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Authors

Liu, Chun
Medina, Pedro
Thomas, Dilip
et al.

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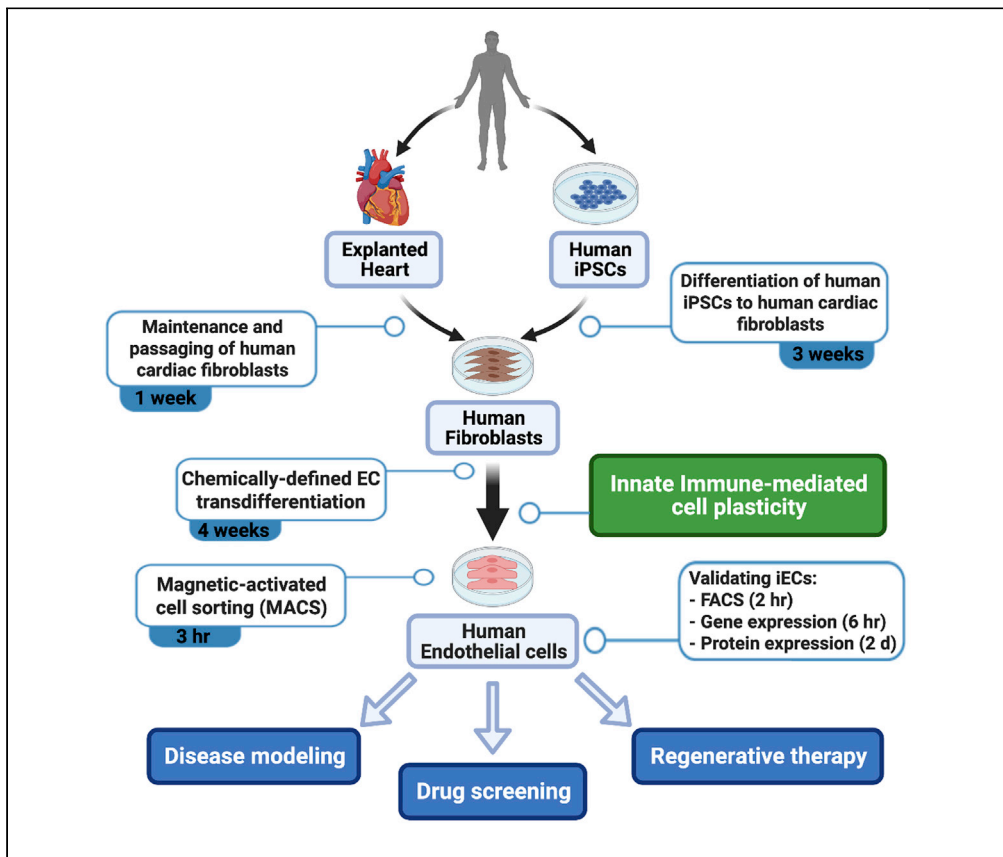
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Protocol

A protocol for transdifferentiation of human cardiac fibroblasts into endothelial cells via activation of innate immunity



Endothelial cells (ECs) have emerged as key pathogenic players in cardiac disease due to their proximity with cardiomyocytes. iPSCs have been employed to generate ECs. However, it may be more clinically relevant to transdifferentiate fibroblasts into ECs directly without introducing pluripotent or virally driven transcription factors. Here, we present a protocol that describes the direct conversion of human cardiac fibroblasts into ECs by leveraging the innate immune system. Our protocol produces bona fide human ECs with 95%–98% purity by first passage.

Chun Liu, Pedro Medina, Dilip Thomas, Ian Y. Chen, Karim Sallam, Danish Sayed, Nazish Sayed

liuchun@stanford.edu (C.L.)
sayedns@stanford.edu (N.S.)

Highlights

An optimized protocol for direct reprogramming of human endothelial cells (ECs)

A clinically relevant protocol devoid of the use of exogenous transcription factors

The protocol leverages the innate immune pathway for EC transdifferentiation

Induced human ECs can be generated with a purity of 95%–98% by first passage

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Protocol

A protocol for transdifferentiation of human cardiac fibroblasts into endothelial cells via activation of innate immunity

Chun Liu,^{1,2,5,*} Pedro Medina,¹ Dilip Thomas,^{1,2} Ian Y. Chen,^{1,2} Karim Sallam,^{1,2} Danish Sayed,³ and Nazish Sayed^{1,2,4,6,*}

¹Stanford Cardiovascular Institute, Stanford University, Stanford, CA 94305, USA

²Department of Medicine, Stanford University, Stanford, CA 94305, USA

³Department of Cell Biology and Molecular Medicine, Rutgers New Jersey Medical School, Newark, NJ 07103, USA

⁴Department of Surgery, Division of Vascular Surgery, Stanford University, Stanford, CA 94305, USA

⁵Technical contact

⁶Lead contact

*Correspondence: liuchun@stanford.edu (C.L.), sayedns@stanford.edu (N.S.)
<https://doi.org/10.1016/j.xpro.2021.100556>

SUMMARY

Endothelial cells (ECs) have emerged as key pathogenic players in cardiac disease due to their proximity with cardiomyocytes. Induced pluripotent stem cells (iPSCs) have been employed to generate ECs. However, it may be more clinically relevant to transdifferentiate fibroblasts into ECs directly without introducing pluripotent or virally driven transcription factors. Here, we present a protocol that describes the direct conversion of human cardiac fibroblasts into ECs by leveraging the innate immune system. Our protocol produces bona fide human ECs with 95%–98% purity by first passage.

For complete details on the use and execution of this protocol, please refer to Liu et al. (2020) and Sayed et al. (2015).

BEFORE YOU BEGIN

Human fibroblasts are commercially obtained and used according to ethical guidelines. All experiments are conducted in Class II biosafety cabinets using standard sterile techniques. All cells are cultured and propagated in a humidified 37°C incubator and 5% CO₂. This transdifferentiation protocol can be applied to other types of fibroblasts including skin fibroblasts. Prior to initiation of the protocol, prepare the below media, solutions and matrigel-coated plates. Similarly, the media should be pre-warmed at 37°C for 30 min before the start of the protocol. Please refer to Key Resources Table for a comprehensive list of reagents and resources. Please refer to Materials and Equipment for complete recipe tables.

1. Human Fibroblast Medium: Dulbecco's modified Eagle's medium (DMEM, Gibco, catalog number: 11320033), supplemented with 20% fetal bovine serum (FBS), and 1% penicillin/streptomycin, filter-sterilized.
2. Human iPSC medium: Essential 8™ Medium (Gibco, catalog number: A1517001).
3. Differentiation medium: RPMI 1640 medium, no glutamine (Gibco, catalog number: 21870076).
4. Human Endothelial medium: EGM™ Endothelial Cell Growth Medium BulletKit™ (Lonza, catalog number: CC-3124).
5. Freezing medium: Bambanker (Fisher Scientific, catalog number: NC9582225).



Matrigel-coated culture plates

6. Thaw growth factor-reduced Matrigel on ice. Once thawed, aliquot the Matrigel (250 μ L, each) in 1.5 mL Eppendorf tubes and store at -20°C freezer for future use.
7. Thaw one Matrigel aliquot on ice and resuspend in cold DMEM/F12 at 1:250 dilution.
8. For a 6-well plate, add 2 mL of diluted Matrigel solution per well.
9. Coat the plates for minimum of 30 min in 37°C cell culture incubator and 5% CO_2 before use. Additional plates can be prepared and stored in 37°C cell culture incubator for up to 1 week or 4°C for up to 1 month after sealing it with parafilm to avoid drying of the plates.

