4b*-EGFP iPSC lines

Animal ID	Cell line	Dox	Tumor diameter (mm)	Ectoderm*	Mesoderm*	Endoderm*
1	#1 iPSC	-	20	√	√	√
			20	ND	ND	ND
2	#1 iPSC	+	19	√	√	√
			10	ND	ND	ND
3	#3 iPSC	-	No tumor	-	-	-
4	#7 iPSC	-	13	-	√	√
5	v6.5 ESC	-	15	√	√	√
			13	ND	ND	ND

*Germ layer identity was determined on histological sections (Fig. 6D). ID=unique animal

identifier; ND=not determined

Supplemental Table 6. Chimera formation from *Kdm4b*-EGFP iPSC lines

iPSC line	sex	No. of born mice	Chimaeras*	% of Chimaeras*
#1 <i>Col1a1</i> ^{4F2a/F-<i>Kdm4b</i>}	♀	18	3	17%
#7 <i>Col1a1</i> ^{4F2a/F-<i>Kdm4b</i>}	♂	2	1	50%

*Chimerism was determined by coat colour (Fig. 6E).

Supplemental Table 7: Primer sequences used for qPCR

Primer		Sequence (5'- 3')	Size (bp)	Reference
<i>2810474019Rik</i>	F:	TTTCTGTCTCGGTCTTCCGC	181	This paper
	R:	CACGCCTCTAATTCTGTTTTTCCA		
<i>Actb</i>	F:	CAGAAGGACTCCTATGTGGG	200	1
	R:	TTGGCCTTAGGGTTCAGGG		
<i>Transgene Kdm4b</i>	F:	AGAAGACACCGGGACCGATC	190	2
	R:	TGAATTCATCCATGGTGGGG		
<i>C430002N11Rik</i>	F:	AGCCCCTTCGGCTGTTTTTA	136	This paper
	R:	GGGCTGAACCTCCTTCTGTC		
<i>Chst13</i>	F:	ATGGGAAGACGCTCCTGTTG	98	This paper
	R:	TTTTCAAATGCGGGACGCAG		
<i>Cola1 (Wild-type)</i>	F:	GCTCGCACGTACTTCATTCC	400	This paper
	R:	GAGAGTTCCTTGAGGGCTGG		
<i>Endo Kdm4B</i>	F:	AGAAGCCTTCCTGTTCTCAG	189	2
	R:	TGTA CTGACTGGCTGTAGGG		
<i>Ephx2</i>	F:	GGGAAAGGAATTTACCCCGCT	72	This paper
	R:	GAGTGGTACCCACTGGGCTG		
<i>Gapdh</i>	F:	TGCACCACCAACTGCTTAG	176	1
	R:	GATGCAGGGATGATGTTC		
<i>Glipr1</i>	F:	CGTGCAGTGATTGCCCAAAA	90	This paper
	R:	CGACAGAGTAGTAACGTGAGACC		
<i>Gpr56</i>	F:	TTTTCTCTGGTGCAAGGTGC	192	This paper
	R:	GAAGCGGGGAATATCTGGGG		
<i>Gusb</i>	F:	ATAAGACGCATCAGAAGCCG	96	3
	R:	ACTCCTCACTGAACATGCGA		
<i>Hprt</i>	F:	GAAATGTCAGTTGCTGCGTC	332	4
	R:	GCCAACACTGCTGAAACATG		
<i>Lrrn1</i>	F:	TCCTCATCCTCCGGCTAGTG	93	This paper
	R:	GAGACGTGTCCTAAGGCTGG		
<i>Ly6a</i>	F:	GGAGCTGCTAGGTTTTATCTGTGC	178	This paper

