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Lab Resource: Multiple Cell Lines

Generation of human induced pluripotent stem cell lines from two down syndrome patients, including a down syndrome/Alzheimer's disease case (FLENIi002-A) and a beta-amyloid-resistant case (FLENIi003-A)



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ABSTRACT

Two human induced pluripotent stem cell (hiPSC) lines, FLENIi002-A and FLENIi003-A, were generated from peripheral blood mononuclear cells (PBMCs) using the lentiviral-hSTEMCCA-loxP vector. FLENIi002-A was derived from a 52-year-old Down syndrome patient with Alzheimer's disease and amyloid-beta brain accumulation. FLENIi003-A was derived from a cognitively unimpaired 51-year-old Down syndrome patient exhibiting no brain amyloid-beta deposition. Both lines retained the trisomy 21 genotype, and their pluripotency and differentiation potential were confirmed.

(continued)

1. Resource table

Unique stem cell lines identifier	FLENII002-A	Unique stem cell lines identifier	FLENIi002-A FLENIi003-A
	FLENII003-A	Genetic Modification	YES
Alternative name(s) of stem cell lines	FBDS1	Type of Genetic Modification	Trisomy 21 (both lines)
	FBDS2	Evidence of the reprogramming	RT-qPCR
Institution	Instituto de Neurociencias, Fleni (INEU- Fleni-CONICET)	transgene loss (including genomic copy if applicable)	
Contact information of distributor	Leonardo Romorini (lromorini@fleni.org. ar); Ezequiel Surace (esurace@fleni.org.	Associated disease	Down syndrome-Alzheimer's disease (FLENIi002-A)
	ar)		Down syndrome (FLENIi003-A)
Type of cell lines	iPSC	Gene/locus	N/A
Origin	Human	Date archived/stock date	8-11/2022
Additional origin info required	Age: 52 (FLENIi002-A), 51 (FLENIi003-A)	Cell line repository/bank	https://hpscreg.eu/cell-line/F
	Sex: Male (both patients)		LENIi002-A
	Ethnicity if known: Caucasian (both		https://hpscreg.eu/cell-line/FL
	patients)		ENIi003-A
Cell Source	Peripheral blood mononuclear cells	Ethical approval	The study was approved by the local
	(PBMCs) (both lines)		Ethics Committee (Ethics Committee for
Clonality	Clonal		Biomedical Research of the Fleni
Method of reprogramming	Lentiviral EF1a-hSTEMCCA-loxP vector		Institute) (code number:21–16). Written
	expressing OCT-4, SOX-2, c-MYC and		informed consent was obtained from each
	KLF4		patient.
	(continued on next column)		

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Fig. 1. Characterization of the hiPSC lines FLENIi002-A and FLENIi003-A. (A) Karyotype analysis. (B) hiPSC morphology and AP activity. (C) Immunocytochemistry for pluripotency markers SSEA-4, TRA1-60, OCT4, NANOG, and nuclear marker DAPI. (D) Expression of pluripotency markers analyzed by RT-qPCR. (E) Silencing of transgenes introduced by the STEMCCA lentiviral vector confirmed by RT-qPCR. (F) Immunocytochemistry for germ layer markers: mesoderm (SMA and NKX2.5), endoderm (AFP and SOX17), and ectoderm (PAX6 and TUJ1).

Table 1

Characterization and validation.

Classification	Test	Result	Data
Morphology	Photography Bright	Normal (both	Fig. 1 panel B
Phenotype	field Immunocytochemistry	lines) Positive for pluripotency markers OCT-4, NANOG, TRA- 1–60, and SSEA4 (both	Fig. 1 panel C
	RT-qPCR	Cells express OCT-4, SOX-2 and NANOG (both lines)	Fig. 1 panel D
Genotype	Karyotype (G-banding) and resolution	47,XY,+21, both cell lines Resolution: 440	Fig. 1 panel A
mtDNA analysis (IF APPLICABLE)	N/A	N/A	N/A
Identity	Microsatellite PCR (mPCR) OR	N/A	N/A
	STR analysis	26 sites tested, matched	Submitted in archive with journal
Mutation analysis (IF APPLICABLE)	N/A	N/A	N/A
Microbiology and virology	Mycoplasma	Mycoplasma testing by PCR (Negative)	Supplementary file
Differentiation potential	Embryoid body formation	Expression of α -smooth muscle actin (SMA), NK2 homeobox 5 (NKX2.5), SRY- Box Transcription Factor 17 (SOX17), α -fetoprotein (AFP), paired box protein Pax- 6 (PAX6) and Class III β -tubulin (TUJ1) was used as a proof of three germ layers formation (both lines)	Fig. 1 panel F

1.0.1. Resource utility

The FLENIi002-A line was derived from a Down syndrome (DS) patient with Alzheimer's disease (AD) and amyloid-beta deposits, while the FLENIi003-A line was generated from a Down syndrome patient resistant to amyloid-beta (A β) deposition and without signs of Alzheimer's disease. These hiPSC lines provide valuable resources for investigating the factors that confer protection against amyloid-beta pathology in DS.