Gelatin-coated culture plates

10. Prepare 0.2% gelatin solution in DPBS, filter-sterilized. Store at 4°C for future use.
11. For a 6-well plate, add 2 mL 0.2% gelatin to ensure entire well is covered.
12. Incubate the plate for minimum of 30 min in 37°C cell culture incubator.

KEY RESOURCES TABLE

| REAGENT or RESOURCE | SOURCE | IDENTIFIER |
|--|-------------------|--------------|
| Antibodies | | |
| PE-conjugated anti-human CD31 (1:400) | Miltenyi Biotec | 130-110-669 |
| Anti-human CD31 (1:100) | Thermo Fisher | BBA7 |
| Anti-human CD144 (1:500) | Abcam | ab33168 |
| Alexa Fluor 488 donkey anti-mouse IgG (1:1000) | Thermo Fisher | A21202 |
| Alexa Fluor 594 donkey anti-mouse IgG (1:1000) | Thermo Fisher | A21203 |
| Chemicals, peptides, and recombinant proteins | | |
| DPBS | Gibco | 14190 |
| DMEM/F12 | Gibco | 11320033 |
| Essential 8 medium and supplement | Gibco | A1517001 |
| RPMI 1640 medium | Gibco | 21870076 |
| B27 Supplement, minus insulin | Gibco | A1895601 |
| EGM2 medium | Lonza | CC-3124 |
| Fibroblast Growth Medium 3 | PromoCell | C23025 |
| Growth Medium 3 SupplementMix | PromoCell | C39345 |
| BD Matrigel | Fisher Scientific | 08-774-552 |
| Gelatin | Sigma | G1393 |
| TrypLE Express dissociation reagent | Gibco | 12605010 |
| FBS | Life Technologies | 26140079 |
| Bambanker | Fisher Scientific | NC9582225 |
| GlutaMax | Thermo Fisher | 35050-038 |
| Penicillin/streptomycin | Life Technologies | 15140-122 |
| DMSO | Sigma | D2650 |
| Rock inhibitor Y27632 | Selleckchem | S1049 |
| CHIR 99021 | Selleckchem | S2924 |
| IWR-1 | Selleckchem | S7086 |
| Retinoic acid | Sigma | R2625 |
| KnockOut™ Serum | Thermo Fisher | 10828010 |
| Polyinosine-polycytidylic [Poly(I:C)] | InvivoGen | tlrl-pic |
| VEGF | PeproTech | 100-20-250UG |
| BMP4 | PeproTech | 120-05ET |
| FGF-basic | Fisher Scientific | 233-FB-025 |
| 8-Bromo-cAMP | Fisher Scientific | 11-401-0 |
| SB431542 | Selleck Chemicals | S1067 |

(Continued on next page)

| <i>Continued</i> | | |
|--|-------------------------------|---------------|
| REAGENT or RESOURCE | SOURCE | IDENTIFIER |
| Paraformaldehyde | Fisher Scientific | 50-980-487 |
| Bovine serum albumin (BSA) | Cell Signaling Technology | 9998S |
| Triton X-100 | Sigma | T8787 |
| Normal donkey serum | Sigma | D9663 |
| Permout™ Mounting Medium | Fisher Scientific | SP15-100 |
| Matrigel Basement Membrane Matrix | Fisher Scientific | CB-40234 |
| Acetylcholine | Sigma-Aldrich | A6625-25G |
| <i>Critical commercial assays</i> | | |
| MACS® LS Columns | Miltenyi Biotec | 130-042-401 |
| MACS® BSA Stock Solution | Miltenyi Biotec | 130-091-376 |
| MACS® autoMACS Rinsing Solution | Miltenyi Biotec | 130-091-222 |
| MACS® Strainers 30 µm | Miltenyi Biotec | 130-098-458 |
| MACS® CD144 MicroBeads | Miltenyi Biotec | 130-097-857 |
| Nitrate/Nitrite Fluorometric Assay Kit | Cayman Chemicals | 780051 |
| LDL Uptake Assay Kit (Fluorometric) | BioVision | K585-100 |
| RNeasy Plus Mini Kit | QIAGEN | 74136 |
| qScript® cDNA Synthesis Kit | VWR | 89168-788 |
| qScript cDNA SuperMix | VWR | 101414-106 |
| TaqMan Universal PCR Master Mix | Fisher Scientific | 4364340 |
| <i>Deposited data</i> | | |
| RNA sequencing data sets | Gene Expression Omnibus (GEO) | GSE142217 |
| RNA sequencing data sets | Gene Expression Omnibus (GEO) | GSE45176 |
| <i>Experimental models: Cell lines</i> | | |
| Human cardiac fibroblasts | ScienCell Research Lab | 6330 |
| Healthy control iPSCs | Stanford CVI Biobank | N/A |
| <i>Oligonucleotides</i> | | |
| Real-time qPCR probes – CD31 | Life Technologies | Hs00169777_m1 |
| Real-time qPCR probes – CD144 | Life Technologies | Hs00901465_m1 |
| Real-time qPCR probes – KDR | Life Technologies | Hs00911700_m1 |
| Real-time qPCR probes – vWF | Life Technologies | Hs01109446_m1 |
| Real-time qPCR probes – NOS3 | Life Technologies | Hs01574659_m1 |
| Real-time qPCR probes – GAPDH | Life Technologies | Hs02758991_g1 |
| <i>Other</i> | | |
| Nalgene™ vacuum filter unit | Fisher Scientific | 09-741-04 |
| 6-Well cell culture plates | Fisher Scientific | 08-772-1B |
| 10 cm Cell culture plates | Fisher Scientific | 08-772-E |
| 15 mL Polypropylene conical tubes | Fisher Scientific | 14-959-53A |
| Cryovials | Fisher Scientific | 13-700-500 |
| CoolCell® freezing container | Corning | CLS432002-1EA |
| FACSAria II cell sorter | BD Biosciences | N/A |
| <i>Software and algorithms</i> | | |
| FlowJo 8.7 software | BD Biosciences | N/A |

MATERIALS AND EQUIPMENT

All media and solutions are prepared using sterile techniques.

Matrigel 1:250

Add 200 µL of matrigel thawed on ice into 50 mL cold DMEM/F12 medium to make a stock solution. Add 2 mL diluted matrigel solution to each well of a 6-well cell culture plate.

Gelatin 0.2%

Add 50 mL 2% gelatin into 450 mL DPBS to obtain a 0.2% gelatin stock solution and filter with 0.22 µm filter.

Y27632 10 mM

Dissolve 10 mg of Y27632 in 3.12 mL DMSO to make a 10 mM stock solution. A working concentration of 5 μ M Y27632 would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 6 months.

CHIR99021 10 mM

Dissolve 5 mg of CHIR99021 in 1.08 mL DMSO to make a 10 mM stock solution. A working concentration of 6 μ M CHIR99021 would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 6 months.

IWR-1 10 mM

Dissolve 10 mg of IWR-1 in 2.44 mL DMSO to make a 10 mM stock solution. A working concentration of 5 μ M IWR-1 would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 6 months.

