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Epithelial Cell Differentiation from Human Induced Pluripotent Stem Cells
Using a Single-Cell Culture Method

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Abstract: The conventional culture method of human induced pluripotent stem cells (hiPSCs) has been performed in colony cultures using feeder cells, such as mouse embryonic fibroblasts, which require setup times and procedural complexity, a potential risk of transmission of animal pathogens. Besides, the colony culture exhibits slow growth rate and often give rise to heterogeneous cellular states. However, developing technical methodologies of hiPSCs remains pivotal for use in medical applications and research. Here, we investigated whether hiPSCs passaged and expanded as single cells under feeder-free conditions could differentiate into ectodermal-epithelial cells as a source of cells for future regenerative medicine research. First, an hiPSC line 253G1 was cultured in colonies maintained on feeders and subsequently transferred to a single cell on feeder-free. hiPSCs were then cultured as single cells for 28 days in an induction medium supplemented with retinoic acid, bone morphogenetic protein 4, and N2 supplement for epithelial cell differentiation. The expression of epithelial markers, tumor protein p63 (P63), cytokeratin (CK) 18, and CK14 in induced cells was evaluated over time using real-time polymerase chain reaction, western blotting, and immunocytochemistry. Results showed that hiPSCs cultured as single cells expressed pluripotency markers, as evidenced by colony cultures maintained on feeders. On day 7 post-induction, hiPSCs assumed a cobblestone-like morphology in the epithelial induction medium. Induced cells displayed increased mRNA expression levels of CK18, P63, and CK14 during the 28-day induction period. Furthermore, the expression levels of CK18, P63, and CK14 were detected via Western blotting and immunocytochemistry. Our findings suggest that hiPSCs cultured as single cells could be differentiated into epithelial cells.

Key words: Epithelial cell differentiation, Human induced pluripotent stem cells, Single-cell culture method, Feeder-free

Introduction

Epithelial tissues cover the body, cavity organs, and glands to function as barriers. Epithelial cells exhibit two polarities to form epithelial tissues: one is facing body or organ surface (the apical surface) and the other is facing connective tissues (the basal surface). While gastrulation occurs, cells are differentiated into three germ layers (ectoderm, mesoderm, and endoderm), followed by epithelial invagination into the primitive streak. The invaginated epithelium then forms the primitive gut, leading to the formation of the digestive tract, pancreas, liver, etc., and becomes the endoderm, leaving the ectoderm on the outer layer of the embryo.

Regenerative therapy is aimed at stimulating recovery from disease, injury, or aging through approaches such as the application of stem cells and/or progenitor cells to repair damaged tissues or organs and restore function. An ideal cell source should be easily obtainable and expandable in vitro. For example, as cell sources for epithelial tissue engineering, autologous cultured epidermis or oral mucosal epithelial cells have been applied in the clinical setting to treat burns or corneal disorders, respectively. However, the typical culture conditions for transplantable epithelial cells commonly require murine 3T3 feeder cells and an addition of fetal bovine serum to the culture because it is difficult to prevent pathogen transmission or infection when using animal-derived materials. In addition, a sheet derived from autologous retinal pigment epithelial cells was transplanted in the early 2000s, however, the procedure of harvesting the retinal pigment epithelial cell sheet is invasive and complex, and is a high risk of severe bleeding and retinal detachment. Furthermore, for tooth regeneration, epithelial cells should ideally be combined with mesenchymal cells to mimic the tooth development process. To date, studies of tooth regeneration have revealed that at the cap stage of embryonic day 14 or 14.5 mice, the application of murine tooth germ cells produced regularly-shaped teeth. However, porcine postnatal tooth germ cells applied during the late bell stage generated unusually shaped dental tissues. These results suggest that embryonic tooth germ cells at the cap stage retain their tooth-forming potential; however, using embryonic tooth germ cells is not feasible for clinical applications because of ethical issues. Thus, more suitable cell sources for applications in epithelial tissue engineering for regenerative therapy research are needed.

