

## Supplementary Methods

### S1.0 Generation of hiPSC from peripheral blood.

All work using hiPSC in this study was approved under REB# 2020-6362 from MUHC institutional review board. Blood was collected from two patients diagnosed with dilated cardiomyopathy and one healthy control volunteer with no known cardio-pulmonary disease (see **Table S2**). To generate hiPSCs from these subjects, peripheral blood mononuclear cells were reprogrammed using an episomal plasmid-based strategy. Briefly the Epi5™ Episomal iPSC Reprogramming Kit (ThermoFisher) and Neon Transfection System (Invitrogen) were used to introduce pluripotency factors Oct4, Sox2, Lin28, Klf4, and L-Myc using electroporation. Successful reprogramming was determined by attachment to the cell culture dish and confirmed by immunostaining expression of pluripotency genes. To ensure genetic integrity, hiPSCs were karyotyped using G-banding technique (Cytogenomics platform, SickKids, Toronto).

### S1.1 Differentiation of hiPSCs into cardiomyocytes.

Differentiation of hiPSCs into cardiomyocytes was accomplished via the modulation of Wnt signaling using the previously published GiWi protocol with slight modifications. [35]. Briefly, on day 0 hiPSCs were treated with 12  $\mu$ M CHIR99021 to activate the WNT signalling pathway. The following day, media was replaced with RPMI1640 supplemented with B27 without insulin. On day 3 the WNT signalling pathway was inhibited using 5  $\mu$ M IPW2. On day 5 media was replaced with RPMI1640 supplemented with B27 without insulin and on day 7, media was replaced with RPMI1640 supplemented with B27 with insulin. The cell population was purified using 1M lactate to select for hiPSC-CMs. Successful differentiation of hiPSCs into cardiomyocytes (CMs) was confirmed through immunofluorescence and via the expression of the cardiomyocyte-specific biosensor RGEco-TnT (Figs. 6 and 7). Experiments were conducted 28-35 days after cells began to beat to allow maturation.

### S1.2 Seeding in a 96-well microwell plate.

*Note: Ideally, the assay microplate should have an optical bottom compatible with fluorescent microscopy and black walls to reduce noise during measurements.*

1. Coat a black optical bottom 96-well plate with 50  $\mu$ L of a human plasma fibronectin dilution (1:100 in PBS). Allow to polymerize for at least 1 hour at 37°C.
2. Add 500  $\mu$ L of pre-warmed Accutase solution per well and incubate at 37°C for 10-30 minutes. Check on cells periodically to see when cells detach.
3. Following incubation, add 500  $\mu$ L of plating media (44 mL RPMI1640 + 5 mL KOSR + 1 mL B27 + 50  $\mu$ L Y-27632ROCK inhibitor) [35]. Gently pipette up and down with a 1000  $\mu$ L filter tip, ensuring all cells detach. Collect the cell suspension in a 50 mL conical tube.
4. Count the cells using trypan blue.
5. Aspirate the human plasma fibronectin coating solution and seed cells at 20,000 hiPSC-CMs per well in a 96-well plate. Complete the volume to 100  $\mu$ L per well with plating media.
6. The next day, rinse cells 3 times with basal RPMI1640 media to remove cell debris. Refresh media with RPMI + B27. Maintain cells in RPMI1640 + B27 by changing the media every 3-4 days.

## Supplemental Figure Legends

**Supplemental Figure S1. Validation of hiPSCs used in this study.** (A) Immunofluorescence staining of core pluripotency markers demonstrating the “stemness” of HID041004 post-reprogramming. (B) Karyotyping analysis of HID041004 validating normal diploid human female karyotype. (C) Immunofluorescence staining of core pluripotency markers demonstrating the “stemness” of HID041020 post-reprogramming. (D) Karyotyping analysis of HID041020 validating normal diploid human male karyotype. (E) Immunofluorescence staining of core pluripotency markers demonstrating the “stemness” of HID041021 post-reprogramming. (F) Karyotyping analysis of HID041021 validating normal diploid human female karyotype.

**Supplemental Figure S2. Images of instruments used in this study.** (A) Zeiss Axio Observer fully automated inverted microscope (B) with the temperature- and gas-controlled stage insert for 96-well plates. (C-E) Auxiliary components to use with the Axio Observer microscope. When initializing the system, the gas- and temperature- controller must be turned on first. Then, the Power Supply unit, SMC stage controller, and Focus Controller are turned on in this order. Finally, the X-Cite LED source is turned on and set to 20% power.