Retinoic acid 100 mM

Dissolve 50 mg of Retinoic acid in 1.66 mL DMSO to make a 100 mM stock solution. A working concentration of 2 μ M Retinoic acid would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 12 months.

Poly(I:C) 1 mg/mL

Dissolve 10 mg of poly(I:C) in 10 mL of sterile endotoxin-free water to make a 1 mg/mL stock solution. Make serial dilutions of poly(I:C) to make a working solution of 1 μ g/mL. A working concentration of 30 ng/mL poly(I:C) would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 12 months.

VEGF 250 μ g/mL

Dissolve 250 μ g of VEGF in 1 mL DPBS and 0.1% MACS BSA solution to make a 250 μ g/mL stock solution. A working concentration of 50 ng/mL VEGF would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 6 months.

BMP4 100 μ g/mL

Dissolve 100 μ g of BMP4 in 1 mL sterile water and 0.1% MACS BSA solution to make a 100 μ g/mL stock solution. A working concentration of 20 ng/mL BMP4 would be needed per well. Make 100 μ l aliquots and store in -80°C for up to 3 months.

FGF-basic 100 μ g/mL

Dissolve 100 μ g of FGF-basic in 1 mL sterile water and 0.1% MACS BSA solution to make a 100 μ g/mL stock solution. A working concentration of 20 ng/mL FGF-basic would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 3 months.

8-Br-cAMP 100 mM

Dissolve 50 mg of 8-Br-cAMP in 1.15 mL sterile water to make a 100 mM stock solution. A working concentration of 100 μ M 8-Br-cAMP would be needed per well. Make 100 μ l aliquots and store in -20°C for up to 6 months.

SB431542 50 mM

Dissolve 10 mg of SB431542 in 520 μ l of DMSO to make a 50 mM stock solution. A working concentration of 10 μ M SB431542 would be needed per well. Make 100 μ l aliquots and store in -80°C for up to 6 months.

Human fibroblast medium

| Reagent | Final concentration | Amount |
|-------------------------|---------------------|---------------|
| DMEM/F12 | n/a | 399 mL |
| FBS | 20% | 100 mL |
| GlutaMax | 1% | 0.5 mL |
| Penicillin/streptomycin | 1% | 0.5 mL |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 1 month

Human iPSC medium

| Reagent | Final concentration | Amount |
|-------------------|---------------------|---------------|
| Essential 8 Media | n/a | 490 mL |
| E8 Supplement | 2% | 10 mL |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 1 month

Human differentiation medium

| Reagent | Final concentration | Amount |
|------------------------------|---------------------|---------------|
| RPMI 1640 medium | n/a | 490 mL |
| B27 supplement minus insulin | 2% | 10 mL |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 1 month

Fibroblast growth medium

| Reagent | Final concentration | Amount |
|----------------------------|---------------------|---------------|
| Fibroblast Growth Medium 3 | n/a | 495 mL |
| SupplementMix | n/a | 5 mL |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 6 weeks

Human endothelial (EGM-2) medium

| Reagent | Final concentration | Amount |
|--|---------------------|---------------|
| Endothelial Cell Growth Basal Medium 2 | n/a | 500 mL |
| Epidermal Growth Factor | 5 ng/mL | Kit aliquot |
| bFGF | 10 ng/mL | Kit aliquot |
| Insulin-like Growth Factor (R3 IGF-1) | 20 ng/mL | Kit aliquot |
| VEGF 165 | 0.5 ng/mL | Kit aliquot |
| Ascorbic Acid | 1 µg/mL | Kit aliquot |
| Heparin | 22.5 µg/mL | Kit aliquot |
| Hydrocortisone | 0.2 µg/mL | Kit aliquot |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 1 month

Induction medium

| Reagent | Final concentration | Amount |
|----------------|---------------------|---------------|
| DMEM/F12 | n/a | 425 mL |
| FBS | 7.5% | 37.5 mL |
| Knockout serum | 7.5% | 37.5 mL |
| Poly(I:C) | 30 ng/mL | 15 µL |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 2 weeks.

Transdifferentiation medium

| Reagent | Final concentration | Amount |
|--------------|---------------------|---------------|
| EGM2 medium | n/a | 500 mL |
| VEGF | 50 ng/mL | 100 μ L |
| bFGF | 20 ng/mL | 100 μ L |
| BMP4 | 20 ng/mL | 100 μ L |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 2 weeks.

Maintenance medium

| Reagent | Final concentration | Amount |
|--------------|---------------------|---------------|
| EGM2 medium | n/a | 500 mL |
| VEGF | 50 ng/mL | 100 μ L |
| bFGF | 20 ng/mL | 100 μ L |
| 8-Br-cAMP | 100 μ M | 500 μ L |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 2 weeks.

Expansion medium

| Reagent | Final concentration | Amount |
|--------------|---------------------|---------------|
| EGM2 medium | n/a | 500 mL |
| SB431542 | 10 μ M | 100 μ L |
| Total | n/a | 500 mL |

Once prepared, store at 4°C for up to 2 weeks.

Cryopreservation medium

| Reagent | Final concentration | Amount |
|-----------|---------------------|-----------|
| BamBanker | n/a | 1 mL/vial |

Store at 4°C for up to 12 months.

MACS buffer

| Reagent | Final concentration | Amount |
|---------------------------------|---------------------|---------------|
| MACS® AutoMACS Rinsing Solution | 95% | 95 mL |
| MACS® BSA Stock Solution | 5% | 5 mL |
| Total | n/a | 100 mL |

Once prepared, use the same day.

Immunofluorescence blocking buffer

| Reagent | Final concentration | Amount |
|---------------------|---------------------|---------------|
| DPBS | 88.75% | 88.75 mL |
| BSA | 1% | 1 mL |
| Normal Donkey Serum | 10% | 10 mL |
| Triton X | 0.25% | 250 μ L |
| Total | n/a | 100 mL |

Once prepared, aliquot and store at -20°C for up to 2 months.



Immunofluorescence antibody buffer

| Reagent | Final concentration | Amount |
|---------------------|---------------------|--------------|
| DPBS | 97.75% | 9.775 mL |
| BSA | 1% | 0.1 mL |
| Normal Donkey Serum | 1% | 0.1 mL |
| Triton X | 0.25% | 25 μ L |
| Total | n/a | 10 mL |

Once prepared, aliquot and store at -20°C for up to 2 months.

STEP-BY-STEP METHOD DETAILS

Human fibroblasts

Cardiac fibroblasts are an essential cell type of the human heart that not only provide structural support to the cardiomyocytes but are also an important source of paracrine growth factors (Tallquist and Molkentin, 2017). Under pathophysiological conditions, FBs become proliferative and synthesize extracellular matrix that result in scar formation and fibrosis (Fu et al., 2018; Sayed et al., 2013). Human FBs can be obtained from commercial vendors (who isolate FBs from human heart tissue) or can be obtained by differentiating human iPSCs (Figure 1). The acquisition of human heart tissue is rare and expensive, and thus iPSC-derived FBs pose as an ideal alternative. In this protocol we obtained human FBs from commercial vendor, ScienCell Research Laboratories. They were passaged and maintained as described below. We obtained human iPSC-derived cardiac fibroblasts (iPSC-CFs) from the laboratory of Dr. Joseph Wu following their established protocol (Zhang et al., 2020). Healthy control iPSCs were obtained from Stanford CVI Biobank (<https://med.stanford.edu/scvibiobank.html>) using an Institutional Review Board (IRB)-approved protocol. A brief summary is provided below.