In 2006, Takahashi and Yamanaka reported the successful derivation of induced pluripotent stem cells (iPSCs) from mouse embryonic and adult fibroblasts, followed by the generation of human iPSCs (hiPSCs) from human dermal fibroblasts. Overcoming potential ethical issues associated with the use of embryonic stem cells (ESCs), iPSCs are derived from various somatic cells and can differentiate into the cell types...
of all three germ layers\textsuperscript{25-28}. Thus, hiPSCs have broad applicability in studies of disease mechanisms, drug screening, and regenerative medicine\textsuperscript{29-31}. Conventional iPSC culture has been performed in colony cultures using feeder cells, such as mouse embryonic fibroblasts (MEFs). However, long setup times are required, which increases the procedural complexity\textsuperscript{32}. Furthermore, human ESCs (hESCs) cultured on MEFs express an immunogenic non-human sialic acid on their surface, which may induce an immune response upon transplantation\textsuperscript{33}. To prevent the transmission of animal pathogens and the subsequent induction of an immune response, the clinical translatability of hiPSCs has recently led to improvements in the development of fully defined and xeno-free media and coating matrices for pluripotency cell (PSC) cultures\textsuperscript{44}. Few single-cell passaging methods have been reported, in which cells are cultured upon the addition of Rho-kinase inhibitors to improve the single-cell plating efficiency\textsuperscript{34-37}. However, the final cellular products of these methods include colony-type hESCs\textsuperscript{38,39}. Single cell-type culture, which prevents colony formation, is preferable because colony cultures result in slower expansion and cellular heterogeneity\textsuperscript{39}. Furthermore, novel medium and coating material supporting single-cell passaging and the expansion of human PSCs (hPSCs) as single-cell cultures in a feeder-free and defined environment have been developed\textsuperscript{44,39,41}.

Recently, Cellartis\textsuperscript{TM} DEF-CS\textsuperscript{®} 500 culture medium (DEF-CS medium) was shown to be very efficient for use in single-cell cloning\textsuperscript{44} and genome editing\textsuperscript{39}. iPSCs cultured in DEF-CS medium are passaged as single cells and expand as a homogeneous monolayer, thereby maintaining pluripotency with a stable karyotype\textsuperscript{39,41}. The efficient and scalable culture methods of hiPSCs and large-scale production of differentiated cells are prerequisites for regenerative therapy research and subsequent applications\textsuperscript{42}. This method was applied to pancreatic epithelium\textsuperscript{40} and neural differentiation\textsuperscript{39}; however, to the best of our knowledge, the differentiation of such PSCs cultured as single cells into surface ectodermal cells (hereafter, ectodermal-epithelial cells) has not been reported. Therefore, this study aimed to evaluate the feasibility of using single-cell and feeder-free cultures of hiPSCs to support differentiation into ectodermal-epithelial cells.

**Materials and Methods**

**Culturing of hiPSCs on feeders**

We purchased pregnant Institute for Cancer research (ICR) mice (n = 2, Chubu Kagaku Shizai Co., Ltd., Nagoya, Japan) and generated MEFs from 12-day mouse embryos with reference to a previously described method\textsuperscript{43}. This animal experiment was approved by the Animal Care and Use Committee for School of Dentistry, Aichi Gakuin University (approval number AGUD420). Animal care and experimental procedures were conducted in accordance with the Regulations on Animal Experimentation at School of Dentistry, Aichi Gakuin University.

The hiPSC line 253G1\textsuperscript{43} was obtained from the RIKEN BRC through the Project for Realization of Regenerative Medicine and the National Bio-Resource Project of the MEXT, Japan. As previously reported, undifferentiated hiPSCs were cultured on mitotically inactivated MEF feeder cells in 1:1 Dulbecco’s modified Eagle’s medium:Ham’s nutrient mixture F-12 (DMEM-F12; Sigma-Aldrich, St. Louis, MO, USA), 20% KNOCKOUT Serum Replacement (Thermo Fisher Scientific, Waltham, MA, USA), 0.1 mM nonessential amino acids (Thermo Fisher Scientific), 0.11 mM 2-mercaptoethanol (Thermo Fisher Scientific), 100 U/mL penicillin, 100 μg/mL streptomycin, and 2 mM L-glutamine (Sigma-Aldrich) supplemented with 5 ng/mL recombinant human basic fibroblast growth factor (Fujifilm Wako, Osaka, Japan)\textsuperscript{43}. Cultures were maintained at 37°C in a 5% CO\textsubscript{2} incubator with daily fresh medium changes.

