ELSEVIER

Contents lists available at ScienceDirect

# Stem Cell Research



journal homepage: www.elsevier.com/locate/scr

Lab Resource: Stem Cell Line

# Establishment of DYT5 patient-specific induced pluripotent stem cells with a *GCH1* mutation



Nagahisa Murakami <sup>a,b</sup>, Taizo Ishikawa <sup>a,c</sup>, Takayuki Kondo <sup>a</sup>, Keiko Imamura <sup>a</sup>, Kayoko Tsukita <sup>a</sup>, Takako Enami <sup>a</sup>, Misato Funayama <sup>a</sup>, Ran Shibukawa <sup>a</sup>, Shinichi Matsumoto <sup>d</sup>, Yuishin Izumi <sup>b</sup>, Etsuro Ohta <sup>e</sup>, Fumiya Obata <sup>e</sup>, Ryuji Kaji <sup>b</sup>, Haruhisa Inoue <sup>a,f,\*</sup>

<sup>a</sup> Center for iPS Cell Research and Application (CiRA), Kyoto University, Kyoto, Japan

<sup>b</sup> Department of Clinical Neuroscience, Institute of Biomedical Sciences, Tokushima University Graduate School, Tokushima, Japan

<sup>c</sup> Sumitomo Dainippon Pharma, Osaka, Japan

<sup>d</sup> Department of Neurology, Shinko Hospital, Kobe, Japan

e Division of Clinical Immunology, Graduate School of Medical Sciences, Kitasato University, Kanagawa, Japan

<sup>f</sup> Drug-Discovery Cellular Basis Development Team, RIKEN BioResource Center, Kyoto, Japan

#### ARTICLE INFO

Article history: Received 13 March 2017 Received in revised form 11 July 2017 Accepted 25 July 2017 Available online 29 July 2017

## ABSTRACT

Peripheral blood mononuclear cells (PBMCs) were collected from a clinically diagnosed 20-year-old dystonia patient with a *GCH1* mutation (DYT5). Episomal vectors were used to introduce reprogramming factors (OCT3/4, SOX2, KLF4, L-MYC, LIN28, and p53 carboxy-terminal dominant-negative fragment) to the PBMCs. The generated iPSCs expressed pluripotency markers, and were capable of differentiating into derivates of all three germ layers *in vitro*. The iPSC line also showed a normal karyotype and preserved the *GCH1* mutation. This cellular model can provide opportunities to perform pathophysiological studies for aberrant dopamine metabolism-related disorders.

© 2017 The Authors. Published by Elsevier B.V. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

	(continued)			
Resource table.		resistance Inducible/constitutive	None	
Unique stem cell line identifier	CIRAi001-A	system Date archived/stock	Oct 2016	
Alternative name of stem cell line Institution	HPS1816 Center for iPS Cell Research and Application (CiRA), Kyoto	date Cell line repository/bank Ethical approval	RIKEN BioResource Center, Japan http://en.brc.riken.jp/index.shtml The Ethics Committee of the Department of Medicine and	
Contact information of distributor	University Haruhisa Inoue haruhisa@cira.kyoto-u.ac.jp		Graduate School of Medicine, Kyoto University (R0091 and G259) and Tokushima University (3172 and H26-9)	
Type of cell line	iPSC			
Origin Additional origin info	Human Age: 20-year-old Sex: male	Resource utility		
Cell Source Method of reprogramming	Peripheral blood mononuclear cells (PBMCs) Episomal vectors	Research investigating dystonia using patient-derived iPSCs is limit- ed, and a single nucleotide polymorphism in <i>GCH1</i> is reportedly associ-		
Associated disease	DYT5	ated with Parkinson's disease. This finding has the potential to provide new insight into not only DYT5 but also Parkinson's disease. <i>Resource details</i>		
Gene/locus Method of modification	GCH1, c.626+1G>C None			
Gene correction Name of transgene or	NO None			
		Autogonal domin	ant CTD avalabudaalaaa I (CCIII) dafaian ay yuhiah	

Autosomal dominant GTP cyclohydrolase I (GCH1) deficiency, which is characterized by dopa-responsive dystonia with marked diurnal fluctuations, was first reported in 1976 by Segawa et al. (1976). GCH1 is involved in the production of tetrahydrobiopterin (BH4), which is an

E-mail address: haruhisa@cira.kyoto-u.ac.jp (H. Inoue).