Maintenance and passaging of human cardiac fibroblasts

⌚ Timing: 1 week

1. Prepare a 6-well matrigel coated plate for the recovery of human cardiac fibroblasts. Incubate the plate for 30 min in a 37°C cell culture incubator.
2. Thaw and quickly recover one cryovial of commercial human fibroblasts in 1-well of the matrigel coated 6-well plate using 2 mL fibroblast medium.
3. Following 24 h incubation, replace the medium with fresh 2 mL fibroblast medium.
4. Change medium every 2 days till confluent.
5. Once confluent, passage the human fibroblasts 1:6 split by incubating with 1 mL TrypLE per well for 5 min in a 37°C cell culture incubator and plate onto a new 6-well matrigel coated plate to achieve 70%–80% confluency.

⚠ **CRITICAL:** We recommend a confluency of 70%–80% before starting the transdifferentiation protocol and a confluency of 100% before cryopreserving them.

⏸ **Pause point:** At this stage of the protocol, the cells can be cryopreserved for future use by resuspending them in Bambanker (freezing medium) and placing in CoolCell® at -80°C for 24 h followed by long-term storage in liquid nitrogen. Commercial human cardiac fibroblasts are usually received from the vendor at passage one and can be expanded for 15 population doublings as per the vendor's recommendations.

Alternate method to obtain human cardiac fibroblasts: Differentiation of human iPSCs into human cardiac fibroblasts

⌚ Timing: 3 weeks

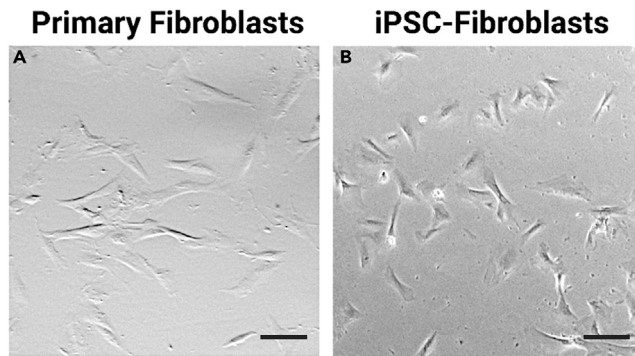


Figure 1. Obtaining human fibroblasts

(A) Representative brightfield image of commercially obtained primary human cardiac fibroblasts.

(B) Representative brightfield image of iPSC-derived human cardiac fibroblasts. Image is of day 21 iPSC-CFs since start of differentiation. Scale bars, 50 μm .

Human induced pluripotent stem cells (iPSCs) have served as an ideal platform to generate patient-specific cardiovascular cells to model human diseases (Sayed et al., 2016; Sayed and Wu, 2017) or assess cardiotoxicity (Sayed et al., 2019; Stack et al., 2019; Thomas et al., 2021). In this protocol, we used healthy control iPSCs cultured in human iPSC medium (E8 medium) to generate cardiac fibroblasts. We include an overview of the protocol:

6. Seed iPSCs in matrigel coated 6-well plate till 80%–85% confluency.
7. Once confluent, initiate the differentiation protocol by replacing the E8 medium with 2 mL RPMI/B27 minus insulin (RPMI/B27-insulin) per well supplemented with 6 μM CHIR99021 for 2 days.
8. On day 2, replace the medium with fresh 2 mL RPMI/B27-insulin per well for 24 h. After the initiation of the protocol, cells are not washed with 1 \times PBS.
9. On day 3, replace the medium with freshly prepared 2 mL RPMI/B27-insulin per well supplemented with 5 μM IWR-1 for 2 days.
10. On day 5, replace the medium with fresh 2 mL RPMI/B27-insulin per well for 24 h.
11. On day 6, passage the differentiating cells with 1 mL TrypLE per well and seed them in new matrigel coated 6-well plates (at 1:12 split ratio) in freshly prepared 2 mL Advanced DMEM/GlutaMAX medium per well with 1% FBS supplemented with 5 μM CHIR99021, 2 μM retinoic acid, and 5 μM of Y27632.
12. On day 7, replace the medium with freshly prepared 2 mL Advanced DMEM/GlutaMAX medium per well supplemented with 5 μM CHIR99021 and 2 μM retinoic acid for 2 days.
13. On day 9, replace the medium with fresh 2 mL Advanced DMEM/GlutaMAX medium per well for 2 days.
14. On day 11, passage the cells with 1 mL TrypLE per well and seed them in new matrigel coated 6-well plates (at 1:3 or 1:6 split ratio) in freshly prepared 2 mL Advanced DMEM/GlutaMAX medium per well supplemented with 2 μM SB431542 for 24 h.
15. On day 12, replace the medium with fresh 2 mL Advanced DMEM/GlutaMAX medium per well supplemented with 2 μM SB431542, and change medium till the cells are confluent.
16. On day 14, passage the cells with 1 mL TrypLE per well and seed them in new matrigel coated 10 cm cell culture dish (at 1:3 split ratio or 2 full 6-well plates to one 10 cm dish) in freshly prepared 10 mL fibroblast growth medium supplemented with 20 ng/mL bFGF and 10 μM SB431542.
17. On day 16, replace the medium with fresh 10 mL fibroblast growth medium per dish supplemented with 20 ng/mL bFGF and 10 μM SB431542.
18. On day 18, replace the medium with fresh 10 mL fibroblast growth medium per dish supplemented with 20 ng/mL bFGF and 10 μM SB431542.

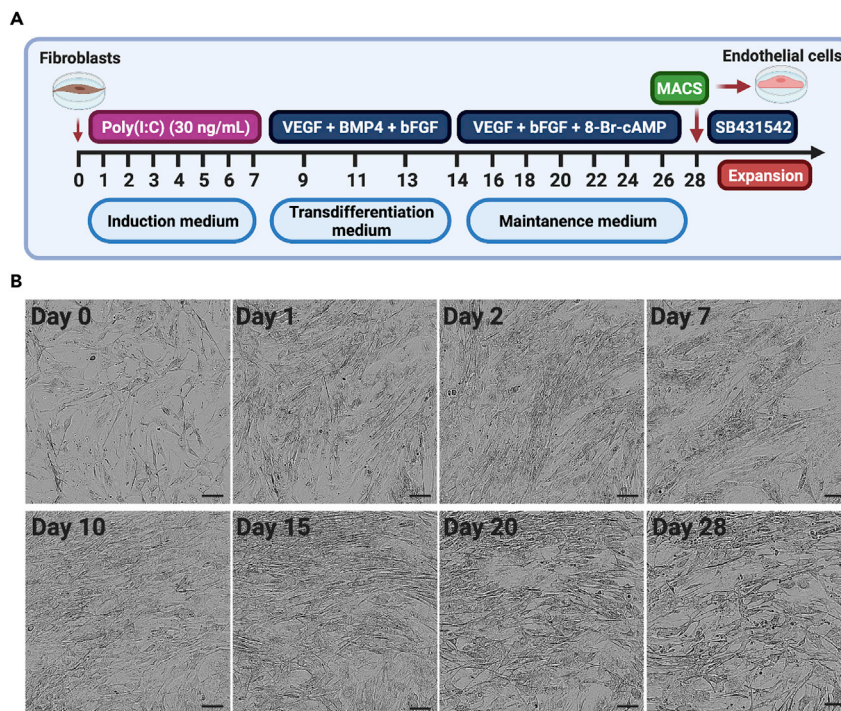


Figure 2. Protocol to transdifferentiate human fibroblasts into human endothelial cells

(A) Schematic of protocol to transdifferentiate human fibroblasts into endothelial cells. The schematic outlines the time course and treatments of growth factors and small molecules for the transdifferentiation of fibroblasts. (B) Representative phase-contrast images of human fibroblasts undergoing transdifferentiation at different time-points (days) using Incucyte® Live-Cell Imaging system. Scale bars, 100 μ m.