**Culturing of hiPSCs as single cells**

Cultured hiPSC colonies as described above were disrupted using TrypLE Select (Thermo Fisher Scientific) and seeded in DEF-CS medium (Takara, Shiga, Japan) at a density of 2 × 10\textsuperscript{4} cells/cm\textsuperscript{2} on dishes coated with COAT-1 (Takara). The medium was replaced every 1–2 days. Feeder cells were eliminated through repeated passaging. The hiPSCs were then seeded at an approximate density of 4–5 × 10\textsuperscript{4} cells/cm\textsuperscript{2} and passaged every 3–4 days for further use. Before commencement of the experiment, the expression levels of pluripotency markers, including OCT3/4, SOX2, KL/F, and c-MYC, in hiPSCs cultured in colonies and as single cells were confirmed by performing polymerase chain reaction (PCR).

**Differentiation of hiPSCs cultured as single cells into epithelial cells**

The induction method reported by Cai et al.\textsuperscript{44}, originally used to differentiate hiPSCs and hESCs cultured in colonies into epithelial sheets, was used to differentiate hiPSCs as single cells into ectodermal-epithelial cells. Prior to the execution of the experiment, to ensure the same degree of epithelial differentiation potency of hiPSCs cultured as single cells and as colonies maintained on feeders, both types of hiPSCs were cultured in an epithelial induction medium for over a period of 7 days. hiPSCs cultured in colonies maintained on feeders were used as positive controls.

The experimental design using hiPSCs cultured in single cells was as indicated in Fig. 1. First, to determine a suitable coating material for differentiation into epithelial cells, hiPSCs as single cells were seeded (4 × 10\textsuperscript{4} cells/cm\textsuperscript{2}) in pre-coated dishes with three different types of coating; Collagen IV (Corning BioCoat Cellware, Mouse collagen type IV, 354416; Corning, Bedford, MA, USA), Matrigel (Corning BioCoat Matrigel Cellware, 354601; Corning), or COAT-1 (Takara). After culturing for 1–2 days, the medium was replaced with epithelial induction medium, DMEM-F12 (Sigma-Aldrich) supplemented with 1X N2 supplement (Thermo Fisher Scientific), 1 mM retinoic acid (RA; Fujifilm Wako), and 25 ng/mL bone morphogenetic protein (BMP4; Fujifilm Wako). hiPSCs were induced for a maximum of 28 days and the induction medium was replaced daily. Thereafter, the cells were observed using a phase-contrast microscope and counted for the indicated periods using a cell counter model R1 (Olympus, Tokyo, Japan). Each test was conducted in triplicate.

**PCR analysis**

Total RNA was extracted after cell differentiation (days 3, 7, 10, 14, 21, and 28) using the RNeasy Plus Mini Kit (Qiagen, Valencia, CA, USA).
USA), following the manufacturer’s instructions. RNA was extracted from the 253G1-derived neural crest cells (NCCs) as previously described and served as the negative control. RNA concentration was determined using a NanoDrop 1000 spectrophotometer (Thermo Fisher Scientific). First-strand cDNA was then synthesized from 100 ng of total RNA and amplified using ReverTra Ace qPCR RT Master Mix with gDNA Remover (Toyobo, Osaka, Japan). Subsequently, few PCR amplicons were separated by electrophoresis at 50 V on a 2% agarose gel on Mupid-2 (Advance Co. Ltd., Tokyo, Japan).

Expression profiling of genes encoding cell adhesion molecules

Expression profiling was performed by using an array of real-time PCR primers for genes encoding cell adhesion molecules (PrimerArray Cell adhesion molecules [Human] PH003; Takara) per manufacturer’s instructions. The primer array contained a mixture of 96 primer pairs for 88 genes encoding cell adhesion molecules and 8 housekeeping genes. Gene expression was quantified using the PrimerArray Analysis Tool version 2.2 (Takara).