2. Resource details

DS results from trisomy 21 and is linked to various phenotypic features, including an increased risk for AD due to the triplication of the *APP* gene (located in 21q21). Individuals with DS over 40 years of age typically have A β deposits in their brains. However, some individuals show resistance to these deposits despite advanced age (Lao et al., 2017; Tudorascu et al., 2019; Zammit et al., 2020; Zammit et al., 2021). The FLENIi002-A cell line was derived from a 51-year-old patient with DS who exhibited both AD and A β plaques. In contrast, the FLENIi003-A line was obtained from a 52-year-old patient with DS who demonstrated resistance to A β pathology and showed no clinical signs of AD.

PBMCs were obtained from both patients, and reprogramming into hiPSCs was performed using the EF1a-hSTEMCCA-loxP lentiviral vector expressing OCT-4, SOX-2, c-MYC, and KLF4 pluripotency genes, following a feeder-free protocol (Somers et al., 2010; Chen et al., 2011). Both hiPSC lines retained the trisomy 21 genotype (Fig. 1A). Morphologically, both lines displayed the typical characteristics of hiPSCs, including compact multicellular colonies with a high nucleus-tocytoplasm ratio and clear colony borders. High levels of Alkaline Phosphatase (AP) activity were observed (Fig. 1B). Immunofluorescence staining further validated the robust expression of pluripotency markers OCT-4, NANOG, SSEA-4 and TRA-1-60 (Fig. 1C). RT-qPCR confirmed expression of stemness markers SOX-2, POU5F1 (OCT-4), and NANOG, with a primary culture of human fibroblasts (HF) serving as a negative control (Fig. 1D). The established hiPSC line FN2.1 (Questa et al., 2016) and H9 human embryonic stem cells were used as positive controls. Silencing of the transgenes inserted by the STEMCCA lentiviral vector was confirmed by RT-qPCR using specific primers for exogenous gene expression (Fig. 1E), with hiPSCs (FN2.1) and transduced fibroblasts from day 6 of reprogramming (HF) as negative and positive controls, respectively. In vitro differentiation assays confirmed the pluripotent potential of both lines. Undirected differentiation through the embryoid body method demonstrated their ability to generate the three germ layers, as evidenced by immunofluorescence staining for Smooth Muscle Actin (SMA, mesoderm), NK2 homeobox 5 (NKX2.5, mesoderm), Alpha-Fetoprotein (AFP, endoderm), SRY-Box Transcription Factor 17 (SOX17, endoderm), paired box protein Pax-6 (PAX6, ectoderm) and Class III β -tubulin (TUJ1, ectoderm) (Fig. 1F). (See Tables 1 and 2).

3. Materials and methods

3.1. Reprogramming and cell culture

PBMCs were isolated from blood using a Ficoll density gradient procedure (HISTOPAQUE® SIGMA, #10771). A total of 2×10^6 cells were cultured in 2 mL of erythrocyte expansion media containing StemPro[™]-34-SFM (Gibco[™], #10639011), 50 µg/mL ascorbic acid, 50 ng/mL SCF, 10 ng/mL IL-3, 2 U/mL EPO, 40 ng/mL IGF-1, and 1 µM dexamethasone in 1 well of a 12-well dish at 37 °C, 5 % CO₂. Media was replaced on days 3 and 6. On day 9, cells were transduced with the EF1ahSTEMCCA-loxP lentiviral reprogramming vector at a MOI of 7. Cells were plated on Geltrex-coated dishes (1 %, Gibco™, #A1413202) and further cultured according to the manufacturer's instructions for TeSR™-E7™ (STEMCELL Technologies™, #05914). At day 28 posttransduction, hiPSC colonies were mechanically isolated and expanded on Geltrex-coated dishes in mTeSRTM Plus medium (STEM-CELL Technologies™, #17187501). Cell cultures were passaged every 3 to 4 days using Versene (GibcoTM, #15040066) at a 1:8 ratio, with 10 µM Y-27632 ROCK inhibitor (Cell Signaling Technology, #13624) added for up to 24 h.

3.2. Karyotyping

G-banded chromosome analysis of 20 metaphases from hiPSCs (passages 18 and 10) was performed according to standard protocols..

3.3. Alkaline phosphatase assay

hiPSCs were subjected to alkaline phosphatase staining following the manufacturer's instructions (Sigma, #86R) (passages 17 and 8).

3.4. RNA isolation and RT-qPCR

RNA was extracted using TRIzol (ThermoFisher Scientific,

Table 2

Reagents details.