19. By day 20, the resulting cardiac fibroblasts are confluent and can be passaged for transdifferentiation.

△ CRITICAL: By day 18, the purity of cardiac fibroblasts differentiated from iPSCs should be >95% (Zhang et al., 2019a). However, we recommend checking the purity of iPSC-CFs before initiating their transdifferentiation to ECs by assessing the expression of CF markers such as COL-1, DDR2, and TCF21 via fluorescence-activated cell sorting (FACS).

▣ Pause point: At this stage the iPSC-CFs can be cryopreserved for future use by resuspending them in Bambanker (freezing medium) and placing in CoolCell® at -80°C for 24 h followed by long-term storage in liquid nitrogen.

Transdifferentiating human fibroblasts to endothelial cells

⌚ Timing: 4 weeks

Activation of innate immune pathways have been shown to play an important role in nuclear reprogramming (Lee et al., 2012; Liu et al., 2021; Liu et al., 2017; Sayed et al., 2017) and transdifferentiation of one cell type to another (Sayed et al., 2015; Zhou et al., 2019). The protocol described below employs a safe and efficient way to directly differentiate human fibroblasts to endothelial cells without the use of any exogenous transcription factors (Sayed et al., 2015) (Figures 2A and 2B). Transdifferentiated ECs (iECs) attain an efficiency of 95%–98% by passage 1 and can be cryopreserved for future research purposes.

Note: Human fibroblasts obtained either commercially or derived using human iPSCs should be cultured at least 3–4 days before initiating the transdifferentiating protocol. The protocol was optimized using a 6-well plate.

20. On day 0 (the day before transdifferentiation), prepare 6-well gelatin-coated plates 30 min prior to seeding the human fibroblasts.
21. Following 30 mins, remove the excess gelatin and seed human fibroblasts in human fibroblast medium at a density of 1×10^5 cells per well of a 6-well plate for 24 h.
22. On day 1, aspirate the spent medium and replace with induction medium (see recipe) containing DMEM/F12 supplemented with 7.5% FBS, 7.5% knockout serum (KSR), and 30 ng/mL poly(I:C).
23. Treat cells daily with fresh induction medium (day 2–6).
24. On day 7, switch the induced human fibroblasts over to transdifferentiation medium (see recipe) containing EGM2 medium supplemented with 50 ng/mL vascular endothelial growth factor (VEGF), 20 ng/mL bone morphogenetic protein 4 (BMP4), and 20 ng/mL basic fibroblast growth factor (bFGF) for 2 days.
25. On day 9, replace the spent medium with fresh transdifferentiation medium for 2 days.
26. On day 11, replace the spent medium with fresh transdifferentiation medium for 2 days.
27. On day 13, replace the spent medium with fresh transdifferentiation medium for 1 day.
28. On day 14, switch the transdifferentiated human fibroblasts over to maintenance medium (see recipe) containing EGM2 medium supplemented with 50 ng/mL VEGF, 20 ng/mL bFGF, and 100 μ M 8-bromo-adenosine-3':5'-cyclic monophosphate sodium salt (8-Br-cAMP) for the next 2 weeks, changing medium every 2 days.
29. On day 28, the cells are ready for CD144 MACS sorting.

⚠ **CRITICAL:** We recommend observing the cells throughout the protocol as distinct morphological changes can be seen, especially after the addition of the transdifferentiation medium (Figure 2).

CD144 magnetic-activated cell sorting (MACS)

⌚ **Timing:** 3 h

This step sorts the iECs from the transdifferentiation protocol. CD144, also known as VE-Cadherin has been widely used for the isolation of ECs derived from embryonic stem cells (ESCs) and iPSCs (Liu et al., 2020; Sayed et al., 2020; Sayed et al., 2015).

30. Aspirate the spent medium and dissociate the cells using 1 mL TrypLE express per well for 10 mins at 37°C.
31. Pipette the cells repeatedly to ensure single cell suspension and collect the cells in 1 mL basal EGM2 medium to neutralize the TrypLE express.
32. Strain the cell suspension through a pre-rinsed 40 μ m strainer into a 50 mL conical tube, centrifuge at 300 \times g for 5 min, and resuspend 1×10^7 cells in 160 μ L ice-cold MACS buffer.

⚠ **CRITICAL:** Pre-rinse the strainer with MACS buffer before use. Typically, one strainer is used for every 6-well plate, however, replace the strainer with a new one if any clogging is observed.

33. Incubate the cell suspension with 40 μ L of CD144 magnetic microbeads at 4°C for 15 min in dark.
34. Wash the cells with 1 mL MACS buffer and centrifuge at 300 \times g for 5 min.
35. During the centrifugation step, setup the LS columns on the magnetic MACS sorting apparatus and pre-rinse the columns with 3 mL MACS buffer.

⚠ **CRITICAL:** Ensure the cells remain on ice before and after sorting to avoid cell death.

