

Characterization of Human Dental Pulp Cells from Supernumerary Teeth by Using Flow Cytometry Analysis

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Abstract

The aim of this study was to analyze cells from human dental pulp tissue of impacted supernumerary teeth as stem cells with flow cytometry. Human dental pulp cells from 15 supernumerary teeth were identified their characteristics as stem cells by expression of mesenchymal stem cell markers through flow cytometry analysis at passage 3 and passage 10. Cluster of differentiation (CD) 73, CD 90, CD 34, CD 45 and STRO-1 cell surface markers were used to figure out characteristics of dental pulp stem cells from supernumerary teeth. At passage 3, the cell population showed positive expression of CD 73, CD90 and STRO-1, lacked expression of CD 34 and CD 45. At passage 10, CD 73, CD 90 and STRO-1 showed positive expression while CD 34 and CD 45 showed negative expression. This study indicated that dental pulp stem cells of supernumerary teeth had the properties of mesenchymal stem cells at both early and late passage. Impacted supernumerary teeth could be considered as a noble source of stem cells because of rapid growth and maintaining characteristics of stem cells until late passage.

Key words : Supernumerary tooth, Dental pulp stem cell, Flow cytometry

I. Introduction

Stem cells in human body have infinite potentials for cure or acceleration of healing in damaged tissue. Nevertheless, sources of stem cells to harvest out from human bodies are limited because of the difficulty of extracting many pure stem cells and the necessity of additional invasive procedure. Recently, it has been trying to find various sources of stem cell[1]. Embryonic stem cells have been treated as conventional stem cells for the last half century[2]. But the embryonic stem cells have limited supply and ethical issues. To replace the embryonic stem cells, studies on adult stem cells from dental tissues had

been reported on various sources including dental pulp, extracted deciduous tooth, periodontal ligament, dental sac and apical follicle[3-7]. Most of stem cell researches in the dental field were related to the cells from wisdom teeth or deciduous teeth. However, the wisdom teeth have limitation of relatively late extraction time. Deciduous teeth exfoliate in an early age, but most of them has a physiologic root resorption and incomplete pulp tissue remnants. Impacted supernumerary teeth (SNT) can removed without damage of pulp tissue during surgery procedure at an early age[8].

The International Society for Cellular Therapy (ISCT) proposed minimal criteria for identifying mesenchymal stem cell

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(MSC) in 2006[9]. Previous studies of adult stem cells confirmed as MSCs by applying these criteria[10-12] while most studies used data obtained from a few or one donor.

The purpose of this study is to analysis characteristics of MSC of dental pulp from 15 impacted SNT.

II. Materials and Methods

1. Materials

Impacted supernumerary teeth in the anterior maxilla were extracted from 15 healthy patients who were between 6 and 9 years old. This study was approved by the Dankook university dental college institutional ethics committee (H-1506/006/001). Supernumerary teeth were kept in α -minimum essential medium (α -MEM, Gibco) with 20% fetal bovine serum (FBS, Gibco), 100 U/mL penicillin, 100 μ g/mL streptomycin (Gibco), 2 mM L-glutamine (Gibco) and 10 nM L-ascorbic acid (Sigma) immediately after extraction.

2. Methods

1) Cell preparation and culture

Supernumerary teeth were cut around at cementoamel junction with dental disc until before exposing dental pulp. The crown and root were split along the groove to obtain dental pulp tissue with sterile dental file. Dental pulp tissue was chopped to fine slices less than 1 mm. Dental pulp cells were extracted using enzymatic digestion method with 3 mg/mL type I collagenase (Sigma-Aldrich) and 4 mg/mL Dispase (Sigma-Aldrich) under the shaking incubator at 37°C for 1 hour. Cells were filtered by 70 μ m Falcon strainer (Corning) and cultured in α -MEM with 20% FBS, 100 U/mL penicillin, 100 μ g/mL streptomycin, 2 mM L-glutamine. Culture solution was changed every 2 - 3 days, and floating particles were rinsed out. The cells were separated for next passage using trypsin-EDTA (Corning). Subculture until passage 10 was lasting for stemness comparison between cells of early and late passages.

2) Flow cytometry analysis

Passage 3 and passage 10 cells were chosen to compare the expression of cell surface markers of early and late passage. Cells were counted to put same number of cells for each cell surface markers. 1×10^5 of the cells and 1 μ g cell surface markers were leaved in refrigerator at 4°C for 1 hour. The sur-

face markers used in this study were cluster of differentiation (CD) 73, CD 90, CD 34, CD 45 (BD Biosciences) and STRO-1 (R&D system). Inactive cell surface markers were removed and the fluorescent antibodies were used for florescence-activated cell sorting (FACS) analysis. The cell-antibody complexes were analyzed with BD FACSCalibur flow cytometry (BD Biosciences). The cells that were not reacted to any markers were used as control group.

