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박지현의 석사 학위논문을 인준함

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2011년 6월 일

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2. Colony forming unit - fibroblast (CFU - F)

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Polymerase Chain Reaction, RT - PCR)

4. (Flow Cytometry Analysis)

5. (Adipogenic Differentiation)

6. (Osteogenic
Differentiation)

7. (Quantitative Reverse
Transcription Polymerase Chain Reaction, qRT - PCR)
8. (Statistical analysis)
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3. RT - PCR
- 4.
5. (Adipogenic Differentiation)
6. (Osteogenic
Differentiation)
- .
- .

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Table 1. RT - PCR and qRT - PCR primers used in this
study6

ALP : alkaline phosphatase

- MEM : - minimum essential medium

BMMSCs : bone marrow mesenchymal stem cells

BSP : bone sialoprotein

CFU - F : colony forming unit - fibroblast

EMSCs : ectomesenchymal stem cells

ES cells: embryonic stem cells

FBS : fetal bovine serum

GAPDH : glyceraldehyde - 3 - phosphate dehydrogenase

LPL : lipoprotein lipase

PCR : polymerase chain reaction

PPAR α : peroxisome - proliferator - activated receptor α

qRT - PCR : quantitative reverse transcription polymerase
chain reaction

RT - PCR : reverse transcription - polymerase chain reaction

sPDLSCs : supernumerary periodontal ligament stem cells

가

가

colony forming unit - fibroblast (CFU - F)

(Oct - 4, Nanog, Nestin, Stro - 1, CD146)

(5.1%)

(6.0%)가

34.1%, 84.8%)

Oct - 4, Nanog, Nestin Stro - 1 CD146 (

, peroxisome - proliferator - activated receptor 2
(PPAR 2) lipoprotein lipase (LPL) 가

alkaline phosphatase (ALP)
bone sialoprotein (BSP) 가 , ALP

가

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1, 2,

3, 4 .

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5-8 ,

9-

14 ,

15 ,

16, 17

가

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, , Malassez , ,
 11 ,
 가
 14, 18 .
 가 ,
 19 9, 12 , 20
 .
 가
 1.0~3.5%²¹ . ,
 , ,
 22 .
 가
 (source) 가
 가 .
 가 23, 24 .
 가 .

1.

6 - 10

6

가

(2 - 2010 - 0015)

explants culture ²⁵

(supernumerary periodontal

ligament stem cells, sPDLSCs)

15

, 5

(bone marrow mesenchymal

stem cells, BMMSCs)

- minimum

essential medium (- MEM; Gibco BRL, Grand Island, NY, USA)

10% fetal bovine serum (FBS; Gibco BRL), 100 U/ml of penicillin and

100 µg/ml of streptomycin (Gibco BRL), 2 mM L - glutamine (Gibco

BRL), 10 mM L - ascorbic acid (Sigma, St. Louis, MO, USA) 가

37 ° C

5% CO2

2

2. Colony forming unit - fibroblast (CFU - F)

가 (mesenchymal stem cells, MSCs) (colony)
 , colony forming unit - fibroblast (CFU - F)
 . 6 - well (BD Falcon, Lincoln Park, NJ, USA) 480 cells/well 10
 . 10% natural - buffered formalin (Sigma) 4
 20 , 0.5% crystal violet (Sigma, St. Louis, MO, USA)
 5 . 2mm CFU -
 F 가 CFU - F

3. (Reverse Transcription - Polymerase Chain Reaction, RT - PCR)

RT - PCR
 (embryonic stem cells, ES cells) Oct - 4
 Nanog (ectomesenchymal stem cells, EMSCs)
 Nestin mRNA . Total RNA

RNeasy mini kit (Qiagen, Valencia, CA, USA)
 . Total RNA Nanodrop ND - 1000[®] (Thermo
 Scientific, Waltham, MA, USA) . cDNA
 total RNA 1 μ g oligo (dT)15 primer
 Maxime RT premix kit (Intron biotechnology, Seoul, Korea)
 20 μ l , 45 1 , 95 5
 . (polymerase chain reaction, PCR)
 Maxime PCR PreMix Kit (i - StarTaq) (Intron biotechnology)
 . cDNA 1 μ l, 10 pmol/ μ l forward reverse
 primer 1 μ l 가 20 μ l가
 . Denaturation 95 2 , 95 20
 가 , annealing 60 10 , extension 72 20 , final
 extension 72 5 PCR (Swift[™] MaxPro Thermal Cyclers;
 ESCO, Singapore) . PCR
 human glyceraldehyde - 3 - phosphate dehydrogenase (GAPDH)
 , primer Table 1
 . PCR 6 \times LoadingStar (DyneBio, Sungnam,
 Korea) 1.5% agarose gel . ChemiDoc XRS
 (BIO - RAD Lab, Richmond, CA, USA) UV light