Western blotting

Proteins were extracted using the Mammalian Protein Extraction Reagent (Thermo Fisher Scientific) per the manufacturer’s instructions after epithelial cell differentiation (days 3, 14, and 28). Protein concentration was determined using a BCA protein assay (Thermo Fisher Scientific) per the manufacturer’s instructions. For western blot analysis, equal amounts of proteins were separated on 4–15% Mini-PROTEAN TGX gels (Bio-Rad Laboratories, Hercules, CA, USA) and electro-transferred onto polyvinylidene difluoride membranes (Bio-Rad) for 1 h at 50 V. After blocking membranes with 1X Tris-buffered saline (TBS) with 1% casein (Bio-Rad), they were probed for 18 h at 4°C with primary antibodies, including rabbit monoclonal anti-P63 (1:500; Abcam, Cambridge, MA, USA), mouse monoclonal anti-cytokeratin (CK) 14 (1:500; Abcam), mouse monoclonal anti-CK18 (1:500; Abcam), and mouse monoclonal anti-GAPDH (1:10,000; Thermo Fisher Scientific) antibodies. The membrane was washed with TBS with Tween 20 and then probed for 1 h at 25°C with either preabsorbed goat anti-mouse IgG (Fujifilm Wako) and rabbit IgG (Gene Tex, Irvine, CA, USA). The secondary antibodies, Alexa Fluor 488 goat anti-mouse IgG (1:200 for anti-P63 antibody; Thermo Fisher Scientific) and Alexa Fluor 594 goat anti-rabbit IgG (1:200 for anti-CK18 antibody; Thermo Fisher Scientific) antibodies, were added and incubated for 1 h at room temperature (25°C). Cells were then counterstained with 1 μg/ml 4',6-diamidino-2-phenylindole (DAPI; Dojindo, Kumamoto, Japan) and observed under a BZX-710 fluorescence microscope (Keyence, Osaka, Japan). To determine the percentage of epithelial marker-positive cells, CK18-, P63-, and CK14-positive and total cells were enumerated and observed under a BZX-710 fluorescence microscope (Keyence, Osaka, Japan).

Immunocytochemistry

Induced cells were fixed with 4% paraformaldehyde in phosphate-buffered saline (PBS) and rendered permeable with 0.1% Triton X-100 (Kanto Chemical, Tokyo, Japan) in PBS. The cells were blocked using 2.5% normal horse serum (Vector Laboratories, Burlingame, CA, USA) for 1 h and then incubated with the following primary antibodies: rabbit monoclonal anti-P63 (1:100; Abcam) and mouse monoclonal anti-CK14 (1:100; Abcam), or CK18 (1:200; Abcam) antibodies for 1 h at room temperature (25°C). Isotype controls were normal mouse IgG (Fujifilm Wako) and rabbit IgG (Gene Tex, Irvine, CA, USA). The secondary antibodies, Alexa Fluor 488 goat anti-mouse IgG (1:200 for anti-CK14 antibody and 1:1,000 for anti-CK18 antibody; Thermo Fisher Scientific), and Alexa Fluor 594 goat anti-rabbit IgG (1:200 for anti-P63 antibody; Thermo Fisher Scientific), were added and incubated for 1 h at room temperature (25°C). Cells were then counterstained with 1 μg/ml 4',6-diamidino-2-phenylindole (DAPI; Dojindo, Kumamoto, Japan) and observed under a BZX-710 fluorescence microscope (Keyence, Osaka, Japan).
and cell proliferation of hiPSCs cultured as single cells during epithelial cell differentiation

Three different coating materials were prepared to induce hiPSCs cultured as single cells into epithelial cells (Fig. 1). When hiPSCs were seeded in the COAT-1-coated dishes (Fig. 5A), the cells were stably cultured without further passaging for 28 days. Cells cultured in the induction media displayed a cobblestone-like morphology, resembling that of epithelial cells, beginning on approximately day 7 of induction (Fig. 5B), and displayed a higher density after day 14 (Fig. 5C, D). In the case of hiPSCs seeded in Collagen IV-coated dishes, the cells adhered to the dish at 12 h (Fig. 5E). However, they detached 24 h after seeding (Fig. 5F). Simultaneously, hiPSCs seeded in the Matrigel-coated dishes adhered to the dish (Fig. 5G). However, all cells detached on day 6 of induction (Fig. 5H). Based on these results, hiPSCs cultured as single cells seeded on COAT-1-coated dishes clearly showed an epithelial cell-like morphology and were used for subsequent experiments. We then observed cell proliferation. The number of induced cells on day 10 increased by 4.8-fold compared to that on day 3, whereas the cell number

Figure 3. Phase-contrast microscopic image of human induced pluripotent stem cells (hiPSCs) cultured in colonies on feeders. (A) hiPSCs were maintained in an undifferentiated state on mouse embryonic fibroblast feeders. (B) Higher magnification of the area demarcated in (A). (C) hiPSCs cultured in an epithelial induction medium for 7 days displayed a cobblestone-like morphology (arrows). Black scale bar, 500 μm; white scale bars, 100 μm.

