

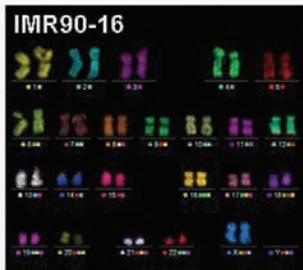
Red blood cell generation from human induced pluripotent stem cells: perspectives for transfusion medicine

Hélène Lapillonne,^{1,2,3*} Ladan Kobari,^{1,2} Christelle Mazurier,^{1,4} Philippe Tropel,^{5,6} Marie-Catherine Giarratana,^{1,2} Isabelle Zanella-Cleon,⁷ Laurent Kiger,^{8*} Marie Wattenhofer-Donzé,⁹ Hélène Puccio,⁹ Nicolas Hebert,^{1,2} Alain Francina,¹⁰ Georges Andreu,¹¹ Stéphane Viville,⁵ and Luc Douay^{1,2,3,4}

¹INSERM, UMR_S938, Proliferation and Differentiation of Stem Cells, Paris, France; ²UPMC Univ Paris 06, UMR_S938, Proliferation and Differentiation of Stem Cells, Paris, France; ³AP-HP, Hôpital Armand Trousseau, Service d'Hématologie biologique, Paris, France; ⁴Etablissement Français du Sang Ile de France, Ivry-sur-Seine, France; ⁵IGBMC, Department of Cell Biology and Development, Illkirch Cedex, France; ⁶INSERM/UEVE UMR-861, I-STEM, AFM, Institute for Stem Cell Therapy and Exploration of Monogenic Diseases, 5 rue Henri Desbruyères, 91030 Evry cedex, France; ⁷Institut de Biologie et de Biochimie des Protéines, CNRS UMR 5086, IFR 128, Université Claude Bernard-Lyon I, Lyon, France; ⁸INSERM U473, Hôpital de Bicêtre, Le Kremlin Bicêtre, France; ⁹IGBMC, Department of Neurobiology and Genetics, Illkirch Cedex, France; ¹⁰Unité de Pathologie Moléculaire du Globule Rouge, Fédération de Biochimie et de Biologie Spécialisée, Hôpital Edouard Herriot, Lyon, France, and ¹¹Institut National de la Transfusion Sanguine (INTS), Paris, France

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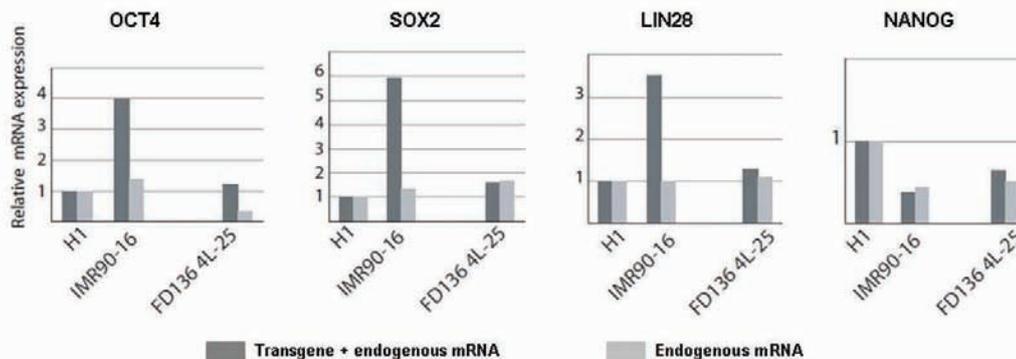
A



B

		oct-04	Oct4A	Nanog	Sox2	KLF4	DNMT	TGDF ₁	GDF3	Cadh	ZFP42
	hEC 2102E P	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00
IMR90	clone 16	0,49	0,12	0,01	0,46	0,01	0,18	0,09	0,34	0,36	0,13