3) Statistical analysis

Data were analyzed using SPSS 23.0 statistical software (SPSS Inc., USA).

III. Results

1. The culture of supernumerary dental pulp stem cells

Dental pulp cells were extracted from SNT of 15 patients; SNT 01 - SNT 15. Dental pulp cells were cultured to passage 10.

2. Immunophenotype

The expression of the cell surface markers in the control group and the experimental group was obtained at passage 3 and passage 10 of human dental pulp cells from SNT 01 and SNT 02 (Fig. 1). CD 73, CD 90, STRO-1 as MSC surface markers were highly expressed in the dental pulp cells from SNT, compared to only a small degree expression of CD 34, CD 45 as hematopoietic and endothelial markers. The expression of STRO-1 at passage 3 of SNT 02 and passage 10 of SNT 01 satisfy the criteria presented by ISCT.

Table 1 and 2 showed the expression of CD markers of the dental pulp cells from SNT at passage 3 and passage 10. CD 73 and CD 90 were positively expressed, while CD 34 and CD 45 were negatively expressed. STRO-1 was expressed. Most of dental pulp cells from SNT were satisfied with the criteria of ISCT.

Table 3 showed the mean value of expression of CD markers at passage 3 and passage 10. At passage 3, dental pulp cells from SNT showed positive expression of CD 73 (94.82%), CD 90 (98.86%) and STRO-1 (20.93%), negative expression of CD 34 (2.25%) and CD 45 (2.52%). At passage 10, dental pulp cells from SNT showed positive expression of CD 73 (96.62%), CD 90 (98.61%) and STRO-1 (35.62%), negative expression of CD 34 (3.86%) and CD 45 (4.14%). The expression of CD 73, CD90