Corresponding author.

http://dx.doi.org/10.1016/j.scr.2017.07.029

1873-5061/© 2017 The Authors. Published by Elsevier B.V. This is an open access article under the CC BY license (http://creativecommons.org/licenses/by/4.0/).

essential cofactor for biosynthesis of dopamine. Heterozygous *GCH1* mutation leads to partial deficiency of BH4 and selective decreases of tyrosine hydroxylase in nigrostriatal dopamine neurons, which causes dopa-responsive dystonia (Segawa et al., 2003), alternatively called as Segawa disease or DYT5. A large-scale genome-wide association study meta-analysis also revealed that a single nucleotide polymorphism at *GCH1* is associated with Parkinson's disease (Nalls et al., 2014). Peripheral blood mononuclear cells (PBMCs) derived from a 20-year-old dystonia patient with a *GCH1* mutation c.626 + 1G>C were reprogrammed by inducing episomal vectors with OCT3/4, SOX2, KLF4, L-MYC, LIN28, and p53 carboxy-terminal dominant-negative fragment, as previously reported (Okita et al., 2013) (Fig. 1A). The generated iPSCs expressed pluripotency markers, including NANOG and SSEA4 (Fig. 1B) and were capable of differentiating into derivates of all three germ layers, namely, endoderm (SOX-17), mesoderm ( $\alpha$ SMA), and ectoderm ( $\beta$ III-tubulin)

(Fig. 1C). Short tandem repeat (STR) analysis of the iPSC line was identical to the parental PBMCs (Supplementary Table 1). The iPSC line had a normal karyotype (46 XY) (Fig. 1D). The reprogramming process did not alter a *GCH1* mutation, as demonstrated in Fig. 1E. By flow cytometry, 66.1% of iPSCs were positive for SSEA4 (Fig. 1F). Collectively, the HPS1816 iPSC line was karyotypically normal, pluripotent, and carried a *GCH1* mutation.

### Materials and methods

#### Ethics statements

Generation and use of human iPSCs was approved by the Ethics Committee of the Department of Medicine and Graduate School of Medicine, Kyoto University, and Tokushima University. All methods were

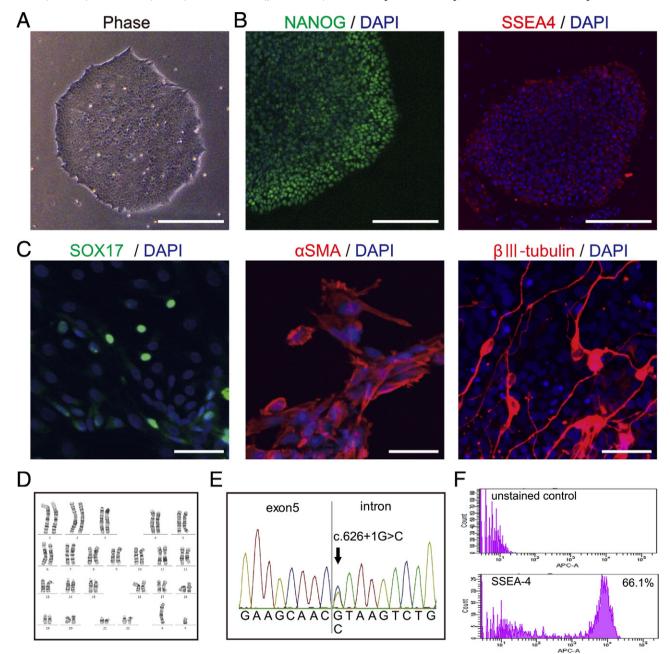


Fig. 1. Characterization of iPSC line. (A) The DYT5 patient iPSCs displayed a typical round-shaped colony under microscopy. Scale bar: 200 µm (B) Immunocytochemical staining of iPSCs showed high expression of pluripotency marker NANOG and SSEA4. Scale bars: 200 µm. (C) Pluripotency of iPSCs was confirmed by *in vitro* three-germ layer assay. Scale bars: 50 µm. (D) Karyotype analysis of patient iPSCs showed a normal karyotype of 46 XY. (E) Sanger sequence of the *GCH1* gene in iPSCs showed a heterozygous c.626+1G>C substitution. (F) FACS analysis with evaluation of SSEA-4 staining.

# Table 1

Characterization and validation.

Classification	Test	Result	Data
Morphology	Photography	Visual record of the line: normal	Fig. 1 panel A
Phenotype	Immunocytochemistry	Assess staining of pluripotency markers: NANOG and SSEA4	Fig. 1 panel B
	Flow cytometry	SSEA4 66.1%	Fig. 1 panel F
Genotype	Karyotype (G-banding) and resolution	46XY	Fig. 1 panel D
		Resolution 300–400	
Identity	Microsatellite PCR (mPCR)	Not performed	Not performed
	STR analysis	16 loci, matched	Supplementary Table 1
Mutation analysis (IF	Sequencing	Heterozygous c.626 + 1G>C mutation	Fig. 1 panel E
APPLICABLE)	Southern Blot OR WGS	Not performed	Not performed
Microbiology and	Mycoplasma	Contamination of Mycoplasma was negative	Not shown but available
virology			from author
Differentiation potential	Embryoid body formation OR Teratoma	Describe expression of genes in embryoid bodies: SOX-17, $\alpha$ SMA,	Fig. 1 panel C
	formation OR Scorecard	and βIII-tubulin	
Donor screening	HIV 1 + 2 Hepatitis B, Hepatitis C	HIV1 (negative), HIV2 (not performed), Hepatitis B (negative), and	Not shown but available
-		Hepatitis C (negative)	from author
Genotype additional info	Blood group genotyping	Not performed	Not performed
	HLA tissue typing	Not performed	Not performed

