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Lab Resource: Single Cell Line

Generation of a human induced pluripotent stem cell line (UALGi001-A) from a patient with Left-Ventricular Noncompaction Cardiomyopathy

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ABSTRACT

Left Ventricular Noncompaction Cardiomyopathy (LVNC) is characterized by excessive trabeculation of the left ventricle. To date, mutations in more than 40 genes have been associated with LVNC, however the exact mechanisms underlying the disease remain unknown. Here, we describe an induced pluripotent stem cell (iPSC) line (UALGi001-A) from a LVNC patient (LVNC-iPSC) that does not present mutations in the genes most commonly associated with the disease (van Waning et al., 2019). The LVNC-iPSC exhibited full pluripotency and differentiation potential, and retained a normal karyotype after reprogramming. This in vitro cellular model will be useful to study the molecular, genetic and functional aspects of LVNC.

1. Resource Table:

Unique stem cell line identifier	UALGi001-A		
Alternative name(s) of stem cell line	VITAL16		
Institution	Universidade do Algarve, Faro, Portugal Algarve		
	Biomedical Center Research Institute (ABC-RI),		
	Faro, Portugal		
Contact information of	jebraganca@ualg.pt		
distributor			
Type of cell line	iPSC		
Origin	Human		
Additional origin info required	Age: 75		
for human ESC or iPSC	Sex: Male		
	Ethnicity: Caucasian/Portuguese		
Cell Source	Peripheral blood mononuclear cells		
Clonality	Clonal		
Method of reprogramming	Non-integrating, Sendai Virus expressing hOCT3/4,		
	hSOX2, hKLF4 and hC-MYC		
Genetic Modification	NO		
Type of Modification	N/A		
Associated disease	Left Ventricular Non-Compaction Cardiomyopathy		
	(LVNC)		
Gene/locus	N/A		
Date archived/stock date	July 2020		
Cell line repository/bank	https://hpscreg.eu/cell-line/UALGi001-A		
	(continued on next column)		

(continued)

Unique stem cell line identifier	UALGi001-A
Ethical approval	Approval ID: 78/19 by Ethics Committee for Health, Centro Hospital e Universitário do Algarve, Faro, Portugal

2. Resource utility

Left Ventricular Noncompaction Cardiomyopathy (LVNC) is a heart disorder characterized by endomyocardial noncompaction. UALGi001-A cell line, derived from a patient with LVNC offers a useful tool to study molecular and cellular mechanisms involved in this cardiomyopathy.

3. Resource details

Mononucleated cells were collected from 4 mL of peripheral blood sample from 75-year-old male. The patient was clinically diagnosed with left ventricular trabeculations located in the middle and apical segments and deep recesses, compatible with Left Ventricular Non-Compaction Cardiomyopathy (LVNC). LVNC is the third most prevalent heart muscle disease, accounting for almost 10% of all cardiomyopathies and it is

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characterized by the presence of a vast trabeculations likely due to endomyocardial morphogenesis arrest during embryogenesis (van Waning et al., 2019). LVNC clinical manifestations include heart failure, thromboembolism, ventricular arrhythmias, and ultimately sudden cardiac death (Finsterer et al., 2017). The human induced pluripotent stem cell (iPSC) line UALGi001-A was generated by using the Cyto-Tune® iPSC-Reprogramming kit (Thermo Fisher Scientific, Invitrogen), encoding for the reprogramming factors hOCT3/4, hc-MYC, hKLF4, and hSOX2 (Takahashi and Yamanaka, 2006), according to manufacturer's instructions. Clonal iPSC lines were established and further characterized (Table 1), formed colonies with a standard stem-like morphology visible by phase contrast, and were positive for Alkaline Phosphatase (AP) activity (Fig. 1A-B). The expression of pluripotency markers was demonstrated by immunofluorescence staining of SOX2, NANOG, OCT4 and SSEA-4 (Fig. 1 C), as well as flow cytometry (Fig. 1D). Viral clearance was confirmed at passage 14 (Supplementary Fig. S1A).

Genomic integrity was assessed by karyotype analysis, showing that UALGi001-A, at passage 21, presented a normal diploid (46, XY) chromosomal content (Fig. 1E). The capacity of iPSCs to differentiate into three germ layers was confirmed by embryoid body (EB) differentiation assay, and expression of ectoderm (TUJ1), endoderm (AFP) and mesoderm (α -SMA) markers (Fig. 1F). DNA fingerprinting was used to prove the genetic identity to parental mononucleated blood cells (archived with journal). Mycoplasma was regularly tested negative throughout cell culture indicating that UALGi001-A line is mycoplasma-free (Supplementary Fig. S1B).