<i>M2-rtTA</i>	R:	GGGCAGGTAATTGATGGGCA	407	This paper
	F:	GAGAGTTCCTTGAGGGCTGG		
<i>Mmp13</i>	R:	CAATTGCTTGTCTCAGAAGTGG	108	This paper
	F:	GGAGCCCTGATGTTTCCCAT		
<i>Nnat</i>	R:	GTCTTCATCGCCTGGACCATA	104	This paper
	F:	CCACCCACTTTCGGAACCAT		
<i>Prss42</i>	R:	ACTGGGGACAGGGTCTGC	130	This paper
	F:	GAAGAGGGGAAGTGGCCTTG		
<i>Rosa26 (Wild-type)</i>	R:	CGTTGTACTGGATTCGGCTGTA	700	This paper
	F:	GCTGTTTTGGAGGCAGGAAG		
<i>Serpina3g</i>	R:	TGCCAATGCTCTGTCTAGGG	139	This paper
	F:	ACAGAGGCTGAAAAGGAGCAG		
<i>Slc29a3</i>	R:	AGAGGCTGAAGGCAAAGTCA	91	This paper
	F:	CTTTGAGAGCTACCTGGCAGT		
<i>Sry</i>	R:	GTGCACCTGGACCCTGTTGA	402	5
	F:	TGGGACTGGTGACAATTGTC		
<i>Trim24</i>	R:	GAGTACAGGTGTGCACCTCT	150	This paper
	F:	GCCATTCGCCACCCAAGT		
<i>Zfp37</i>	R:	CAGCTTGTACATACCTGATTAGACT	198	This paper
	F:	GGACCTGACAAAGCCAGAGG		
	R:	TGATTTCCCATGGAGGCTGG		

Supplemental Table 8: Overview of MEF lines used for the various assays

Assay	Induction	MEF lines													
		<i>F-Kdm4b</i>							<i>M-Kdm4b</i>	<i>F-Kdm4b/iPS</i>			<i>M-Kdm4b/iPS</i>		<i>WT/iPS</i>
	Dox (days)	#1 (♂)	#3 (♀)	#4 (♀)	#5 (♀)	#6 (♀)	#9 (♂)	#11 (♀)	#13 (♀)	#1 (♀)	#2 (♂)	#7 (♂)	#1 (♀)	#3 (♀)	#1 (♂)
Flow cytometry	1-2	√	-	√	√	√	√	√	√	-	-	-	-	-	-
qPCR	2	-	-	√	√	√	-	√	-	-	-	-	-	-	-
Microarray	2	-	-	√	√	√	-	-	-	-	-	-	-	-	-
mRNA-seq	2	-	-	√	√	√	-	-	-	-	-	-	-	-	-
Immunofluorescence	2	√	√	√	√	-	-	√	-	-	-	-	-	-	-
Western blot	2	-	-	-	√	-	-	√	-	-	-	-	-	-	-
Demethylase activity	2	-	-	-	√	√	-	-	-	-	-	-	-	-	-
NT	1	-	-	√	√	√	-	√	√	-	-	-	-	-	-
NT ± HDACi	1	√	√	-	-	-	√	-	-	-	-	-	-	-	-
iPS colony formation	21	-	-	-	-	-	-	-	-	√	√	√	√	√	√
Teratoma or chimera	21	-	-	-	-	-	-	-	-	√	√	√	-	-	-

√=tested for this assay; -=not determined; NT=Nuclear transfer, WT=wild-type

Supplemental Table 9: Primary and secondary antibodies

<i>Antibody</i>	<i>Dilution</i>		<i>Manufacturer (Cat.-no.)</i>
	IF	WB	
Alexa Fluor donkey anti mouse 488	1:1000	NA	Thermo Fisher (Cat-A11008)
Alexa Fluor goat anti rabbit 568	1:1000	NA	Thermo Fisher (Cat-A11011)
Goat anti rabbit HRP	NA	1:10000	Thermo Fisher (Cat-G21234)
Mouse anti GFP	1:1000	NA	Thermo Fisher (Cat-A11120)
Rabbit anti H3K9Ac	1:250	NA	Millipore (Cat-04-1003)
Rabbit anti H3K9me1	1:1000	1:6000	Gift by T Jenuwein (Freiburg, Germany)
Rabbit anti H3K9me2	1:1000	1:3000	Gift by T Jenuwein (Freiburg, Germany)
Rabbit anti H3K9me3	1:1000	1:3000	Gift by T Jenuwein (Freiburg, Germany)
Rabbit anti H3K27me3	1:1000	1:3000	Gift by T Jenuwein (Freiburg, Germany)
Rabbit anti H3K36me3	1:500	1:1000	Cell Signaling (#9763)
Rabbit anti NANOG	1:100	NA	Abcam (ab80892)

NA=not applied

SUPPLMENTAL REFERENCES

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