Figure 2. Expression of pluripotency markers in human induced pluripotent stem cells (hiPSCs) cultured as single cells and in colonies on feeders. hiPSCs cultured as single cells expressed pluripotency markers including OCT3/4, SOX2, KLF4, and c-MYC, similar to those cultured in colonies maintained on feeders. To rule out the presence of genomic DNA contamination, the same procedure was conducted without performing reverse transcription. All PCR products, except for GAPDH (28 cycles), were subjected to 32 cycles. RT (-): negative control.

**Results**

**Expression of pluripotency markers in hiPSCs cultured as single cells**

Electropherograms of the PCR products are illustrated in Fig. 2. The expression levels of pluripotent markers, including OCT3/4, SOX2, KLF4, and c-MYC, were observed in hiPSCs cultured as single cells or in colonies maintained on feeders.

**Differentiation potential of hiPSCs cultured as single cells and in colonies maintained on feeders**

After hiPSCs cultured in colonies (Fig. 3A, B) were maintained in the epithelial induction medium for 7 days, the induced cells displayed a cobblestone-like morphology (Fig. 3C). Thereafter, the expression levels of epithelial cell markers in induced cells derived from hiPSCs cultured in colonies on feeders and as single cells were compared. In both cases, the mRNA expression levels of epithelial cell markers in induced cells were significantly higher than those in undifferentiated hiPSCs (Fig. 4).

**Morphology and cell proliferation of hiPSCs cultured as single cells during epithelial cell differentiation**

Three different coating materials were prepared to induce hiPSCs cultured as single cells into epithelial cells (Fig. 1). When hiPSCs were seeded in the COAT-1-coated dishes (Fig. 5A), the cells were stably cultured without further passaging for 28 days. Cells cultured in the induction media displayed a cobblestone-like morphology, resembling that of epithelial cells, beginning on approximately day 7 of induction (Fig. 5B), and displayed a higher density after day 14 (Fig. 5C, D). In the case of hiPSCs seeded in Collagen IV-coated dishes, the cells adhered to the dish at 12 h (Fig. 5E). However, they detached 24 h after seeding (Fig. 5F). Simultaneously, hiPSCs seeded in the Matrigel-coated dishes adhered to the dish (Fig. 5G). However, all cells detached on day 6 of induction (Fig. 5H). Based on these results, hiPSCs cultured as single cells seeded on COAT-1-coated dishes clearly showed an epithelial cell-like morphology and were used for subsequent experiments. We then observed cell proliferation. The number of induced cells on day 10 increased by 4.8-fold compared to that on day 3, whereas the cell number

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Figure 4. Expression of epithelial marker genes in human induced pluripotent stem cells (hiPSCs) on day 7 of culture in epithelial induction medium as single cells compared with that in colonies on feeders. The relative expression levels of P63, CK18, and CK14 were significantly higher in induced cells than those in hiPSCs cultured as single cells, and were similar to those in colonies. GAPDH was used as an internal control. Data are represented as mean ± SD (n = 3). *P < 0.05, **P < 0.01.

Figure 5. Phase-contrast microscopic image of human induced pluripotent stem cells (hiPSCs) cultured as single cells on three different types of coated dishes for epithelial cell differentiation. (A–D) Cells seeded on COAT-1-coated dishes. (A) hiPSCs were cultured in the DEF-CS medium for 48 h. (B) hiPSCs cultured in an epithelial induction medium for seven days were morphologically similar to epithelial cells (arrows). (C) The density of the induced cells increased on post-induction day 14. (D) Induced cells were slightly detached on post-induction day 28. (E, F) Cells seeded in Collagen IV-coated dishes. (E) hiPSCs adhering to the bottom of the dish after 12 h of seeding before induction. (F) hiPSCs detached from the bottom of the dish after 24 h of seeding before induction. (G, H) Cells seeded in Matrigel-coated dishes. (G) hiPSCs adhered to the bottom of the dish after 48 h of seeding. (H) Induced cells detached from the bottom of the dish 6 days after replacement of the DEF-CS medium with the epithelial induction medium. White scale bars, 100 μm; black scale bars, 500 μm.
Characteristics of induced cells derived from hiPSCs cultured as single cells during epithelial cell differentiation