**Supplemental Figure S3. Visualization of OriginPro software.** User interface shown here.

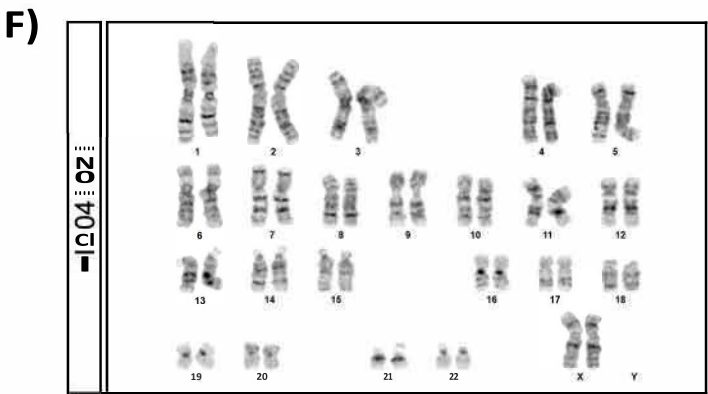
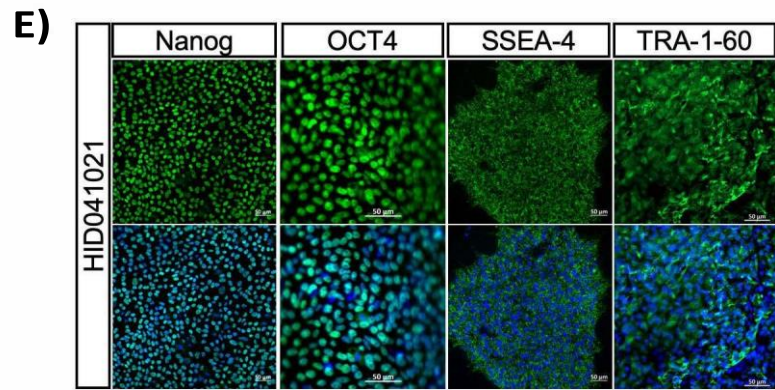
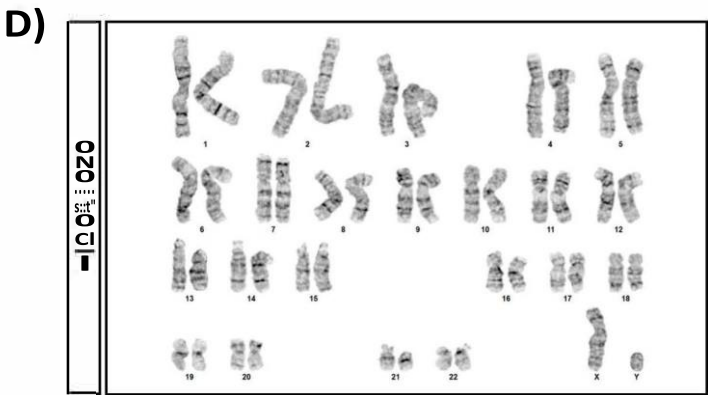
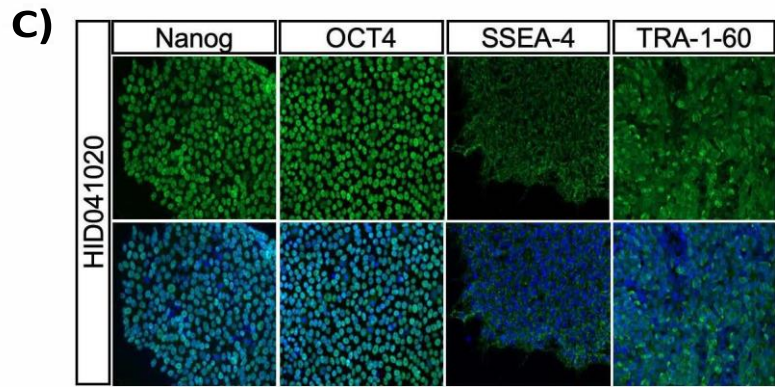
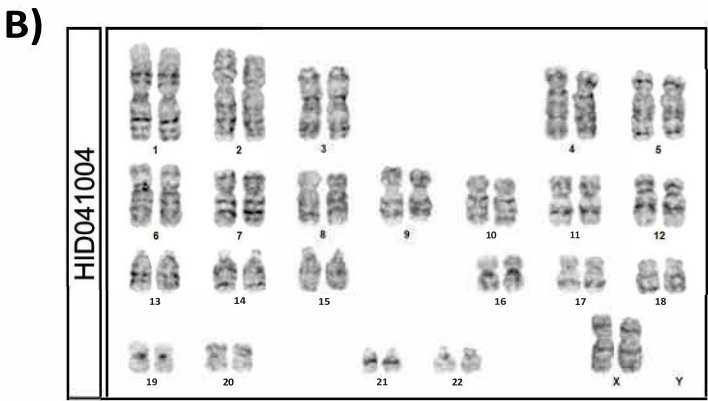
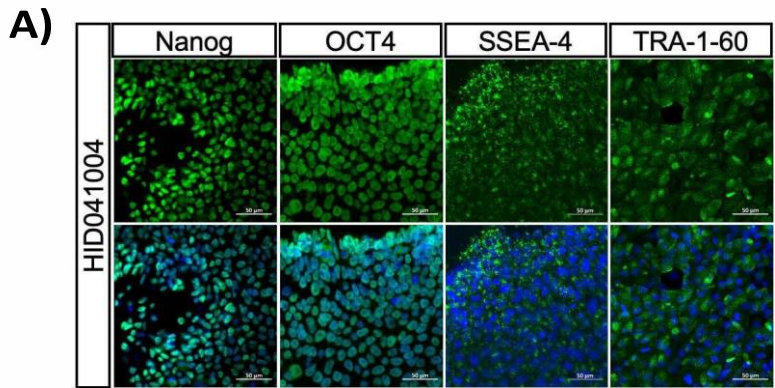
**Supplemental Table S1.** Equipment and software used in calcium imaging and analysis.