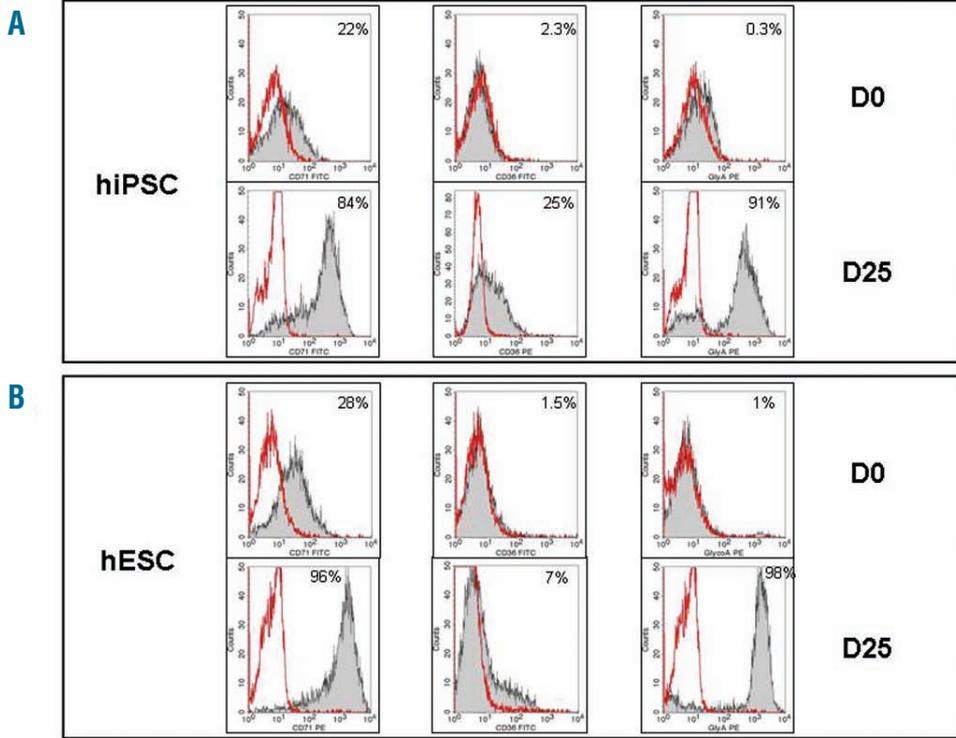
C



D

		Ectoderm		Mesoderm					Endoderm			
		Pax 6	ck18	Brachyury	Gata4	RunX 1	CD34	Nkx2. 5	KDR	ck17	AFP	Gata 4
		hEC 2102E P	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00	1,00
IMR90	clone 16	2,66	1,06	4,26	7,16	12,47	2,04	17,15	0,59	12,30	9,51	7,16

Online Supplementary Figure S1. hiPSC characterization. (A) hiPSC (IMR90-16, passage 10) maintain a normal karyotype. High resolution, multi-FISH karyotype indicates a normal, diploid, female chromosomal content. (B) Expression of hESC-marker genes in hiPSC (IMR90-16) clones by quantitative reverse transcription - PCR analyses (qRT-PCR). Individual PCR reactions were normalized against GAPDH and standardized relative to the expression level in human embryonic carcinoma 2102EP cells. (C) qRT-PCR analyses of OCT4, SOX2, LIN 28, and NANOG expression in fetal hiPSC lines (IMR90-16, passage 10), adult foreskin fibroblasts (FD136), adult hiPSC FD136 4L 25 passage 11 and hESC (H1). Dark gray bars represent amplification of both transgene and endogenous mRNA. Light gray bars represent the amplification of endogenous mRNA only. Data were normalized to H1 level that was set at 1. (D) Gene expression in differentiated hiPSC (IMR90-16) clones by qRT-PCR analyses. qRT-PCR performed on embryoid bodies-differentiated hiPSC shows up-regulated expression of lineage markers from all three embryonic germ layers (ectoderm: PAX6 and CK18; endoderm: GATA4, AFP and CK17; mesoderm: brachyury, GATA4, RUNX1, CD34, NKX2.5 and KDR).



Online Supplementary Figure S2. Phenotypic analyses during erythroid differentiation from hiPSC-IMR90-16 (A) and hESC-H1 (B) by flow cytometry. Expression of erythroid antigens: CD36, CD71 and Glyco-A, from day 0 after hEB dissociation to day 25 – the final point of the erythroid differentiation.

Online Supplementary Table S1. Primer sets for PCR reactions. “Endogenous” indicates that primers included in the 3’ untranslated region measure expression of the endogenous gene only, whereas “total” indicates that primers in coding regions measure expression of both the endogenous gene and the transgene if present.