Real-time PCR

The mRNA expression levels of genes encoding pluripotency marker \((OCT3/4)\), epithelial markers \((P63, CK14, and CK18)\), and cell adhesion molecule markers \((cadherin 1; CDH1, and claudin 1; CLDN1)\) in the induced cells were determined via the \(\Delta\Delta C_t\) method (Fig. 7). \(OCT3/4\) mRNA was expressed on day 3; however, the expression was downregulated on day 7. \(CK18\) mRNA expression was upregulated by days 3–7, prior to \(P63\) and \(CK14\) upregulation. \(P63\) expression was progressively upregulated after day 7, and \(CK14\) expression was upregulated on day 21, followed by an upregulated expression of \(P63\). Furthermore, \(CLDN1\) expression was progressively upregulated after day 21. \(CDH1\) expression was slightly downregulated on day 28 compared to that on days 3–21. The experiments were conducted in triplicate. The expression patterns of these markers displayed similar trends. Genes encoding pluripotency, epithelial, and cell adhesion molecule markers were examined using RNA extracted from hiPSC-derived NCCs as the negative control.

Figure 6. Proliferation potential of cells induced from human induced pluripotent stem cells cultured as single cells in epithelial induction medium for 28 days. The cells were counted using a cell counter model R1 (Olympus) on the indicated days. The induced cells proliferated in a linear manner from post-induction day 3 to 10. Data are presented as mean ± SD (n = 3).

Figure 7. Quantification of the expression of pluripotency and epithelial marker genes in cells induced from human induced pluripotent stem cells cultured as single cells on days 3, 7, 10, 14, 21, and 28 after epithelial induction as evaluated using real-time PCR. Pluripotency marker \(OCT3/4\) expression was downregulated; in contrast, expression levels of epithelial cell markers \(P63, CK14, and CK18\) and cell adhesion markers \(CDH1\) and \(CLDN1\) were upregulated during epithelial induction. Neural crest cells were used as the negative control. Data are presented as mean ± SD of triplicate experiments.
Expression profiling of genes encoding cell adhesion molecules

The expression of 88 cell adhesion molecule-related genes using the primer array was examined. The differences in gene expression between induced cells on days 14 and 28 were compared. Furthermore, genes encoding adhesion molecules with >2-fold differences in expression levels on days 14 and 28 were selected and further investigated using real-time PCR. The expression levels of the three selected genes (inducible T-cell co-stimulator ligand ICOSLG, neural cell adhesion molecule 1 NCAM1, and syndecan 1 SDC1) on day 28 were >2-fold compared to those observed on day 14 (Table 2).

Western blotting

Epithelial marker protein expression was assessed via western blotting. CK18 expression was detected on day 3 and was remarkably expressed on days 14 and 28. P63 expression was detected on days 14 and 28. CK14 expression was only detected on day 14 (Fig. 8).

Immunocytochemistry

Immunocytochemistry was performed to determine the proportion of P63-, CK14-, and CK18-positive cells. CK18 expression was detected on day 7 of induction (Fig. 9A), whereas P63 and CK14 expression levels were detected on day 14 (Fig. 9B, C). The proportion of CK18-positive cells on day 7 was 48.0%; on day 14, the proportion of P63-positive cells was 67.4%, and that of CK14-positive cells was 18.9%.

Table 2. Expression profiling of adhesion molecule genes

| Symbol | Name                                | Accession Number |
|--------|-------------------------------------|------------------|
| ICOSLG | Inducible T-cell co-stimulator ligand | NM_01529         |
| NCAM1  | Neural cell adhesion molecule 1     | NM_181351        |
| SDC1   | Syndecan 1                          | NM_002997        |

| Symbol | Relative Expression Day 14 | Day 28 |
|--------|----------------------------|--------|
| ICOSLG | 2.9                        | 10.3   |
| NCAM1  | 4.7                        | 164.7  |
| SDC1   | 4.0                        | 29.9   |

Figure 8. Immunoblots of epithelial marker proteins in cells on post-induction days 3, 14, and 28 during epithelial differentiation. GAPDH was used as the loading control.