performed in accordance with approved guidelines. Formal informed consent was obtained from the patient.

#### Generation of iPSCs

Human complementary DNAs for reprogramming factors were transduced in PBMCs with episomal vectors (OCT3/4, SOX2, KLF4, L-MYC, LIN28 and p53 carboxy-terminal dominant-negative fragment). The generated iPSCs were grown under feeder-free conditions on laminin-511 E8 (Nippi, Tokyo, Japan)-coated plates with StemFit (AK02N or AK03N, Ajinomoto, Tokyo, Japan) (Nakagawa et al., 2014) (Table 1).

#### Karyotyping and genotyping of GCH1

Karyotype analysis was performed by LSI Medience (Tokyo, Japan). Genotyping of the *GCH1* mutation was performed by PCR amplification of genomic DNA and direct sequencing (3500xL Genetic Analyzer, Applied Biosystems, Waltham, MA) (Table 2).

#### In vitro three-germ layer assay

iPSCs were dissociated with 0.5× TrypLE Select (Gibco, Waltham, MA) and used for embryoid body (EB) formation. Cells were transferred to suspension plates in Dulbecco's modified Eagle's medium/Ham's F12 (DMEM/F12, Sigma-Aldrich, St. Louis, MO) containing 20% knockout serum replacement (KSR, Life Technologies, Waltham, MA), 2 mM L-

#### Table 2

Summary of antibodies and primers.

glutamine, 0.1 mM non-essential amino acids (NEAA, Invitrogen, Waltham, MA), 0.1 mM 2-mercaptoethanol (2-ME, Life Technologies), and 0.5% penicillin and streptomycin. On Day 8, EBs were seeded onto gelatin-coated plates, and cultivated for 8 days.

#### Immunocytochemical staining

Cells were fixed in 4% paraformaldehyde in PBS for 20 min at room temperature, washed three times with PBS after each step, and blocked with 5% bovine serum albumin for 1 h at room temperature followed by incubation with primary antibodies at 4 °C overnight. The primary antibodies used for staining are listed in Table 2. Suitable secondary antibodies were incubated at room temperature for 1 h. Nuclear staining was also performed with DAPI for 5 min. Images were collected using BIOREVO BZ-9000 (Keyence, Osaka, Japan) or Olympus CKX41 (Olympus, Tokyo, Japan) (Tables 1 and 2).

#### Flow cytometry analysis

iPSCs were dissociated into single cells using Accumax (Innovative Cell Technologies, San Diego, CA) and were incubated at  $1.0 \times 10^6$  cells/ml in PBS with 2% FBS and 20 µl SSEA4 APC conjugated monoclonal antibody (BD Biosciences, Franklin Lakes, NJ) for 30 min at 4 °C. After staining, the cells were washed twice in PBS with 2% FBS, and then stored in PBS with 2% FBS on ice. Cells were analyzed on a FACS Aria II (BD Biosciences) high-speed cell sorter using a 647 nm excitation