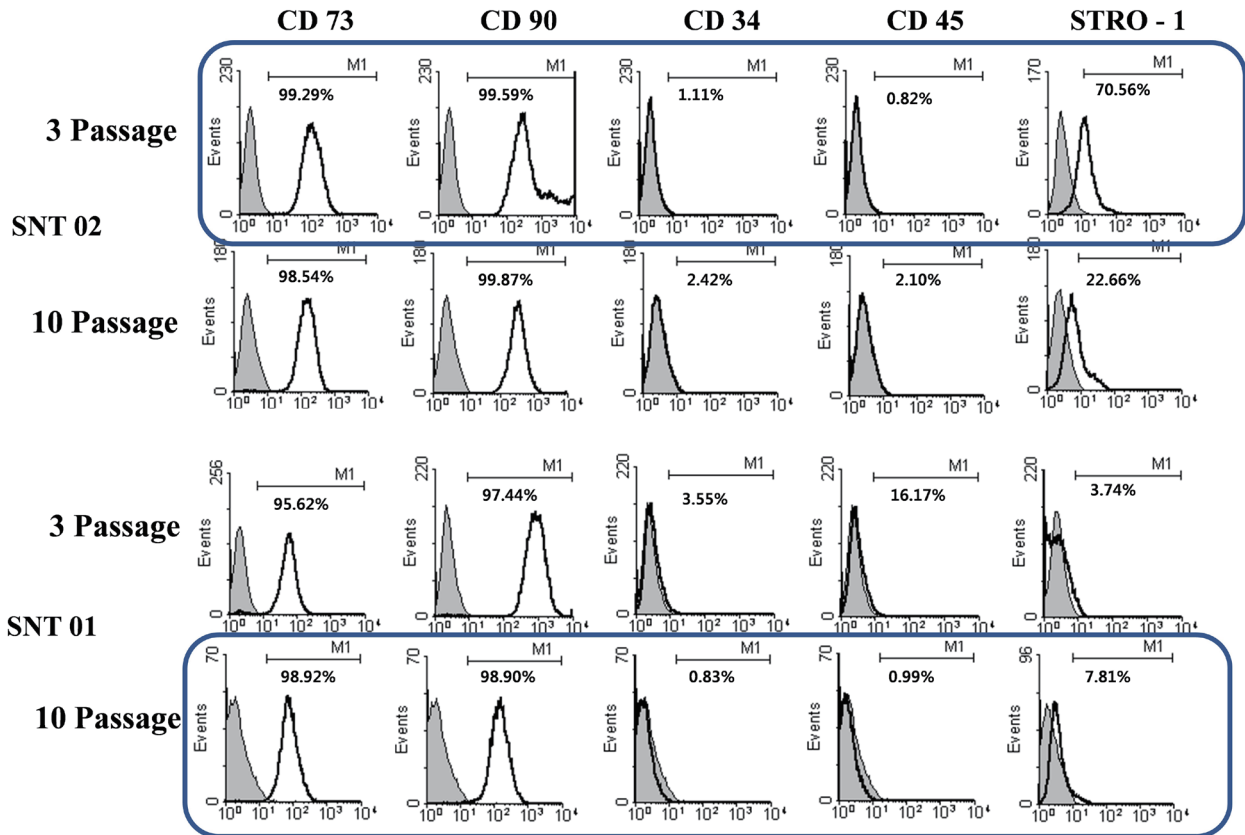


Fig. 1. Expression of cell surface markers in flow cytometry analysis of supernumerary tooth 01 and supernumerary tooth 02 at passage 3 and passage 10.

SNT : Supernumerary tooth, CD : Cluster of differentiation

Table 1. Flow cytometry analysis of the mean percentile values of the expression of CD marker at passage 3

	CD73	CD90	CD34	CD45	STRO-1
SNT01	95.62	97.44	3.55	16.17	3.74
SNT02	99.29	99.59	1.11	0.82	70.56
SNT03	95.83	98.66	0.88	0.91	12.81
SNT04	98.73	99.56	8.34	4.13	6.51
SNT05	99.33	99.41	1.56	1.69	19.77
SNT06	99.26	99.70	1.40	1.07	7.91
SNT07	97.83	99.82	1.54	1.44	13.17
SNT08	99.25	99.48	0.49	0.18	32.76
SNT09	73.21	95.50	2.79	3.02	10.84
SNT10	99.12	99.33	2.19	1.40	16.96
SNT11	99.68	99.76	3.36	2.89	45.40
SNT12	99.59	99.67	1.28	1.11	21.08
SNT13	97.72	98.26	2.33	1.59	29.15
SNT14	95.21	98.26	1.54	1.13	8.55
SNT15	72.63	98.48	1.40	0.25	14.77

Table 2. Flow cytometry analysis of the mean percentile values of the expression of CD marker at passage 10

	CD73	CD90	CD34	CD45	STRO-1
SNT01	98.92	98.90	0.83	0.99	7.81
SNT02	98.54	99.87	2.42	2.10	22.66
SNT03	96.20	99.70	4.89	3.86	47.06
SNT04	98.52	98.77	2.52	2.17	14.59
SNT05	92.41	97.58	1.14	1.11	39.01
SNT06	96.64	96.26	3.41	2.90	16.17
SNT07	85.02	98.72	3.41	2.63	15.26
SNT08	95.90	96.10	6.79	6.27	18.00
SNT09	98.54	99.56	3.71	3.23	11.29
SNT10	97.81	98.52	6.37	5.41	84.66
SNT11	99.69	99.35	6.50	5.00	9.36
SNT12	98.60	98.87	3.45	11.61	74.37
SNT13	95.79	99.49	3.57	2.34	59.61
SNT14	98.81	99.57	3.91	7.37	30.62
SNT15	97.90	97.86	5.03	5.08	83.78

CD : Cluster of differentiation, SNT : Supernumerary tooth

CD : Cluster of differentiation, SNT : Supernumerary tooth

and STRO-1 between two passages was not significantly different. There was a significant difference between two passage in expression of CD 34 and CD 45.

The average values of G-mean were calculated in flow cytometry results (Table 4).

IV. Discussion

The adult stem cells exist as undifferentiated cells in human organs[2,13,14]. Pittenger *et al.*[15] were classified the adult stem cells into MSCs and hematopoietic stem cells, and proved that the adult stem cells could be differentiated into osteoblasts, chondroblasts, myoblasts and nerve cells by controlling culture condition. The other studies reported the adult stem cells could be obtained from various tissues including bone marrow, pancreas, adipose, muscle, blood, hair follicles, skin and dental tissues[16].

In 2006, ISCT presented the minimal 3 criteria of MSCs[9]. First, they have characteristic of adhesion to plastic culture flask. Second, they must show positive expression of CD 105, CD 73 and CD 90, while as well as showing negative expres-

sion of CD 45, CD 34, CD 14 or CD 11b, CD 79alpha or CD 19 and HLA-DR. Thirdly, they should have a possibility of differentiation to osteoblasts, adipocytes and chondroblasts in vitro.

The aim of this study was to identify MSCs in 15 SNT with flow cytometry analysis. CD 73, CD 90 and STRO-1 were used as positive CD markers, and CD 34 and CD45 were used as negative CD markers.