Table 1

Characterization and validation.

Classification	Test	Result	Data
Morphology	Photography Bright field	Normal	Fig. 1A
Phenotype	Qualitative analysis (Immunocytochemistry)	Positive pluripotency markers: Alkaline Phosphatase, Oct3/4, Nanog, Sox2, and SSEA- 4	Fig. 1B and C
	Quantitative analysis (Flow cytometry)	Percentage of NANOG positive cells: 95%; SSEA- 4 positive cells: 93,3%	Fig. 1D
Genotype	Karyotype (G-banding) and resolution	46XY, Resolution 400–500	Fig. 1E
Identity	Microsatellite PCR (mPCR)	Not performed	N/A
	STR analysis	16 loci analyzed, all matched	Supplementary table 1
Mutation	Sequencing	N/A	N/A
analysis (IF APPLICABLE)	Southern Blot OR WGS	N/A	N/A
Microbiology and virology	Mycoplasma	Mycoplasma testing by RT- PCR: Negative	Supplementary Fig. 1
Differentiation potential	e.g. Embryoid body formation	Positive staining for α-fetoprotein (AFP), βIII- tubulin (TUBB3), α-smooth muscle actin (SMA).	Fig. 1F
Donor screening (OPTIONAL)	HIV 1 + 2 Hepatitis B, Hepatitis C	Not performed	N/A
Genotype additional	Blood group genotyping	Not performed	N/A
info (OPTIONAL)	HLA tissue typing	Not performed	N/A

4. Materials and methods

4.1. Reprogramming of PBMCs

PBMCs were isolated using the Vacutainer® CPTTM tubes (BD Biosciences). PBMCs were cultured (1.0 \times 10⁶ cells) in PBMCs Medium (StemPro-34 SFM medium; 1% GlutaMAX; 1% Pen/Strep, 100 ng/mL SCF, 100 ng/mL FLT-3; 20 ng/mL IL-3, 20 ng/mL 20 ng/mL IL-6; 25 ng/ mL GM-CSF), for one week before transduction with CytoTune®-iPS 2.0 Sendai Reprogramming Kit. Briefly, 0.25×10^5 cells were transduced using MOI of 5-5-3 (hKOS, hc-MYC, hKLF4, respectively). After 24 h of incubation, cells were collected, centrifuged, and seeded in a 24-well plate containing PBMCs Medium. Two days later, cells were transferred onto a 6-well plate coated with Geltrex LDEV-Free Reduced Growth Factor Basement Membrane Matrix (Gibco) in ReproTeSR (STEMCELL Technologies), supplemented with 1% Pen/Strep. Individual colonies with stem-like morphology were manually isolated and expanded 14 to 21 days post-transduction. iPSC clones were cultured and expanded on 6-well plates coated with Geltrex LDEV-Free Reduced Growth Factor Basement Membrane Matrix, using a slip ratio of 1:5 and were maintained at 37 °C in humidified atmosphere containing 5% CO₂.

4.2. In vitro differentiation assay

In vitro differentiation was performed by EB formation. iPSC colonies were lifted manually and cultured in non-adherent conditions in mTeSR1 medium, containing 0.4% of polyvinyl alcohol, for 48 h. Thereafter, the EBs were seeded on glass coverslips coated Geltrex LDEV-Free Reduced Growth Factor Basement Membrane Matrix and cultured for 3 weeks in differentiation medium (DMEM, 10% FBS, 1% Pen/Strep, 1% GlutaMAX, 1% MEM-NEAA). The coverslips were fixed with 4% paraformaldehyde (PFA) for 15 min and analyzed by immunofluorescence (Table 2) on Axioimager Z2/Apotome fluorescence microscope (Carl Zeiss).