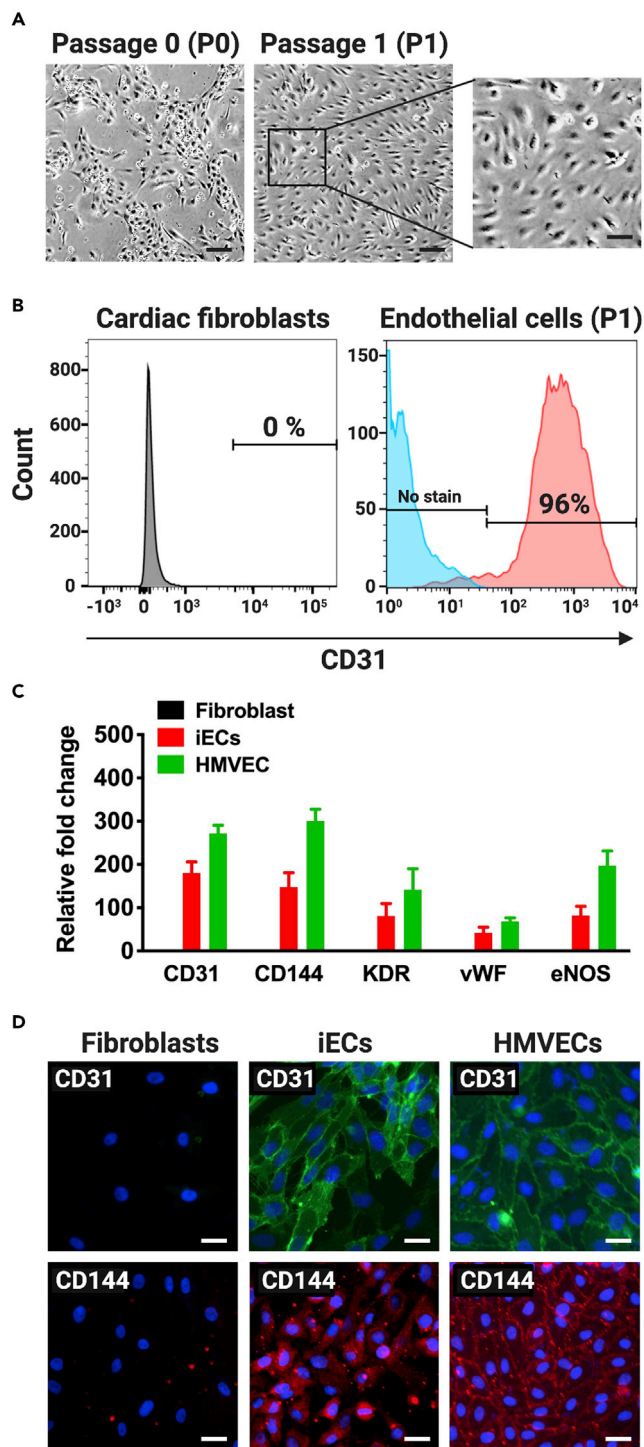


Figure 3. Isolation and validation of induced endothelial cells (iECs)

(A) Representative brightfield images of iECs transdifferentiated from human fibroblasts at passage 0 (P0, left panel) and passage 1 (P1, right panel) show typical cobblestone appearance. Scale bars, 100 μ m. Inset shows high magnification of P1 iECs. Scale bars, 50 μ m.

(B) FACS analysis of cardiac fibroblasts (left panel) and iECs (right panel) with CD31 antibody shows ~96% purity at P1.

Figure 3. Continued

(C) Quantitative PCR (qPCR) data of transdifferentiated iECs show expression of EC markers, CD31, CD144, KDR, vWF, and eNOS as relative fold change when compared to human fibroblasts and normalized to GAPDH, a housekeeping gene. Human microvascular endothelial cells (HMVEC) were used as positive control.

(D) Representative immunofluorescent images of transdifferentiated iECs show expression of CD31 and CD144 protein levels compared to human fibroblasts. Scale bars, 25 μ m.

36. After centrifugation, remove supernatant and resuspend the cell pellet with 1 mL MACS buffer.
37. Pass the cell suspension through the LS column.
38. Wash the column with 3 mL MACS buffer, repeating it 3 times to remove unlabeled cells.
39. After washes, remove the column from the magnetic separator and place on a 15 mL conical tube.
40. Add 5 mL MACS buffer to the column and firmly push the column plunger to elute CD144+ labeled cells.

▣ **Pause point:** CD144+ iECs can be cryopreserved using Bambanker in liquid nitrogen at this stage.

△ **CRITICAL:** Follow the steps listed in the Miltenyi Biotech MACS sorting protocol (<https://www.miltenyibiotec.com/US-en/products/macs-cell-separation.html>). Prewash the LS columns with MACS buffer to ensure removal of any bubbles. When washing the columns, wait till all the previous wash volume has passed through the column before adding the next volume. When plunging the cells from the column ensure steady and even pressure is applied on the column plunger.

Re-plating iECs

⌚ **Timing:** 30 min

△ **CRITICAL:** As the yield of transdifferentiated iECs at passage 0 (P0) is $\sim 2.5 \times 10^5$ cells for each 6-well plate of human fibroblasts, seed the iECs in 1-well of a 6-well plate.

41. Centrifuge the CD144+ labeled cells at $300 \times g$ for 5 min. Resuspend the cell pellet in expansion medium (see recipe) containing EGM2 and 10 μ M SB431542 and seed the cells in 1-well of a 6-well plate precoated with 0.2% gelatin.
42. Ensure cells are evenly dispersed and place them in 37°C incubator.

Maintaining and passaging iECs

⌚ **Timing:** up to passage 8

Transdifferentiated iECs can be maintained with medium changes every 2 days and expanded up to 8 passages (Figure 3A).

43. Once 100% confluent, dissociate the P0 iECs with 1 mL TrypLE for 5 min at 37°C. Neutralize the cell suspension with basal EGM2 medium, centrifuge at $300 \times g$ for 5 min, and resuspend the cell pellet with expansion medium.

△ **CRITICAL:** Passage P0 iECs on gelatin-coated plates at a 1:3 split ratio.

Cryopreserving iECs

⌚ **Timing:** 30 mins

▮▮ **Pause point:** Passage 1 (P1) iECs can be cryopreserved for future use.

44. Aspirate the spent medium from cells and dissociate them with 1 mL TrypLE for 5 min at 37°C. Neutralize the cells with media and centrifuge at 300 × g for 5 min.
45. Aspirate the supernatant and resuspend the cells gently with 1 mL Bambanker.
46. Transfer the cell suspension into cryogenic storage vials and place them in CoolCell® freezing container to be placed in –80°C for 24 h.
47. Transfer the cryogenic vials to liquid nitrogen for long-term storage.

Validating iECs: Fluorescence-activated cell sorting (FACS)

⌚ **Timing:** 2 h

Once the iECs have been expanded they can be characterized and assessed for their purity using FACS (Figure 3B). CD31, also known as platelet endothelial cell adhesion molecule (PECAM-1), is highly expressed at the intercellular junctions of ECs and plays an important role in maintaining vascular permeability. It is a ubiquitous surface marker of ECs.

48. Dissociate iECs from gelatin-coated plates with 1 mL TrypLE for 5 min at 37°C. Neutralize TrypLE with media and centrifuge at 300 × g for 5 min.
49. Aspirate the supernatant and resuspend the cell pellet gently in 1 × PBS containing 2% FBS at 5 × 10⁵ cells per 100 μL volume per tube.
50. Incubate the cells with PE-conjugated antibody against CD31 at 1:400 dilution for 30 min on ice.
51. Wash cells to remove excess antibody, centrifuge and resuspend in 1 × PBS containing 2% FBS.
52. Run FACS.

Note: A FACSAria II cell sorter was used for sorting and data was analyzed using FlowJo 8.7 software (TreeStar).

Validating iECs: Characterizing gene expression

⌚ **Timing:** 6 h

The gene expression profile of iECs using quantitative PCR can be used to confirm the identity of the transdifferentiated cells (Figure 3C). CD31 has been primarily used to demonstrate the presence of ECs. Similarly, CD144, also known as vascular endothelial cadherin (VE-cadherin) is a marker for ECs (Liu et al., 2020; Sayed et al., 2015).

