Endodermal differentiation of human induced pluripotent stem cells using simple dialysis culture system in suspension culture

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A differentiation of human induced pluripotent stem cells (hiPSCs) into definitive endoderm lineage is required for a preparation of metabolic organ derived cells. The differentiation consumed high-priced cytokines and small molecules, which have hampered the manufacturability of differentiated cells. Although the cytokines and small molecules are remained or cells produce the autocrine factors, daily culture medium change should be proceeded to remove toxic metabolites generated from cells. In this study, we developed a simple dialysis culture system to refine the medium during definitive endodermal differentiation. We demonstrated that dialysis culture prevented cell damage to remove lactate. The hiPSCs cultured with dialysis also differentiated similarly as usual differentiation without dialysis even if they were not supplied Activin A for latter culture days in the differentiation. With this dialysis culture system, hiPSCs were differentiated into endodermal lineage with medium refinement and recycling and autocrine factors as well as cytokines, which may lead to reduce differentiation cost.

The required transplanted cell number of pancreatic β cells which is expected to be a replacement of donor derived pancreatic islet is more than tens of million cells. A suspension culture is profitable compared to the adherence culture which depends on the cultured surface area. However, the increase of cultured cell number causes increment of toxic metabolite including lactate and ammonium. A energy metabolic pathway related to lactate production of hiPSCs is activated during differentiation, which is expected actively production of lactate from differentiated cells. Considering the removal toxic metabolite, daily medium change is generally performed.

Definitive endodermal differentiation is generally performed to prepare metabolic organ derived cells; hepatocyte, pancreatic β cells, kidney podocytes, and intestine epithelial cells are generally prepared with definitive endodermal differentiation from hiPSCs using Activin A. The existence of autocrine factors produced by hiPSCs has been reported. Nodal is one of the autocrine factor and perform similarly as Activin A in their pathway.
The recycling of autocrine factors is promising to reduce the usage of costly cytokines for the differentiation. However, daily medium change looses those autocrine factors with removal of waste including toxic metabolites produced by cells.

Dialysis culture is a potential solution to refine the culture medium with retaining of cytokines and autocrine factors. Previous study focused on mass production and their reactors were generally large scale using hollow fiber or perfusion culture [14,15]. The application dialysis culture to hiPSCs culture was reported for expansion of hiPSCs [15]. However, the definitive endoderm differentiation using dialysis culture has not been studied yet.

In this study, we developed a plate-based dialysis culture system to differentiate hiPSCs into definitive endoderm. We also demonstrated that the differentiation without Activin A addition at the latter culture days of differentiation to evaluate the existence of autocrine factor influences toward the differentiation. Our dialysis culture suggested that there were continuous glucose supply, lactate removal, and autocrine factors retaining. However, further study to remove the higher molecular toxic metabolite than 3 kDa should be performed to improve the culture condition. This study is expected to contribute the manufactural application to prepare the plenty number of cells with keeping the usage of costly cytokines down.

2. Materials and methods

2.1. Preparation of dialysis culture plate

A dialysis culture plate was prepared with deep well plate (Corning, USA) and a culture insert (Corning, USA), which had a dialysis membrane (Spectrum, USA). The polycarbonate membrane of the culture insert was removed and replaced with a dialysis membrane, which had a molecular weight cutoff (MWCO) of 3 kDa and a surface area of 0.53 cm² (hereinafter, this is called “dialysis cup”). The bottom of the dialysis membrane was covered with a polyethylene film (Diversified Biotech, USA) to avoid mass transfer as the comparison to dialysis culture (hereinafter, this is called “blocked dialysis cup”) (Fig. 1a).

2.2. Maintenance of hiPSC and formation of hiPSC aggregates

The human iPS cell line TkDN4-M was provided by Stem Cell Bank at Institute of Medical Science, The University of Tokyo. Cells were maintained on a tissue-culture treated 6 well plate coated with 10 µg/cm² vitronectin human recombinant protein (VTN-N; Thermo Fisher, USA) in Essential 8 medium (EB; Thermo Fisher Scientific, USA) containing 1% penicillin–streptomycin–amphotericin B suspension (PSA; FUJIFILM Wako Pure Chemical Corporation, Japan). The hiPSCs were seeded at 1.0 × 10⁴ cell/cm² with EB medium containing 1% PSA and 10 µM Rho-Associated Kinase inhibitor (Y-27632) and were cultured at 37 °C in a humidified atmosphere with 5% CO². Culture medium was changed every day with EB medium containing 1% PSA. Cells were detached by Accutase (Innovative Cell Technologies, Inc., USA) after washing with phosphate buffered saline (PBS; FUJIFILM Wako Pure Chemical Corporation, Japan) for passaging and starting aggregate formation.