Antibodies used for immunocytochemistry/flow-cytometr

	Antibody	Dilution	Company Cat #	RRID
Pluripotency Markers	mouse anti-OCT-4 IgG	1:200	Santa Cruz Biotechnology sc-5279	AB_628051
Pluripotency Markers	rabbit anti-NANOG IgG	1:200	Cell Signaling Technology 4903	AB_10559205
Pluripotency Markers	mouse anti-SSEA4 IgG	1:200	Santa Cruz Biotechnology sc-21704	AB_628289
Pluripotency Markers	mouse anti-TRA-1-60 IgM	1:200	Santa Cruz Biotechnology sc-21705	AB_628385
Differentiation Markers	mouse anti-AFP IgG	1:200	Santa Cruz Biotechnology sc-166325	AB_2305278
Differentiation Markers	rabbit anti-SMA IgG	1:200	Abcam ab124964	AB_11129103
Differentiation Markers	rabbit anti-SOX17 IgG	1:100	Invitrogen PA5-72815	AB_2718669
Differentiation Markers	rabbit anti-NKX2.5 IgG	1:100	Invitrogen PA5-49431	AB_2634885
Differentiation Markers	rabbit anti-PAX6 IgG	1:100	Abcam ab195045	AB_2750924
Differentiation Markers	rabbit anti-TUJ1 IgG	1:400	Covance Antibody Products MMS-435P	AB_2313773
Secondary antibodies	Goat anti-Mouse IgG Alexa Fluor 594	1:400	Thermo Fisher Scientific A-11005	AB_2534073
Secondary antibodies	Goat anti-Mouse IgG Alexa Fluor 488	1:400	Thermo Fisher Scientific A-11001	AB_2534069
Secondary antibodies	Goat anti-Mouse IgM Alexa Fluor 488	1:400	Thermo Fisher Scientific A-21042	AB_2535711
Secondary antibodies	Goat anti-Rabbit IgG Alexa Fluor 594	1:400	Thermo Fisher Scientific A-11012	AB_2534079
Secondary antibodies	Goat anti-Rabbit IgG Alexa Fluor 488	1:400	Thermo Fisher Scientific A-11008	AB_143165
	Primers			
	Target	Size of band	Forward/Reverse primer (5'-3')	
Exogenous factors (RTPCR)	STEMCCA plasmid OCT-4/KLF4	561 bp	CAACGAGAGGATTTTGAGGC/ATCGTTGAACTCCTCGGTCTCTCT	
Exogenous factors (RTPCR)	STEMCCA plasmid SOX-2/c-MYC	550 bp	TTGGCTCCATGGGTTCGGTG/AAGGGTGTGACCGCAACGTAGG	
Pluripotency Markers (qPCR)	POU5F1 (OCT-4)	105 bp	GCAGGCCCGAAAGAGAAAGCGA/TGGCTGATCTGCTGCAGTGTGG	
Pluripotency Markers (qPCR)	uripotency Markers (aPCR) SOX-2 110 bp AGCATGGAGAAAACCCGGTACGC/		AGCATGGAGAAAACCCGGTACGC/	
		-	CGTGAGTGTGGATGGGATTGGTGT	
Pluripotency Markers (qPCR)	NANOG	120 bp	TCCTTCCTCCCCCTCCCCAT/TAGGCTCCAACCATACTCCACCCTC	

138 bp

#15596026), and cDNA was synthesized from total RNA using Oligo (dT) primers and M–MLV Reverse Transcriptase (Promega, #M170B) following the manufacturer's instructions. qPCR amplification (95 °C for 15 s, 60 °C for 1 min, 40 cycles) and analysis were performed using the StepOnePlus Real-Time PCR System (PE Applied Biosystems). The FastStart Universal SYBR Green Master Mix (ROX) (Roche, #04913850001) was used for all reactions. Transgene expression was assessed using primers spanning two of the four transgenes on the STEMCCA lentiviral vector sequence, allowing amplification of only cDNA generated from the exogenous mRNA expression of these genes. Cells were analyzed at passages between 10 and 18. Values were normalized to levels of the housekeeping gene *RPL7*.

RPL7

3.5. In vitro differentiation

House-Keeping Genes (aPCR)

hiPSCs (passages 12 and 20) were detached using Dispase (GibcoTM, #17105041) and transferred to non-adherent Petri dishes in DMEM/F12 with 1x GlutaMAX (Gibco, #35050061), 20 % KnockOutTM Serum Replacement (GibcoTM, #10828), 1x non-essential amino acids (Sigma, #M7145), 0.1 μ M 2-mercaptoethanol, 100 U/ml penicillin, and 50 μ g/ml streptomycin to induce the formation of embryoid bodies (EBs). On day 4, EBs were plated onto 0.1 % gelatin-coated 24-well plates and cultured for an additional 17 days in DMEM/F12 supplemented with 20 % FBS (Gibco, #10270106), 1x GlutaMAX, 100 U/ml penicillin, and 50 μ g/ml streptomycin.