Table 1. RT - PCR and qRT - PCR primers used in this study

Genes	Primer sequence (5' - 3')	Size (bp)	Cycles	Gene Bank Accession No.	Ref
Oct - 4	F: CGACCATCTGCCGCTTTGAG R: CCCCTGTCCCCATTCCTA	573	30	NM_002701.4	26
Nanog	F: TGCAAATGTCTTCTGCTGAGAT R: GTTCAGGATGTTGGAGAGTTC	287	32	NM_024865.2	26
Nestin	F: GCCCTGACCACTCCAGTTTA R: GGAGTCCTGGATTTCTTCC	200	33	NM_006617.1	27
GAPDH	F: AGGTGAAGGTCGGAGTCAACG R: GCTCCTGGAAGATGGTGATGG	231	25	NM_002046.3	28
PPAR 2	F: ACAGCAAACCCCTATTCCATGCTGT R: TCCCAAAGTTGGTGGGCCAGAA	159	45*	NM_015869.4	25
LPL	F: TGGACTGGCTGTCACGGGCT R: GCCAGCAGCATGGGCTCCAA	167	45*	NM_000237.2	25
ALP	F: GGACCATTCCACGTCTTCAC R: CCTTGTAGCCAGGCCATTG	137	45*	NM_000478.4	26
BSP	F: CTGGCACAGGGTATACAGGGTTAG R: ACTGGTGCCGTTTATGCCTTG	182	45*	NM_004967.3	29
GAPDH	F: TCCTGCACCACCACTGCTT R: TGGCAGTGATGGCATGGAC	100	45*	NM_002046.3	29

* qRT - PCR

4. (Flow Cytometry Analysis)

Stro - 1 CD146 .
(phosphate buffered saline, PBS, pH=7.2)
cell dissociation buffer (Invitrogen, Carlsbad, CA,
USA) dish 1×10^6 cells/ml flow cytometry
staining buffer (eBioscience, San Diego, CA, USA) . Stro -
1 $5 \mu\text{g}/1 \times 10^6$ cells
monoclonal anti - human Stro - 1 (IgM; R&D system, Minneapolis, MN,
USA) 4°C 1 . flow cytometry
staining buffer R - phycoerythrin (R -
PE) goat anti - mouse IgM (SouthernBiotech, Birmingham, AL,
USA) $0.1 \mu\text{g}/1 \times 10^6$ cells 4°C 30 .
CD146 , fluorescein isothiocyanate (FITC)가
anti - human CD146 (eBioscience) $20 \mu\text{l}/1 \times 10^6$ cells
 4°C 1 . Stro - 1
가 . FACSCalibur™ Flow
Cytometer (BD Biosciences) FCSExpress V3 software (De Novo
Software, Thornhill, ON, Canada)

5. (Adipogenic Differentiation)

3 -

5 12 - well 1×10^4 cells/cm²

가 .

10 - MEM 10% FBS, 1% antibiotics, 1 μ M dexamethasone (Sigma), 10 μ g/ml human insulin (Sigma), 100 μ M indomethacin (Sigma), 500 μ M 3 - isobutyl - l - methylxanthine (IBMX; Sigma) 가 , 10 (-

MEM 10% FBS 1% antibiotics, 10 μ g/ml human insulin 가) 10 .

Oil red O

. 10% natural - buffered formalin

4 ° C 30 0.2% Oil Red O (Sigma)

10 ,

. peroxisome proliferator - activated receptor 2 (PPAR 2) lipoprotein lipase (LPL) mRNA (Quantitative Reverse Transcription - Polymerase Chain Reaction, qRT - PCR) .

6. (Osteogenic Differentiation)

3 -

5 12 well 1×10^4 cells/cm²

가 MEM 10%
 FBS 1% antibiotics, 0.1 μM dexamethasone, 2 mM -
 glycerolphosphate (Sigma) 50 μM ascorbic acid 2-phosphate
 (Sigma) 가 5
 Alizarin Red S
 10% natural-buffered
 formalin 2% Alizarin Red S (pH=4.2; Sigma) 5
 alkaline phosphatase (ALP) bone
 sialoprotein (BSP) mRNA

**7. (Quantitative Reverse Transcription
 Polymerase Chain Reaction, qRT - PCR)**

Total RNA cDNA RT - PCR
 7 μl, forward reverse primer (10
 pmol/μl) 1 μl, cDNA 1 μl 2x SYBR Premix Ex Taq™
 (Takara Bio, Otsu, Japan) 10 μl 20 μl Thermal
 Cycler Dice™ real time system (Takara Bio)
 denaturation 95 °C 10
 denaturation 95 °C 5 , annealing 60 °C 15 ,
 extension 72 °C 10 45 60 °C 95 °C
 dissociation primer Table 1

primer dissociation curve peak가 band가
 mRNA GAPDH
 2^{-CT}

8. (Statistical analysis)

CFU - Fs qRT - PCR
 mRNA SPSS (version 17.0; Chicago, IL, USA)
 Mann - Whitney U test ($p < 0.05$)

1.