For aggregate formation, hiPSCs aggregates were seeded at 7.5 × 10⁵ cell/mL in 4 mL of EB medium containing 5 mg/mL bovine serum albumin (BSA; Proliand Biologicals, USA), 1% PSA, and 10 µM Y-27632 in a 6 well plate for suspension culture (Sumitomo Bakelite Co., Ltd., Japan). The plate was rotated at 90 rpm on an orbital shaker. The aggregates were formed within 24 h at 37 °C in a humidified atmosphere with 5% CO².

2.3. Differentiation of hiPSC into definitive endoderm lineage

One day before starting hiPSC differentiation in a suspension culture, the culture medium was replaced with DMEM/Ham’s F12 medium (FUJIFILM Wako Pure Chemical Corporation, Japan) containing 20% knockout serum replacement (KSR; Thermo Fisher Scientific, USA), 1% non-essential amino acids (NEAA; FUJIFILM Wako Pure Chemical Corporation, Japan). Cells were detached by Accutase (Innovative Cell Technologies, Inc., USA) after washing with phosphate buffered saline (PBS; FUJIFILM Wako Pure Chemical Corporation, Japan) for passaging and starting aggregate formation.

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Fig. 1. The culture condition in this study. (a) Schematic images of culture design. (b) Schedule of cytokines addition for the differentiation.
Wako Pure Chemical Corporation, Japan) and 7.5 ng/mL recombinant human fibroblast growth factor 2 (bFGF; Reprocell Inc., Japan). Right before differentiation, hiPSCs aggregates in a 6 well plate were collected and partially dissociated using Accutase for cell counting. The hiPSCs aggregates were then resuspended into 2.5 mL at 1 x 10^6 cell/mL and transferred in the dialysis cup or the blocked dialysis cup.

The differentiation of iPSCs into definitive endoderm (DE) was performed as described previously with some modification [9]. The hiPSCs aggregates were cultured in a differentiation medium consisted of a basal medium with cytokines and small molecules for four days. The basal medium was also fed in the deep well and changed every day with fresh basal medium; 3 μM CHIR-99021 (FUJIFILM Wako Pure Chemical Corporation, Japan) was added first day. The basal medium was prepared with RPMI1640 (FUJIFILM Wako Pure Chemical Corporation, Japan) containing 5 mg/mL BSA, 1% sodium pyruvate, 1% NEAA, 1% PSA, and 55 μM 2-mercaptoethanol (Thermo Fisher Scientific, USA).

We compared four different culture conditions to evaluate the culture medium refinement and autocrine existence which promoted differentiation (Fig. 1b). First, the hiPSCs aggregates were cultured in the blocked dialysis cup for four days with every day a differentiation medium change using the basal medium containing 80 ng/mL recombinant human/mouse/rat Activin A (R&D Systems, USA); 3 μM CHIR-99021, 20 ng/mL human bone morphogenetic protein (BMP)-4 (Miltenyi Biotec K.K., Japan) and 50 ng/mL bFGF were added for first day (herein after, this is called “Control”). Second, the hiPSCs aggregates were cultured in the dialysis cup for four days with every day a differentiation medium change using basal medium containing 80 ng/mL recombinant human/mouse/rat Activin A; 3 μM CHIR-99021, 20 ng/mL human BMP-4 50 ng/mL bFGF were added for first day (herein after, this is called “DC1”). Third, the hiPSCs aggregates were cultured in the dialysis cup for four days with first two days a differentiation medium change using basal medium containing 80 ng/mL recombinant human/mouse/rat Activin A; 3 μM CHIR-99021, 20 ng/mL human BMP-4 50 ng/mL bFGF were added for first day. The hiPSCs aggregates were cultured for latter two days without a differentiation culture medium change but with removal of dead cells by centrifugation and added 80 ng/mL recombinant human/mouse/rat Activin A (herein after, this is called “DC2”). Fourth, the hiPSCs aggregates were cultured in the dialysis cup for four days with first two days a differentiation medium change using basal medium containing 80 ng/mL recombinant human/mouse/rat Activin A; 3 μM CHIR-99021, 20 ng/mL human BMP-4 50 ng/mL bFGF were added for first day. The hiPSCs aggregates were cultured for latter two days without a differentiation culture medium change but with removal of dead cells by centrifugation (herein after, this is called “DC3”).