Name	Details	Supplier
<i>Equipment</i>		
Inverted microscope	Zeiss Axio Observer	Zeiss
Camera	Zeiss 506 monochrome CCD	Zeiss
Temperature and gas controller	LCI CU-501	Live Cell Instrument
Power supply unit	Power Supply 232	Zeiss
Stage controller	SMC 2009	Zeiss
Definite focus controller	Focus Controller.2	Zeiss
LED source	X-Cite® 120 LED	Excelitas Technologies
Filter cube	Filter Set 14 (item no. 488014-0000-000)	Zeiss
20x objective lens	20x PLAN APOCHROMAT, NA 0.8	Zeiss
<i>Software</i>		
Microscopy software	Zen Blue 3.3	Zeiss
Image visualization	ImageJ (1.53p) with the Bio-Formats plugin	National Institutes of Health
Transient analysis, graphing, and statistics	OriginPro 2022 (9.9.0.225; Student version)	OriginLab

**Supplemental Table S2.** Detailed information on hiPSCs used in this study.

	Biological sex	Age at study entry	Disease status
HID041004	Female	45	Control
HID041020	Male	65	DCM
HID041021	Female	44	DCM

Supplemental Figure S1



Supplemental Figure S2





# Supplemental Figure S3

Transient data  
A(X1) = Time  
B(Y1) = dF/F0

Computed transient data  
(Peak max, peak begin,  
peak end)

Python packages required  
to run the application

Transient identification parameters:  
Window size (adjust to change number of  
peaks recognized)  
Tol\_Peakstart (beginning of the transient)  
Tol\_peakend (end of transient)

Install Python packages

Window Size (2~100) 12 Tol\_peakstart(0~20) 3 Tol\_peakend (0~20) 3

Find Max/Min Calculate

Find Max/Min  
(apply identification  
settings above)  
Calculate  
(recalculate the  
computed results)

Computed results

Graphed  
transient

	A(X1)	B(Y1)	C(Y1)	D(X2)	E(Y2)	F(Y2)	G(X3)	H(Y3)	I(Y3)	J(X4)	K(Y4)	L(Y4)	M(Y4)	N(Y4)
Long Name		Cell	dx_Max		Max	idx_b		peak b	idx_e		peak e			
Units														
Comments														
F(x)														
1	0.096	0.449	32	3.1731	0.6409	22	2.2115	0.2934	63	6.154	0.292	time to peak	1.02564 sec	
2	0.192	0.408	73	7.1154	0.629	63	6.1538	0.2920	103	10	0.292	time to 90% bottom	1.18931 sec	
3	0.288	0.375	115	11.154	0.623	103	10	0.2921	143	13.85	0.295	time to 50% bottom	0.59813 sec	
4	0.385	0.345										time between maximum	3.99038 sec	
5	0.481	0.335										transient duration	3.87821 sec	
6	0.577	0.321										average peak height	0.33778 dF/F0	
7	0.673	0.314										average peak area	0.48823 dF/F0	
8	0.769	0.310										decay tau	0.79722 sec	
9	0.865	0.3057										upstroke velocity	0.33229 dF/(F0*sec)	
10	0.962	0.305										transient frequency	15.03614 min-1	
11	1.058	0.293												
12	1.154	0.300												
13	1.25	0.295												
14	1.346	0.295												
15	1.442	0.297												
16	1.538	0.288												
17	1.635	0.293												
18	1.731	0.2956												
19	1.827	0.299												
20	1.923	0.293												
21	2.019	0.295												
22	2.115	0.293												
23	2.212	0.293												
24	2.308	0.346												
25	2.404	0.454												
26	2.5	0.532												
27	2.596	0.557												
28	2.692	0.580												
29	2.788	0.598												
30	2.885	0.612												
31	2.981	0.617												
32	3.077	0.628												
33	3.173	0.6409												
34	3.269	0.639												
35	3.365	0.638												
36	3.462	0.623												
37	3.558	0.607												
38	3.654	0.562												
39	3.75	0.5099												
40	3.846	0.468												
41	3.942	0.428												
42	4.038	0.401												