53. RNA extraction
 - a. Dissociate iECs from 1-well of a 6-well plate using 1 mL TrypLE for 5 min at 37°C. Neutralize the cells with basal EGM2 media and centrifuge at 300 × g for 5 min.
 - b. Extract RNA using an RNA extraction kit such as Qiagen RNeasy Plus Mini Kit.
 - c. Resuspend the pellet with lysis buffer (buffer RLT) to disrupt them by pipetting up and down.
 - d. Add equal volume of 70% ethanol to the homogenized cell lysate.
 - e. Transfer the cell lysate to RNeasy spin column® to capture the RNA, followed by washes with RW1 and RPE buffers.
 - f. Elute RNA with RNase-free water.
 - g. Determine the RNA concentration using a Nanodrop.
54. cDNA preparation
 - a. Use qScript® cDNA Synthesis Kit for cDNA synthesis.
 - b. Thaw the RNA and qScript® cDNA SuperMix on ice and prepare 0.2 mL thin-walled PCR tube for the PCR reaction.

| Reagent | 1 × (volume) |
|-------------------------------|--------------|
| RNA (1 µg to 10 pg total RNA) | variable |
| Nuclease-free water | variable |
| qScript Reaction Mix (5×) | 4.0 µL |
| qScript RT | 1.0 µL |
| Final volume | 20.0 µL |

- c. Once mixed, vortex gently and centrifuge to pool the reaction volume.
- d. Place the tubes in a thermal cycler and follow the recommended PCR program.

| Number of cycles | Temperature | Time |
|------------------|-------------|--------|
| 1 cycle | 22°C | 5 min |
| 1 cycle | 42°C | 30 min |
| 1 cycle | 85°C | 5 min |
| Hold | 4°C | 5 min |

55. Quantitative PCR

- a. Dilute the cDNA samples (1:5 ratio) with RNase/DNase-free water.
- b. Thaw the gene-specific primers (CD31, CD144, KDR, vWF, and NOS3) on ice and prepare the reaction mix.

| Reagent | 1 × (volume) |
|---------------------------------|--------------|
| cDNA sample | 3 µL |
| TaqMan Universal PCR Master Mix | 10 µL |
| Gene-specific primer | 1 µL |
| Water | 6 µL |
| Final volume | 20.0 µL |

- c. Turn on the ABI Real time PCR machine (ABI 7300) 30 mins prior to running the qPCR.
- d. Follow the recommended qPCR program.

| PCR cycling conditions | | | |
|------------------------|-------------|---------|--------|
| Steps | Temperature | Time | Cycles |
| Initial denaturation | 94°C | 2 min | 1 |
| Denaturation | 94°C | 30 s | 40 |
| Annealing | 58°C | 30 s | |
| Extension | 68°C | 1 min | |
| Final extension | N/A | N/A | |
| Hold | 4°C | Forever | |

56. Analyze the expression of EC genes by normalizing the data to a housekeeping gene (GAPDH) and express the data as relative fold changes with the Δ Ct method of analysis.

Validating iECs: Characterizing protein expression

⌚ Timing: 2 days

In addition to gene expression profiling, the protein expression of endothelial markers such as CD31 and CD144 can also be used to characterize the transdifferentiated iECs (Figure 3D).

57. Aspirate the media from 1-well of a 6-well plate and wash cells with DPBS.
58. Fix cells with 4% paraformaldehyde (PFA) for 1 h at RT.
59. Wash cells 3 times with DPBS.
60. Block cells with blocking buffer containing 10% normal donkey serum and 1% BSA for 1 h at RT.
61. Apply CD31 (1:100) and CD144 (1:500) primary antibodies diluted in antibody buffer containing 1% normal donkey serum and 1% BSA for 16 h at 4°C.
62. Wash cells 3 times with DPBS.
63. Apply secondary antibodies (1:1000) diluted in antibody buffer containing 1% normal donkey serum and 1% BSA for 1 h at RT. Add DAPI at 5 µg/mL during this incubation.
64. Wash cells 5 times with DPBS.
65. Capture images using a fluorescent microscope.
66. Cells can be stored in PBS at 4°C for future imaging. Seal the edges of the plate with parafilm to prevent the cells from drying out.

EXPECTED OUTCOMES

Induced endothelial cells (iECs) transdifferentiated from human fibroblasts can be available for downstream phenotypic characterization within 4 weeks. Importantly, these transdifferentiated iECs can be generated from different fibroblasts including cardiac, skin, and iPSC-derived fibroblasts and are devoid of virally-driven transcription factors. To maximize the output of iECs, we recommend starting with three 6-well plates of human fibroblasts. In this scenario, the expected outcome would be collection of at least two 6-well plates of iECs by passage 2. We recommend dedicating 2-wells for each downstream phenotypic characterization. The most widely used assays to assess the function of ECs are listed here: a) *In vitro* angiogenesis assay to assess the capacity to form network of tubular structures; b) Nitric oxide assay to assess the capacity to produce NO; c) Acetylated LDL-uptake assay to assess the capacity to uptake Ac-LDL; d) *In vitro* scratch assay to assess the migration capacity; e) *In vivo* angiogenesis assay to assess the capacity to form capillaries (Huang et al., 2021; Liu et al., 2018; Paik et al., 2018; Sayed et al., 2020; Zhang et al., 2019b).

LIMITATIONS

Despite being a simple, safe protocol, which is devoid of the use of exogenous transcription factors, to directly differentiate human fibroblasts to endothelial cells, there are a few limitations and concerns. First, the efficiency of generating iECs using this protocol is low. Every 6-well plate of human fibroblasts that are transdifferentiated yields about 4%–5% of iECs. Second, the differentiation efficiency could vary between which human fibroblasts are being used for transdifferentiation. Lastly, the isolation of iECs from the transdifferentiated cell suspension is heavily dependent on endothelial cell surface markers.

TROUBLESHOOTING

Problem 1

Poor proliferative capacity of human fibroblasts before initiating the transdifferentiation protocol (step 5).

Potential solution

Culturing human fibroblasts long-term in tissue culture plates might lead to cellular senescence, which can render the cells non-proliferative. We recommend seeding the human fibroblasts at a much higher seeding density so that the transdifferentiation protocol can be initiated within a day.

Problem 2

Extensive iPSC detachment during iPSC-CF differentiation (step 7).

Potential solution

For most iPSC lines, 6 μ M CHIR99021 for 2 days is sufficient to induce differentiation to cardiac progenitor cells. However, if massive cell death is observed after addition of CHIR99021, the concentration and time of exposure for CHIR99021 should be adjusted to ensure successful differentiation of iPSC-CFs.

Problem 3

Low yield of iPSC-CFs after differentiation (step 19).

Potential solution

The protocol to generate iPSC-CFs involves three stages, which include: i) differentiation of iPSCs into cardiac progenitors, ii) differentiation of cardiac progenitors into proepicardial cells, and iii) differentiation of iPSC-derived epicardial cells into cardiac fibroblasts. Passing through these stages, it is possible to obtain a low yield of iPSC-CFs. This could be due to low quality of iPSC-derived epicardial cells. We recommend to carefully review the morphology of iPSC-derived epicardial cells before differentiating them to iPSC-CFs. In case, the cells fail to exhibit the typical cobblestone-like morphology, we recommend discarding the cells and stop the differentiation process to iPSC-CFs.

Problem 4

Low yield of transdifferentiated iECs after MACS sorting (step 40).

Potential solution

There are a few steps that need to be followed to avoid a very low yield of iECs. 1) Before MACS sorting, an important step is to ensure complete and total dissociation of the cell suspension. For this, we recommend adding TrypLE for 10 min at 37°C and vigorously pipetting the cell suspension for 5 mins. 2) During MACS sorting, avoid clogging of the LS columns by ensuring the cells remain in single cell suspension throughout the MACS protocol. 3) Additional washes of the LS columns can also increase the yield on iECs.

Problem 5

Low maturity of transdifferentiated iECs (step 41).