2.4. Biochemical assays

After days of culture, the glucose and lactate concentration of the culture medium in the dialysis, blocked dialysis, and deep well were measured by bioanalyzer (Oji analytical, Japan) to evaluate the changing because of dialysis. Lactic dehydrogenase (LDH) in the medium was measured with a cytotoxicity LDH assay kit-WST (FUJIFILM Wako Pure Chemical Corporation, Japan) following the manufacture protocol. The differentiated hiPSCs aggregates were dissociated and counted their number.

2.5. RT-qPCR analyses

Total RNAs of hiPSCs aggregates sampled before and after differentiation were isolated by using TRIzol reagent (Thermo Fisher Scientific, USA). Their RNAs were purified by adding chloroform (FUJIFILM Wako Pure Chemical Corporation, Japan) and the centrifugation of 15,000 x g for 15 min. The supernatants were collected and purified by adding 2-propanol (FUJIFILM Wako Pure Chemical Corporation, Japan) and the centrifugation of 15,000 x g for 15 min. Following washing with 75% ethanol (FUJIFILM Wako Pure Chemical Corporation, Japan), purified RNA samples were qualitative and quantitative analyzed by Nano drop (Shimadzu, Japan). 100 ng of purified RNA from each sample was reverse-transcribed by PrimeScriptTM Reverse Transcriptase (Takara Bio Inc., Japan). The transcribed complementary DNA samples were evaluated their gene expression related to pluriptenotity (Oct4), definitive endoderm (Sox17, Foxa2) and primitive gut tube (Hnf4α) markers using SYBR Green (TOYOBO CO., LTD.) and detected by StepOne Plus (Thermo Fisher Scientific, USA). The specific primers were used as follows. β-actin: forward, 5’-CCTCATGAA-GATCCTCACC-3’; reverse, 5’-TTGCAATGTTGATGACCTGG-3’. Oct4: forward, 5’-AGTTGGTGAGAAGTCAAC-3’; reverse, 5’-TCCTGTTGATGTCGTCGTGTA-3’. Sox17: forward, 5’-CGACATT- CAGACCTGACAACA-3’; reverse, 5’-TCTGCTCTCCTCACAAGG-3’. Foxa2: forward, 5’-GGTTGATCTGTGTTGTTGTTG-3’; reverse, 5’- GCGCACTGTCATGATGTCGTT-3’. Hnf4α: forward, 5’-CCAGA- GATCCTAGTTGTCCTCAA-3’; reverse, 5’-TTGAATGACTCCTCCAGTCTCA-3’.

2.6. Statistical analyses

Results were expressed as the mean ± standard deviation of the mean (SD). The Student’s t-test was performed to assess the statistical significance of the results. A value of P< 0.05 was considered significant.

3. Results and discussion

3.1. Formation and differentiation of hiPSCs aggregates

The hiPSCs aggregates were formed with rotational suspension culture (Fig. 1a). Whereas the morphology of each condition during differentiation seemed similarly, slight larger number of dead cells was found in DC2 on day 3. Activin A is known as the cytokine which promotes survival and proliferation [16]. The proinflammatory, tumor-promoting, and metastasis-promoting effects of Activin A have been also reported [17-19]. In DC2, Activin A was added daily without medium change for two days from second day of differentiation, which might cause cellular damages because of the negative effect of Activin A.

3.2. Effects of dialysis culture

The remaining glucose and produced lactate of the medium in the dialysis cup and blocked dialysis cup for every day to evaluate the dialysis effect on glucose supply and lactate removal (Fig. 2). Whereas Control culture consumed all of the glucose in the differentiation medium, DC1, DC2, and DC3 culture with dialysis maintained near the original glucose concentration of the medium (Fig. 2a). The glucose concentration in the deep well were also decreased as the same level of the differentiation medium but still enough glucose was remained in dialysis culture group; DC1, DC2 and DC3. The glucose concentration of the medium in the deep well didn’t change in the Control culture, which suggested that dialysis membrane was completely blocked and didn’t leak (Fig. 2c). The lactate concentration in the differentiation medium of Control was four times higher than that of in dialysis culture; DC1, DC2, and DC3 (Fig. 2b). The lactate concentration of the medium in the deep well was the same as in the differentiation medium in dialysis culture; DC1, DC2, and DC3, but there was no difference in the control.
culture because the dialysis membrane was completely sealed (Fig. 2d). Therefore, continuous glucose supply to hiPSCs and lactate removal from differentiation medium though a dialysis membrane were successfully performed by dialysis culture until they were equilibrium between differentiation medium and the medium in the deep well.