Figure 9. Immunocytochemistry of epithelial marker proteins in cells on post-induction days 7 and 14 during epithelial cell differentiation. (A) CK18 was detected on post-induction day 7 in hiPSCs. (B, C) P63 and CK14 were detected on post-induction day 14 in hiPSCs. The induced cells were not stained with normal mouse IgG (D, E) and rabbit IgG (E) as isotype controls. The cells on post-induction day 14 (B, C, E, and F) were denser than those on day 7 (A, D). Overlaid images of DAPI-stained slides are shown. Scale bars, 100 μm.
Discussion

In this study, we evaluated the feasibility of growing hiPSCs as single-cell derived, feeder-free cultures and investigated whether hiPSCs cultured as single cells differentiated into ectodermal-epithelial cells. Results showed that single cell-derived hiPSCs cultured in an epithelial induction medium supplemented with RA, BMP4, and N2 displayed a cobblestone-like morphology and exhibited key characteristics of ectodermal-epithelial cells in vitro.

The hiPSC line 253G1 cultured in colonies on MEFs was transferred and cultured as single cells using the DEF-CS medium and the accompanying coating material COAT-1. The hiPSCs cultured as single cells expressed pluripotency markers similar to those using a conventional colony culture method with MEFs. PCR revealed that hiPSCs cultured from single cells expressed OCT3/4, SOX2, KLF4, and c-MYC at levels comparable to those present in colonies on MEF. Maintenance of pluripotency is governed by a set of transcription factors primarily involving OCT3/4, SOX2, and NANO1 in ESCs50. Our findings indicated that hiPSCs cultured as single cells retained their pluripotency.

Feeder cells may contribute to xenogenic contamination and lot-to-lot variability, impacting the final results of culturing51. The direct differentiation of hiPSCs using defined chemical components is preferable. Therefore, we investigated whether hiPSCs cultured as single cells could be directly differentiated into ectodermal-epithelial cells, as per previously reported protocols for hiPSCs cultured in colonies50,52, wherein media supplemented with 1 μM RA, 25 ng/ml BMP4, and 1X N2, were used. We found that these cells could differentiate.

Epithelial development is characterized by the expression of specific markers. The body surface is first covered by a single layer of basal epithelial cells during fetal development. This primitive epithelium activates a differentiation program leading to stratification during development; CK18-positive cells in the primitive epithelium are substituted with CK14-positive basal epithelial cells. Furthermore, P63 regulates the development of the stratified epithelia and is required for the commitment of PSCs to CK14-positive stratified epithelial cells53,54. Our results indicated similar patterns with normal stratified epithelial development markers associated with the EMT, namely ICOSLG, NCAM1, and SDC1, peaked on post-induction day 28, concurrent with a previous report which stated that NCAM1 expression was upregulated and CDH1 expression was downregulated during EMT progression55.

We also found that the cells on post-induction day 28 were negative for CK14 expression, although this protein was detected on post-induction day 14, different from the results of the real-time PCR results. We attributed this to the presence of extensive disulfide cross-linking of keratins that yields insoluble proteins56.

One limitation of this study is that only one hiPSC line, 253G1, was used. It is necessary to differentiate several hiPSC lines cultured as single cells into ectodermal-epithelial cells, as the epigenetic memory of their original phenotypes may limit their differentiation potential57,58.

In conclusion, this study demonstrated the derivation of ectodermal-epithelial cells from hiPSCs cultured as single cells in a simple manner and found that hiPSCs from single-cell derived cultures maintain their pluripotency and differentiation potential. To the best of our knowledge, this is the first to report of the differentiation of passaged hiPSCs expanded as single cells into ectodermal-epithelial cells. The combination of this single cell-based culture method for hiPSCs with an ectodermal-epithelial differentiation protocol may facilitate the large-scale generation of epithelial cells in regenerative therapy research and their subsequent applications. Further studies are needed to analyze the molecular-level events related to odontogenic potential and determine the optimal epithelial cell differentiation period for use in tooth regeneration studies.

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Conflict of Interests

The authors have declared that no COI exists.

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