4.3. Immunocytochemistry

Cells were allowed to grow in glass coverslips coated with Geltrex LDEV-Free Reduced Growth Factor Basement Membrane Matrix and washed in ice-cold PBS before fixation in 4% PFA, for 15 min. Fixed cells were washed twice in PBS and placed in blocking solution (2% bovine serum albumin in 0.2% Triton-X100/PBS) for 1 h, at room temperature. Cells were incubated for 1 h at room temperature with the primary antibody (Table 2). After incubation, samples were washed 3 times with 0.2% Triton X100/PBS, and incubated with the secondary antibodies for 1 h, at room temperature (Table 2). After 3 washes, sections were mounted with Fluoromount G mounting medium (Thermo Fisher Scientific) containing 4,6-diamidino-2-phenylindole (DAPI) and analyzed on a Axioimager Z2 fluorescence microscope (Carl Zeiss)

4.4. Flow cytometry

iPSCs were dissociated using TrypLE Select (Gibco) for 3 min at RT, centrifuged at 300 g for 5 min and 100,000 cells resuspended in 200 μ l of ice-cold 0.5 % PFA in PBS, for 20 min. Fixed cells were washed twice in PBS/0.5% BSA/0.1% Triton X100 and incubated with the primary antibody for 1 h at 4 °C (Table 2). After incubation, samples were washed 3 times with PBS/0.5% BSA/0.1% Triton X100, and incubated with the secondary antibodies for 1 h, at 4 °C (Table 2), protected from light. The cells were analyzed using a FACScalibur cell analyzer (BD Biosciences) and data was analyzed by CytExpert 2.0 software.

4.5. Alkaline phosphatase activity

Alkaline phosphatase staining was carried out using Alkaline Phosphatase Staining Kit II (Stemgent, MA).



Fig. 1. Characterisation of the induced pluripotent stem cell line (UALGi001-A) from a patient with LVNC. (A) Phase contrast micrograph of UALGi001-A colony cultured in feeder-free conditions. (B) Alkaline phosphatase positive staining. CC) Flow cytometry of nuclear (NANOG) and surface (SSEA-4) pluripotency markers. (D) Immunofluorescence for pluripotency markers OCT4, SOX2, NANOG, and SSEA-4. Nuclei were counterstained with DAPI (blue). (E) Representative metaphase showing normal diploid 46, XY karyotype. (F) Immunocytochemistry for ectodermal (TUJ1), endodermal (AFP) and mesodermal (αSMA) markers. Nuclei were counterstained with DAPI (blue).

4.6. Karyotype analyses

Genome integrity of the iPSC was evaluated by G-banding at 400–550 band resolution, with a minimum of 30 metaphase spreads analyzed (Genomed, Lisbon, Portugal).

4.7. RT-PCR for detection of viral clearance

Total RNA was isolated from cultured iPSC with RNeasy Mini Kit (Qiagen). 1 μ g of total RNA was used as template to obtain cDNA, using NZY First-Strand cDNA Synthesis Kit (nzytech). Viral clearance was

Table 2 Reagents details.