The cellular damages were evaluated by measurement of LDH changes in the differentiation medium (Fig. 3). DC2 and DC3 showed higher value than other two conditions because of the non-medium change. Cellular damage of DC1 might lower than that of Control throughout the whole differentiation days.

The differentiated cell number seemed to have correlation with the results of LDH assay (Fig. 4). The cell number of DC1 was larger than that of other conditions; Control, DC2, and DC3. The cell number of Control, DC2, and DC3 had no statistical differences. The higher cell damage of DC2 and DC3 between days 3 and 4 suggested that there were toxic macromolecule larger than dialysis MWCO; 3 kDa in the medium. The energy metabolism of hiPSCs shift oxidative phosphorylation from glycolysis during a differentiation [20]. The oxidative phosphorylation shift of hiPSCs has potential to provoke reactive oxygen spices, which reduce hemo oxygenase-1. The lack of hemo oxygenase-1 promotes cellular damage and differentiation of pluripotent stem cells [21]. When the cells are induced apoptosis or autophagy cytotoxic macromolecules are eliminated, e.g. caspase (30–60 kDa) or cytochrome c (12–15 kDa) [22,23]. To avoid the effect those macromolecules, MWCO of

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**Fig. 2.** Glucose and lactate concentration transitions in the culture medium during differentiation. (a) glucose concentration of differentiation medium, (b) lactate concentration of differentiation medium, (c) glucose concentration of medium in deep well, and (d) lactate concentration of medium in deep well. Data represent mean ± SD of $n = 5$–12 from two to four independent experiments.

**Fig. 3.** Lactate dehydrogenase (LDH) cytotoxicity assay on day 4. Data represents mean ± SD of $n = 5$–6 from two independent experiments.

**Fig. 4.** Differentiated cell number on day 4. Data represents mean ± SD of $n = 5$–9 from two to four independent experiments ("**P < 0.01)."
dialysis membrane should be increased or further refinement to remove cytotoxic macromolecule is necessary.

3.3. Gene expression of differentiated hiPSCs aggregates

The gene expression of differentiated hiPSCs aggregates was evaluated (Fig. 5). There were no statistical differences between all conditions; Control, DC1, DC2, and DC3 for the gene expression of OCT4, SOX17 and FOXA2. OCT4 expression of differentiated hiPSCs aggregates were statistically lower than hiPSCs sampled right before starting differentiation, which suggests that the hiPSCs aggregates were successfully differentiation, whereas HNF4a expression of DC2 was lower than DC1. The transition of FOXA2 and SOX17 expression is similarly increased at the early stage of endodermal differentiation [24,25]. The HNF4a expression starts to increase behind the expression of FOXA2 and SOX17 during primitive gut tube development [26]. The decreased expression of HNF4a in DC2 suggested that the excess amount of Activin A inhibited the differentiation. DC3 was similarly differentiated as the Control and DC1 according to the gene expression even if Activin A was not added for two days. Therefore, it was suggested that dialysis culture enabled Activin A usage reduction by recycling of retaining cytokines and autocrine factors.

4. Conclusion

We developed a plate-based simple dialysis culture system for a differentiation of hiPSCs into definitive endodermal lineage. The dialysis culture refined the differentiation medium, which achieved the increase of differentiated cells. We also demonstrated that the reduction of Activin A usage didn’t no difference on the gene expression related to definitive endoderm and primitive gut tube markers. The retaining of cytokines and autocrine factor enabled to promote definitive endodermal differentiation without daily Activin A addition. This study will be expected to contribute to improve the manufacturability of hiPSCs differentiated into endodermal lineage with medium refinement and recycling autocrine factors as well as cytokines.

Author contributions

M.S., H.C., M.I. and Y.S. conceived and designed the study. M.S., H.C. and M.I. conducted the experiments. H.C. and M.S. provided all the figures. M.S., M.I, SG.Y acquired the hiPS culture and differentiation technique. M.S., H.C., M.I, SG.Y, H.O, A.M, and Y.S interpreted and analyzed the data. M.S., H.C., M.I, SG.Y, H.O, A.M, and Y.S revised the manuscript.

Conflict of interest

The authors have no conflict of interest to declare.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.reth.2019.05.004.
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