STRO-1 is early mesenchymal stem cell marker, which was detected around blood vessels or nerve fascicles while not detected in size-sieved stem cells[12,17,18]. In 2007, Kolf *et al.*[19] reported STRO-1 was the best CD marker to identify mesenchymal stem cells. Gay *et al.*[20] reported that STRO-1 showed 27% expression in the cells from periodontal ligament of impacted wisdom teeth, and Park *et al.*[21] presented that STRO-1 showed 33.4% expression in the cells from periodontal ligament of supernumerary teeth. In this study, STRO-1 showed 20.93% expression at passage 3 and 35.62% at passage 10. The expression of the STRO-1 was even slightly increased in passage 10 comparing to passage 3. However, there was no significant difference between the two passages.

The expression of positive CD markers was not statistically different between passage 3 and passage 10 in flow cytometry analysis. This meant later passage of dental pulp cells from SNT still contained stemness and could have value of use as a source of stem cells. The expression of negative CD markers was a statistical significance between passage 3 and passage 10. According to ISCT's recommend, the negative CD marker should be detected lower, further study is required.

The dental pulp stem cells (DPSCs) of SNT from 15 donors showed characteristic of MSCs by flow cytometry using CD markers. The characteristic of MSCs was maintained until passage 10. However, FACS results alone are not enough to verify the stem cell function of cells. The differentiation to hard tissues or fat should also be examined. Also additional researches including real time qPCR, would be needed to confirm gene expression in follow-up studies.

Table 3. The mean percentile values of flow cytometry analysis at passage 3 and passage 10

	Passage	Mean	SD
CD 73	3	94.82	9.02
	10	96.62	3.69
CD 90	3	98.86	1.17
	10	98.61	1.18
CD 34	3	2.25 ^{*a}	1.90
	10	3.86 ^{*a}	1.80
CD 45	3	2.52 ^{*b}	3.92
	10	4.14 ^{*b}	2.79
STRO-1	3	20.93	17.69
	10	35.62	27.69

Mann-Whitney test (* : $p < 0.05$), MSC : Mesenchymal stem cell, CD : Cluster of differentiation, SD : Standard deviation

Table 4. Average G-mean values of flow cytometry analysis at passage 3 and passage 10

	CD 73		CD 90		CD 34		CD 45		STRO-1	
Passage	3	10	3	10	3	10	3	10	3	10
G-Mean	114.95	144.82	338.40	493.86	0.11	3.22	-0.01	3.29	3.15	9.23
SD	81.76	60.15	214.38	475.05	0.41	0.60	0.37	0.85	3.00	5.88

CD : Cluster of differentiation, SD : Standard deviation

V. Conclusions

The result indicated that DPSCs of impacted SNT had the characteristics of MSCs. Comparing passage 3 and passage 10 in flow cytometry analysis, the cell marker expressions were similar. There was no significantly difference between the positive CD marker values of passage 3 and passage 10. There was a significant difference in the negative CD marker values between the two passages. DPSCs maintained stemness until the late passage. It is confirmed that the possibility of using DPSCs of impacted SNT from young children as a donor of stem cells.

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국문초록

유세포 분석을 통한 과잉치 치수 유래 세포의 줄기세포 특성 연구

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이 연구의 목적은 발치한 매복 과잉치에서 얻은 치수 유래 세포의 줄기세포 특성을 유세포 분석을 통해 알아보는 것이다. 15명의 환자로부터 채취한 정중 과잉치의 치수 세포를 계대 배양하였고, 3계대와 10계대 세포들을 유세포 분석을 이용하여 분석하였다. 간엽성 줄기세포 표지자 관찰에 사용된 항체 표지자(CD)는 양성 표지자로는 CD 73, CD 90, 그리고STRO-1 와 음성 표지자로는 CD 34, CD 45 이었다. 3계대에서 CD 73, CD 90은 각각 94.82%, 98.86%의 양성반응을, CD 34, CD 45는 각각 2.25%, 2.52%로 음성 반응을 보였으며, STRO-1은 20.93%를 나타냈다. 10계대에서는 CD 73, CD 90은 각각 96.62%, 98.61%의 발현을 보였지만, CD 34, CD 45는 각각 3.86%, 4.14%를 나타냈다. STRO-1은 35.62%로 발현되었다. 이상의 결과에서 과잉치 치수 유래 세포는 간엽성 줄기세포의 특성을 가지며, 3계대와 10계대 모두에서 간엽성 줄기세포의 특성을 유지하고 있다고 사료된다. 이에 빠른 성장 속도와 늦은 계대까지 유지되는 줄기 세포능을 고려할 때, 치아 유래 줄기세포의 공여부로서 매복 과잉치의 충분한 활용 가능성을 확인하였다.