Genes	Sequences (5' to 3')
OCT4 - Total	F CTGGAGCAAACCCGGAGGAG R GCAGATGGTCGTTTGGCTGAATACC
OCT4 - Endogenous	F AGTTTGTGCCAGGGTTTTTG R ACTTCACCTTCCCTCCAACC
SOX2 - Total	F TACCTCTTCTCCCACTCCA R GGTAGTGCTGGGACATGTGA
SOX2 - Endogenous	F AGTCTCAAGCGACGAAAAA R TTTCACGTTTGCAACTGTCC
LIN28 - Total	F AAGCGCAGATCAAAGGAGA R CTGATGCTCTGGCAGAAGTG
LIN28 - Endogenous	F AGTGGCCTGGATAGGGAAGT R CTTGGCTCCATGAATCTGGT
NANOG - Total	F CAACATCCTGAACCTCAGCTAC R ATTGTTCCAGGTCTGGTTGC
NANOG - Endogenous	F TTTGGAAGCTGCTGGGGAAG R GATGGGAGGAGGGGAGAGGA
GAPDH	F GTGGACCTGACCTGCCGTCT R CTGTAGCCAAATTCGTTGTC

Online Supplementary Table S2. Analysis of stem cell and differentiation markers in hiPSC (FD136-25) in normal culture conditions and after differentiation in embryoid bodies (FD136-25 iEB). Stem cell and differentiation marker expression was investigated with TaqMan[®] Human Stem Cell Pluripotency Array (Applied Biosystems). Results were analyzed using the comparative $\Delta\Delta Ct$ method with *GAPDH* as the endogenous control and a normal hESC sample as calibrator (HUES-24 line; kindly provided by M. Puc at), ND: not detected. Genes showing at least a 2-fold increase/decrease in differentiating cells compared to the undifferentiated state were considered up- or down-regulated, respectively. The formation of embryoid bodies allows a rapid assessment of pluripotent cell lines. Results from TaqMan low density arrays show a clear activation of a wide panel of genes characteristic of all three germ layers, demonstrating pluripotency of adult-derived hiPSC (blood, bone, cardiac muscle, cartilage, endothelium, muscle; mesoderm; dopaminergic neurons, motor neurons, neurons, neural, astrocytes, oligodendrocytes: ectoderm; hepatocytes, pancreas, visceral endoderm: endoderm).