3.6. Immunofluorescence staining

Cells were fixed with 4 % paraformaldehyde for 30 min and permeabilized with 0.1 % Triton X-100 for 30 min. Primary antibody incubation was performed overnight at 4 °C in PBS with 1 % BSA. Fluorescent dye-conjugated secondary antibodies were incubated in PBS with 1 % BSA for 1 h at room temperature. Cells were counterstained with DAPI and examined under a EVOS FLoid (Thermo Fisher Scientific) inverted microscope. Pluripotent markers were analyzed at passages 17 and 9.

3.7. STR analysis

Genomic DNA was isolated using the Wizard Genomic DNA Purification kit (Promega). STR analysis of 26 *loci* was performed at the Laboratorio de Huellas Digitales Genéticas (Facultad de Farmacia y Bioquímica, UBA, Buenos Aires, Argentina) (passages 18 and 9).

3.8. Statistical analysis

Results are expressed as mean \pm SEM (n = 3). Comparisons were analyzed by Two-tailed Student's *t*-test (* p < 0.05; ** p < 0.01; **** p < 0.0001).

CRediT authorship contribution statement

AATGGCGAGGATGGCAAG/TGACGAAGGCGAAGAAGC

Giulia S. Clas: Validation, Methodology, Formal analysis. Mariela Marazita: Validation, Methodology, Formal analysis. Lucía Pertierra: Resources, Investigation. Silvia Vazquez: Investigation. Luciana Isaja: Methodology. Soledad Rodríguez-Varela: Methodology. Sofía Mucci: Methodology. Belén Helou: Investigation. Fernanda Tapajóz: Investigation. Nahuel Magrath: Investigation. Fernanda Tapajóz: Investigation, Investigation, Resources. Ricardo Allegri: Investigation. Gustavo E. Sevlever: Investigation. Maria E. Scassa: Methodology, Investigation, Ezequiel I. Surace: Writing – review & editing, Supervision, Investigation, Conceptualization. Leonardo Romorini: Writing – review & editing, Supervision, Project administration, Investigation, Funding acquisition.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.scr.2025.103728.

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References

- Chen, G., Gulbranson, D.R., Hou, Z., Bolin, J.M., Ruotti, V., Probasco, M.D., Smuga Otto, K., Howden, S.E., Diol, N.R., Propson, N.E., Wagner, R., Lee, G.O., Antosiewicz Bourget, J., Teng, J.M.C., Thomson, J.A., 2011. Chemically defined conditions for human iPSC derivation and culture. Nat. Methods. 8 (5), 424–429.
- Lao, P.J., Handen, B.L., Betthauser, T.J., et al., 2017. Longitudinal changes in amyloid positron emission tomography and volumetric magnetic resonance imaging in the nondemented Down syndrome population. Alzheimer's Dement: Diagn Assess Dis. Monit. 9, 1–9.
- Questa, M., Romorini, L., Blüguermann, C., Solari, C.M., Neiman, G., Luzzani, C., Scassa, M.E., Sevlever, G.E., Guberman, A.S., Miriuka, S.G., 2016. Generation of iPSC line iPSC-FH2.1 in hypoxic conditions from human foreskin fibroblasts. Stem Cell Res. 16 (2), 300–303.
- Somers, A., Jean, J.C., Sommer, C.A., Omari, A., Ford, C.C., Mills, J.A., Ying, L., Sommer, A.G., Jean, J.M., Smith, B.W., Lafyatis, R., Demierre, M.F., Weiss, D.J., French, D.L., Gadue, P., Murphy, G.J., Mostoslavsky, G., Kotton, D.N., 2010. Generation of transgene-free lung disease-specific human iPS cells using a single excisable lentiviral stem cell cassette. Stem Cells. 28 (10), 1728–1740.

Tudorascu, D.L., Anderson, S.J., Minhas, D.S., et al., 2019. Comparison of longitudinal $A\beta$ in nondemented elderly and Down syndrome. Neurobiol Aging. 73, 171–176.

- Zammit, M., Laymon, C.M., Betthauser, T.J., et al., 2020. Amyloid accumulation in Down syndrome measured with amyloid load. Alzheimer's Dement: Diagn Assess Dis Monit. 12, e12020.
- Zammit, M.D., Tudorascu, D.L., Laymon, C.M., et al., 2021. PET measurement of longitudinal amyloid load identifies the earliest stages of amyloid-beta accumulation during Alzheimer's disease progression in Down syndrome. Neuroimage 228, 117728.