Potential solution

Primary human endothelial cells are significantly more mature when compared to transdifferentiated iECs, especially with respect to their gene and protein expression (Sayed et al., 2015). Even though, it is unlikely that transdifferentiated iECs could fully mimic primary ECs, mainly due to differences in their environmental milieu, we recommend two steps that can improve some aspects of iEC maturity. These include: i) culturing iECs on specific substrates such as fibronectin, and ii) passaging iECs to passage 2 before subjecting them to *in vitro* characterization.

RESOURCE AVAILABILITY

Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Nazish Sayed (sayedns@stanford.edu).

Materials availability

This study did not generate new unique reagents.

Data and code availability

The datasets (RNA sequencing) generated during this study are available at Gene Expression Omnibus (GEO) under accession numbers GEO: GSE142217 and GSE45176.

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AUTHOR CONTRIBUTIONS

C.L., P.M., D.T., and I.Y.C. performed the experiments and analyzed the data. C.L., K.S., D.S., and N.S. wrote the manuscript.

DECLARATION OF INTERESTS

Dr. Sayed is an inventor of the intellectual property, assigned to Stanford University, related to the use of innate immune signaling for nuclear reprogramming and transdifferentiation.

REFERENCES

- Fu, X., Khalil, H., Kanisicak, O., Boyer, J.G., Vagnozzi, R.J., Maliken, B.D., Sargent, M.A., Prasad, V., Valiente-Alandi, I., Blaxall, B.C., et al. (2018). Specialized fibroblast differentiated states underlie scar formation in the infarcted mouse heart. *J. Clin. Invest.* *128*, 2127–2143.
- Huang, X., Shen, W., Veizades, S., Liang, G., Sayed, N., and Nguyen, P.K. (2021). Single-cell transcriptional profiling reveals sex and age diversity of gene expression in mouse endothelial cells. *Front. Genet.* *12*, 590377.
- Lee, J., Sayed, N., Hunter, A., Au, K.F., Wong, W.H., Mocarski, E.S., Pera, R.R., Yakubov, E., and Cooke, J.P. (2012). Activation of innate immunity is required for efficient nuclear reprogramming. *Cell* *151*, 547–558.
- Liu, C., Ameen, M., Himmati, S., Thomas, D., and Sayed, N. (2021). Generation of human iPSCs by protein reprogramming and stimulation of TLR3 signaling nuclear reprogramming. *Methods Mol. Biol.* *2239*, 153–162.
- Liu, C., Cheng, L., Chen, C.L., and Sayed, N. (2018). Generation of endothelial cells from human induced pluripotent stem cells. *Bio-Protocol* *8*, e3086.
- Liu, C., Himmati, F., and Sayed, N. (2017). Paying the toll in nuclear reprogramming. *Front. Cell Dev. Biol.* *5*, 70.
- Liu, C., Ruan, H., Himmati, F., Zhao, M.-T., Chen, C.C., Makar, M., Chen, I.Y., Sallam, K., Mocarski, E.S., Sayed, D., et al. (2020). HIF1 α regulates early metabolic changes due to activation of innate immunity in nuclear reprogramming. *Stem Cell Rep.* *14*, 192–200.
- Paik, D.T., Tian, L., Lee, J., Sayed, N., Chen, I.Y., Rhee, S., Rhee, J.W., Kim, Y., Wirka, R.C., Buikema, J.W., et al. (2018). Large-scale single-cell RNA-seq reveals molecular signatures of heterogeneous populations of human induced pluripotent stem cell-derived endothelial cells. *Circ. Res.* *123*, 443–450.
- Sayed, N., Ameen, M., and Wu, J.C. (2019). Personalized medicine in cardio-oncology: the role of induced pluripotent stem cell. *Cardiovasc. Res.* *115*, 949–959.
- Sayed, N., Liu, C., Ameen, M., Himmati, F., Zhang, J.Z., Khanamiri, S., Moonen, J.-R., Wnorowski, A., Cheng, L., Rhee, J.-W., et al. (2020). Clinical trial in a dish using iPSCs shows lovastatin improves endothelial dysfunction and cellular cross-talk in LMNA cardiomyopathy. *Sci. Transl. Med.* *12*, eaax9276.
- Sayed, N., Liu, C., and Wu, J.C. (2016). Translation of human-induced pluripotent stem cells: from clinical trial in a dish to precision medicine. *J. Am. Coll. Cardiol.* *67*, 2161–2176.
- Sayed, N., Ospino, F., Himmati, F., Lee, J., Chanda, P., Mocarski, E.S., and Cooke, J.P. (2017). Retinoic acid inducible gene 1 protein (rig1)-like receptor pathway is required for efficient nuclear reprogramming. *Stem Cells* *35*, 1197–1207.
- Sayed, N., Wong, W.T., and Cooke, J.P. (2013). Therapeutic transdifferentiation: can we generate cardiac tissue rather than scar after myocardial injury? *Methodist Debakey Cardiovasc. J.* *9*, 210–212.
- Sayed, N., Wong, W.T., Ospino, F., Meng, S., Lee, J., Jha, A., Dexheimer, P., Aronow, B.J., and Cooke, J.P. (2015). Transdifferentiation of human fibroblasts to endothelial cells: role of innate immunity. *Circulation* *131*, 300–309.
- Sayed, N., and Wu, J.C. (2017). Towards cardio-precision medicine. *Eur. Heart J.* *38*, 1014–1016.
- Stack, J.P., Moslehi, J., Sayed, N., and Wu, J.C. (2019). Cancer therapy-induced cardiomyopathy: can human induced pluripotent stem cell modelling help prevent it? *Eur. Heart J.* *40*, 1764–1770.
- Tallquist, M.D., and Molkenin, J.D. (2017). Redefining the identity of cardiac fibroblasts. *Nat. Rev. Cardiol.* *14*, 484–491.
- Thomas, D., Shenoy, S., and Sayed, N. (2021). Building multi-dimensional induced pluripotent stem cells-based model platforms to assess cardiotoxicity in cancer therapies. *Front. Pharmacol.* *12*, 607364.
- Zhang, H., Shen, M., and Wu, J.C. (2020). Generation of quiescent cardiac fibroblasts derived from human induced pluripotent stem cells. *Methods Mol. Biol.* *1–7*.
- Zhang, H., Tian, L., Shen, M., Tu, C., Wu, H., Gu, M., Paik, D.T., and Wu, J.C. (2019a). Generation of quiescent cardiac fibroblasts from human induced pluripotent stem cells for in vitro modeling of cardiac fibrosis. *Circ. Res.* *125*, 552–566.
- Zhang, J.Z., Termglinchan, V., Shao, N.Y., Itzhaki, I., Liu, C., Ma, N., Tian, L., Wang, V.Y., Chang, A.C.Y., Guo, H., et al. (2019b). A human ipsc double-reporter system enables purification of cardiac lineage subpopulations with distinct function and drug response profiles. *Cell Stem Cell* *24*, 802–811.e805.
- Zhou, Y., Liu, Z., Welch, J.D., Gao, X., Wang, L., Garbutt, T., Keepers, B., Ma, H., Prins, J.F., Shen, W., et al. (2019). Single-cell transcriptomic analyses of cell fate transitions during human cardiac reprogramming. *Cell Stem Cell* *25*, 149–164.e149.