Antibodies used for immunocytochemistry/flow-cytometry				
Antibody	Dilution	Company Cat #	RRID	
Rabbit anti-OCT4	1:400	Cell Signaling Technology Cat# 2840	RRID: AB_2167691	
Rabbit anti-NANOG	1:400	Cell Signaling Technology Cat# 4903	RRID: AB_10559205	
Rabbit anti-SOX2	1:400	Cell Signaling Technology Cat# 3579	RRID: AB_2195767	
Mouse anti-SSEA-4	1:200	BD Biosciences Cat# 560,073	RRID: AB_1645601	
Mouse anti-TUBB3	1:500	BioLegend Cat#801213	RRID: AB_2728521	
Mouse anti-ASM	1:500	Sigma-Aldrich Cat#A5228	RRID: AB_262054	
Rabbit anti-AFP	1:100	Dako Cat# A0008	RRID: AB_2650473	
Donkey anti-Mouse 488	1:500	Molecular Probes Cat# A-21202	RRID: AB_141607	
Donkey anti-Rabbit 594	1:500	Molecular Probes Cat# A-21207	RRID: AB_141637	
Donkey anti-Rabbit 488	1:200	Molecular Probes Cat# A-21206	RRID: AB_2535792	
Primers				
Target	Size of band	Forward/Reverse primer (5'-3')		
SeV Plasmid	181 bp	GGATCACTAGGTGATATCGAGC/		
hKLF4 Plasmid	410 bp	ACCAGACAAGAGTTTAAGAGATATGTA	ГС	
hKOS Plasmid	528 bp	TTCCTGCATGCCAGAGGAGCCC/		
hC-MYC Plasmid	532 bp	AATGTATCGAAGGTGCTCAA		
		ATGCACCGCTACGACGTGAGCGC/		
		ACCTTGACAATCCTGATGTGG		
		TAACTGACTAGCAGGCTTGTCG/		
		TCCACATACAGTCCTGGATGATGATG		
M. arginine	500 bp	Forward primers:		
M. boris		CGC CTG AGT AGT ACG TTC GC		
M. fermentans		CGC CTG AGT AGT ACG TAC GC		
M. hominis		TGC CTG AGT AGT ACA TTC GC		
M. hyorhinis		CGC CTG GGT AGT ACA TTC GC		
M. orale		CGC CTG AGT AGT ATG CTC GCTGC CTG GGT AGT ACA TTC GC		
		Reverse primers:		
		GCG GTG TGT ACA AGA CCC GA		
		GCG GTG TGT ACA AAA CCC GA		
		GCG GTG TGT ACA AAC CCC GA		
	Antibodies used for imm Antibody Rabbit anti-OCT4 Rabbit anti-NANOG Rabbit anti-NANOG Rabbit anti-SOX2 Mouse anti-SEA-4 Mouse anti-SEA-4 Mouse anti-AEP Donkey anti-ASM Rabbit anti-AFP Donkey anti-Rabbit 594 Donkey anti-Rabbit 488 Primers Target SeV Plasmid hKLF4 Plasmid hKOS Plasmid hC-MYC Plasmid hC-MYC Plasmid hC-MYC Plasmid M. arginine M. boris M. fermentans M. horininis M. hyorhinis M. orale	Antibodies used for immunocytochemistAntibodyDilutionRabbit anti-OCT41:400Rabbit anti-NANOG1:400Rabbit anti-SOX21:400Mouse anti-SSEA-41:200Mouse anti-ASM1:500Mouse anti-ASM1:500Donkey anti-Mouse 4881:500Donkey anti-Rabbit 5941:500Donkey anti-Rabbit 5941:500PrimersTargetSize of band528 bphKLF4 Plasmid410 bphKOS Plasmid532 bpM. arginine500 bpM. borisM. fermentansM. hyorhinisM. orale	Antibodies used for immunocytochemistry/flow-cytometryAntibodyDilutionCompany Cat #Rabbit anti-OCT41:400Cell Signaling Technology Cat# 2840Rabbit anti-NANOG1:400Cell Signaling Technology Cat# 4903Rabbit anti-SOX21:400Cell Signaling Technology Cat# 3579Mouse anti-SSEA-41:200BD Biosciences Cat# 560,073Mouse anti-TUBB31:500BioLegend Cat#801213Mouse anti-ASM1:500Sigma-Aldrich Cat#A5228Rabbit anti-AFP1:100Dako Cat# A0008Donkey anti-Mouse 4881:500Molecular Probes Cat# A-21202Donkey anti-Rabbit 5941:500Molecular Probes Cat# A-21207Donkey anti-Rabbit 5941:500Molecular Probes Cat# A-21206PrimersTargetSize of bandForward/Reverse primer (5'-3')Sev Plasmid181 bpGGATCACTAGGTGGTATAGAGAC/hKLF4 Plasmid410 bpACCAGACAAGAGTTTAAGAGATATGTAChKOS Plasmid532 bpATTGCTGCATGCCAGAGGGGC/hC-MYC Plasmid532 bpATGCACGCTACGAGCGC/AACCTGGATGATGGM. arginine500 bpForward primers:M. borisCGC CTG AGT AGT ACG TTC GCM. hominisTG CCTG AGT AGT ACG TTC GCM. nominisCGC CTG AGT AGT ACA TTC GCM. oraleCGC CTG AGT AGT ACA TTC GCM. oraleCGC CTG AGT AGT ACA TTC GCM. oraleCGC CTG AGT AGT ACA CCC GAGCG GTG TGT ACA AAA CCC GA	

analyzed using the primers described in Table 2. RT-PCR reaction was performed using DreamTaq DNA Polymerase (Thermo Scientific) and PCR products were visualized on a 2% agarose gel.

4.8. Fingerprinting

Genomic DNA from PBMC and iPSC was extracted using QIAamp DNA Blood mini kit (Qiagen). Fingerprinting analyses was performed using Promega's PowerPlex 16 kit and analyzed on ABI PRISM 3100 using GeneMapper (Thermo Fisher) by STABVida, Lisbon, Portugal.

4.9. Mycoplasma detection

The presence of mycoplasma was tested regularly by PCR (Uphoff and Drexler, 2001) using the Primers listed in Table 2.

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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.2021.102302.

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