Expression profile	Gene	FD136-25	FD136-25 iEB	Expression profile	Gene	FD136-25	FD136-25 iEB
Astrocytes	GFAP-H600157674_m1	0,18	0,05	Stem Cell	BRX-H600217848_m1	0,30	0,41
Blood	HEB-H600247223_g1	ND	ND	Stem Cell	CD9-H60023521_m1	0,12	0,09
Blood	HBZ-H600744391_g1	2,49	17,08	Stem Cell	COMM3-H60001350_m1	0,42	3,93
Bone	COL1A1-H600164004_m1	0,19	7,52	Stem Cell	CRAFP2-H600275636_m1	1,22	7,25
Bone	RUNX2-H600231892_m1	32,98	328,65	Stem Cell	DNM1T3B-H600171876_m1	0,21	0,02
Cardiac Muscle	ACTC-H600606316_m1	0,34	0,78	Stem Cell	EBAF-H600745761_g1	0,00	0,07
Cardiac Muscle	NPPA-H600383230_g1	ND	24,57	Stem Cell	FGF4-H600173564_m1	0,15	0,07
Cartilage	COL2A1-H600156589_m1	1,71	65,68	Stem Cell	FGF5-H600170454_m1	4,23	51,08
Dopaminergic Neurons	TH-H600165341_m1	ND	13,52	Stem Cell	FOXO3-H600255287_g1	0,16	0,02
Endogenous Control	18S-H609999001_g1	1,73	2,11	Stem Cell	GABRB3-H600241459_m1	0,24	0,04
Endogenous Control	ACTB-H609999803_m1	0,77	2,00	Stem Cell	GAL-H600544955_m1	0,02	0,01
Endogenous Control	CTNND1-H600170025_m1	0,37	1,42	Stem Cell	GBX2-H600230965_m1	0,19	0,65
Endogenous Control	EEF1A1-H600742749_g1	0,50	0,50	Stem Cell	GOP3-H600220996_m1	0,13	0,04
Endogenous Control	GAPD-H609999905_m1	1,00	1,00	Stem Cell	GRB7-H600917999_g1	0,25	0,14
Endogenous Control	RAF1-H600234119_m1	0,32	0,57	Stem Cell	IFTM1-H600705137_g1	0,16	0,07
Endothelial	CD34-H600156373_m1	0,26	8,00	Stem Cell	IFTM2-H600294965_gH	0,10	0,67
Endothelial	CDH5-H600174344_m1	0,40	259,89	Stem Cell	IL6ST-H600174360_m1	0,29	19,88
Endothelial	FLT1-H600176573_m1	1,73	3,11	Stem Cell	IMP2-H600538956_m1	0,65	1,07
Endothelial	PECAM1-H600169777_m1	0,27	45,01	Stem Cell	KIT-H600174029_m1	0,26	0,45
Extraembryonic endoderm	FOXA2-H600232764_m1	0,09	0,38	Stem Cell	LEFTB-H600764128_g1	0,01	0,00
Extraembryonic endoderm	GATA4-H600174603_m1	0,15	0,91	Stem Cell	LFR-H600190730_m1	0,17	0,79
Extraembryonic endoderm	PITF1A-H600603996_g1	ND	ND	Stem Cell	LINC8-H600702808_g1	0,93	0,42
Germ Cell	DDX4-H600251859_m1	ND	ND	Stem Cell	Nanog-H602387400_g1	0,19	0,03
Germ Cell	SYCP3-H600538143_m1	8,18	25,31	Stem Cell	NODAL-H600415443_m1	0,07	0,04
Hepatocytes	TAT-H600366830_m1	2,44	5,06	Stem Cell	NOG-H600271522_g1	0,08	0,96
Mesoderm	GATA6-H600230018_m1	0,03	0,96	Stem Cell	NRS42-H600187067_m1	0,15	0,36
Mesoderm	T-H600510080_m1	0,08	1,73	Stem Cell	NRS41-H600265986_m1	0,60	0,35
Mesoderm	WFI-H600240913_m1	ND	25,31	Stem Cell	POU5F1-H600742896_g1	0,49	0,23
Muscle	HLXB9-H600232126_m1	0,03	0,55	Stem Cell	PITEN-H60022813_g1	0,66	0,03
Muscle	DES-H600157258_m1	0,26	0,69	Stem Cell	REST-H600194498_m1	0,73	0,07
Muscle	MYF5-H600271574_m1	1,77	ND	Stem Cell	SEMA3A-H600173610_m1	0,41	0,43
Muscle	MYO01-H600159228_m1	ND	ND	Stem Cell	SFRP2-H600293258_m1	0,29	0,37
Neural	NES-H600707120_g1	0,39	1,14	Stem Cell	SOX2-H600602736_g1	1,10	0,41
Neural	NEUROD1-H600155998_m1	4,68	34,09	Stem Cell	TGDF1-H602339499_g1	0,17	0,00
Neural	PAIS-H60240671_m1	1,28	3,05	Stem Cell	TERT-H600162669_m1	0,39	0,12
Neurons	SYP-H600300631_m1	0,58	0,33	Stem Cell	TF-CP2L1-H600232708_m1	0,18	0,18
Oligodendrocytes	OLIG2-H600377600_m1	11,98	62,78	Stem Cell	UTF1-H600247497_g1	0,02	0,02
Pancreas	GCG-H600174867_m1	ND	2,33	Stem Cell	Xist-H601079624_m1	1,92	5,76
Pancreas	IAPP-H600163095_m1	ND	ND	Stem Cell	ZFP42-H600399279_m1	0,04	0,00
Pancreas	INS-H600365773_m1	ND	ND	Stem Cell	CDX2-H600230919_m1	10,81	1729,46
Pancreas	IPF1-H600236830_m1	ND	ND	Trophoblast	CGB-H600361224_gH	0,11	1,49
Pancreas	PAX4-H600173014_m1	ND	ND	Trophoblast	EOMES-H600128972_m1	0,24	0,11
Pancreas	SST-H600174949_m1	5,05	808,59	Trophoblast	GCM1-H600172692_m1	ND	28,87
Parietal Endoderm	FNI-H600277509_m1	0,40	7,20	Trophoblast	KRT1-H600198158_m1	ND	ND
Parietal Endoderm	LAMAI-H600300650_m1	0,40	2,56	Trophoblast	ISL1-H600158126_m1	0,53	32,75
Parietal Endoderm	LAMB1-H600168620_m1	0,24	2,14	V3 Interneurons	AFP-H600173480_m1	166,57	123419,89
Parietal Endoderm	LAMC1-H600267056_m1	0,51	2,37	Visceral Endoderm	SERPINA1-H600165475_m1	3,57	1532,01
Parietal Endoderm	SOX17-H600751752_g1	0,08	0,49	Visceral Endoderm			