Antibodies used for immunocytochemistry/flow cytometry					
	Antibody	Dilution	Company Cat# and RRID		
Pluripotency markers	Rabbit anti-NANOG	1:500	Cosmo Bio Co Cat# REC-RCAB0003P, RRID: AB_1962353		
Pluripotency markers	Mouse anti-SSEA4	1:1000	Millipore Cat# MAB4304, RRID: AB_177629		
Differentiation markers	Goat anti-SOX-17	1:1000	R and D Systems Cat# AF1924, RRID: AB_355060		
Differentiation markers	Mouse anti- $lpha$ SMA	1:500	Dako Cat# M0851, RRID: AB_2223500		
Differentiation markers	Rabbit anti-BIII-tubulin	1:2000	Cell Signaling Technology Cat# 5568P, RRID: AB_10692510		
Secondary antibodies	Donkey anti-Goat IgG Alexa Fluor 488	1:1000	Thermo Fisher Scientific Cat# A-11055, RRID: AB_142672		
Secondary antibodies	Goat anti-Rabbit IgG Alexa Fluor 488	1:1000	Thermo Fisher Scientific Cat# A-11034, RRID: AB_2576217		
Secondary antibodies	Goat anti-Mouse IgG Alexa Fluor 546	1:1000	Thermo Fisher Scientific Cat# A-11030, RRID: AB_2534089		
Secondary antibodies	Goat anti-Rabbit IgG Alexa Fluor 546	1:1,000	Thermo Fisher Scientific Cat# A-11010, RRID: AB_10584649		
Primers					
	Target	Forward/revers	Forward/reverse primer (5'-3')		
Genotyping	GCH1 exon 1	CAGGTGCAGCA	CAGGTGCAGCAATGGGTTCC/CGCACTGACCTGAGATGGTCTCC		
Genotyping	GCH1 exon 2	TTCTCCTTCCTC	TTCTCCTTCCTCTCCATACTGC/TGAGAGCCTTCTGCTACTTTGG		
Genotyping	GCH1 exon 3	CCGCATTTACCAACGGACAAC/CAGGTAAAGAGAGAGAAAGCCTGATG			
Genotyping	GCH1 exon 4	TTCCTCTTGCAGCCCACTTGCTTC/CACAGTGGCTCATGCCTGTAATCC			
Genotyping	GCH1 exon 5	GGGCTTCAGGGTGTTCTGAGA/AGCTTTAGGCTCAGGGATGG			
Genotyping	GCH1 exon 6	ACTTGGTAACTGTGAGCTGAG/GGTGGCAAGAAGAAGAAGTAGAGG			

wavelength and 100 µm nozzle. Unstained cells were also analyzed as negative controls to exclude data from non-specific fluorescence.

#### DNA fingerprinting

STR analysis was performed by BEX (Tokyo, Japan) from PBMCs and the iPSC line.

#### Mycoplasma test

iPSCs were confirmed to be mycoplasma-negative using the MycoAlert kit (Lonza, Basel, Switzerland) in accordance with the manufacturer's instructions.

Supplementary data to this article can be found online at http://dx. doi.org/10.1016/j.scr.2017.07.029.

#### Acknowledgments

We would like to express our sincere gratitude to all of our coworkers and collaborators. We would also like to express our sincere gratitude to Noriko Endo and Ruri Taniguchi for their administrative support. This research was supported in part by a grant from the Program for Intractable Diseases Research utilizing disease-specific iPS cells from the Japan Agency for Medical Research and Development (AMED) to H.I., and from the grant for the Core Center for iPS Cell Research of the Research Center Network for Realization of Regenerative Medicine from AMED to H.I.

#### References

- Nakagawa, M., Taniguchi, Y., Senda, S., Takizawa, N., Ichisaka, T., Asano, K., Morizane, A., Doi, D., Takahashi, J., Nishizawa, M., Yoshida, Y., Toyoda, T., Osafune, K., Sekiguchi, K., Yamanaka, S., 2014. A novel efficient feeder-free culture system for the derivation of human induced pluripotent stem cells. Sci. Rep. 4, 3594.
- Nalls, M.A., Pankratz, N., Lill, C.M., Do, C.B., Hernandez, D.G., Saad, M., DeStefano, A.L., Kara, E., Bras, J., Sharma, M., Schulte, C., Keller, M.F., Arepalli, S., Letson, C., Edsall, C., Stefansson, H., Liu, X., Pliner, H., Lee, J.H., Cheng, R., Ikram, M.A., Ioannidis, J.P.A., Hadjigeorgiou, G.M., Bis, J.C., Martinez, M., Perlmutter, J.S., Goate, A., Marder, K., Fiske, B., Sutherland, M., Xiromerisiou, G., Myers, R.H., Clark, L.N., Stefansson, K., Hardy, J.A., Heutink, P., Chen, H., Wood, N.W., Houlden, H., Payami, H., Brice, A., Scott, W.K., Gasser, T., Bertram, L., Eriksson, N., Foroud, T., Singleton, A.B., 2014. Large-scale meta-analysis of genome-wide association data identifies six new risk loci for Parkinson's disease. Nat. Genet. 56, 1–7.
- Okita, K., Yamakawa, T., Matsumura, Y., Sato, Y., Amano, N., Watanabe, A., Goshima, N., Yamanaka, S., 2013. An efficient nonviral method to generate integration-free human-induced pluripotent stem cells from cord blood and peripheral blood cells. Stem Cells 31, 458–466.
- Segawa, M., Hosaka, A., Miyagawa, F., Nomura, Y., Imai, H., 1976. Hereditary progressive dystonia with marked diurnal fluctuation. Adv. Neurol. 14, 215–233.
- Segawa, M., Nomura, Y., Nishiyama, N., 2003. Autosomal dominant guanosine triphosphate cyclohydrolase I deficiency (Segawa disease). Ann Neurol. 54, S32–S45.