(Fig 1).

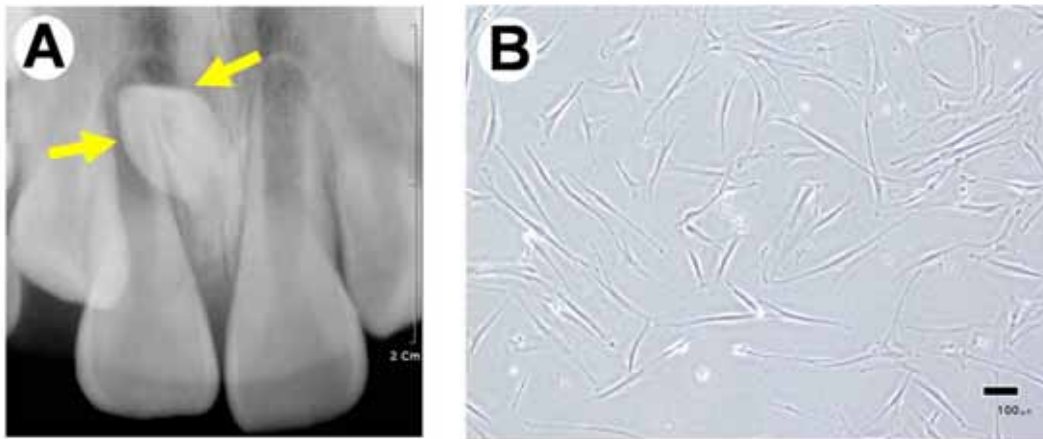


Figure 1. (A) Radiographic view of a supernumerary tooth (arrows). (B) Morphology of stem cells obtained from the periodontal ligament of a supernumerary tooth (sPDLSCs). Scale bar: 100 μ m.

2. Colony forming unit - fibroblast (CFU - F)

	10	CFU -
F		6.0%
	5.1%	.

(Fig 2).

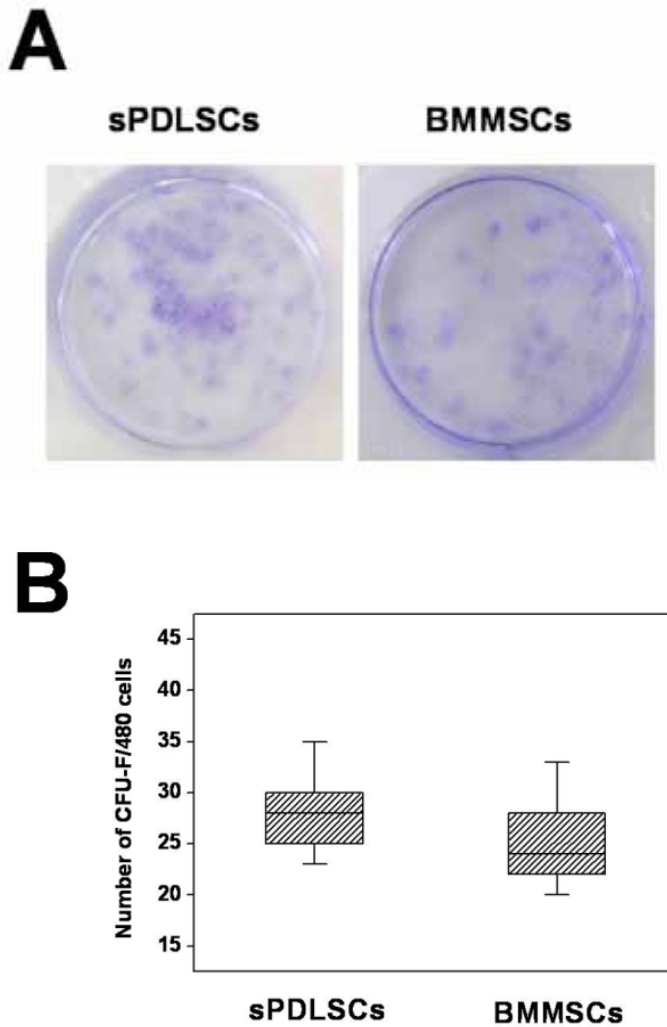


Figure 2. Colony forming unit–fibroblast (CFU - F) assay for the sPDLSCs and BMMSCs. (A) Crystal violet staining. (B) The numbers of colonies per 480 cells after 10 days of culture did not differ significantly between the

two cell types (Mann - Whitney U test: $p > 0.05$; $n = 8$). Data are presented as box and whisker plots showing median, quartiles, and range.

3. (Reverse Transcription - Polymerase Chain Reaction, RT - PCR)

PCR, ES cells, mRNA RT - Oct - 4
Nanog, EMSCs, Nestin (Fig 3A).

4. (Flow Cytometry Analysis)

Stro - 1 CD146
Stro - 1 CD146 (Fig 3B).

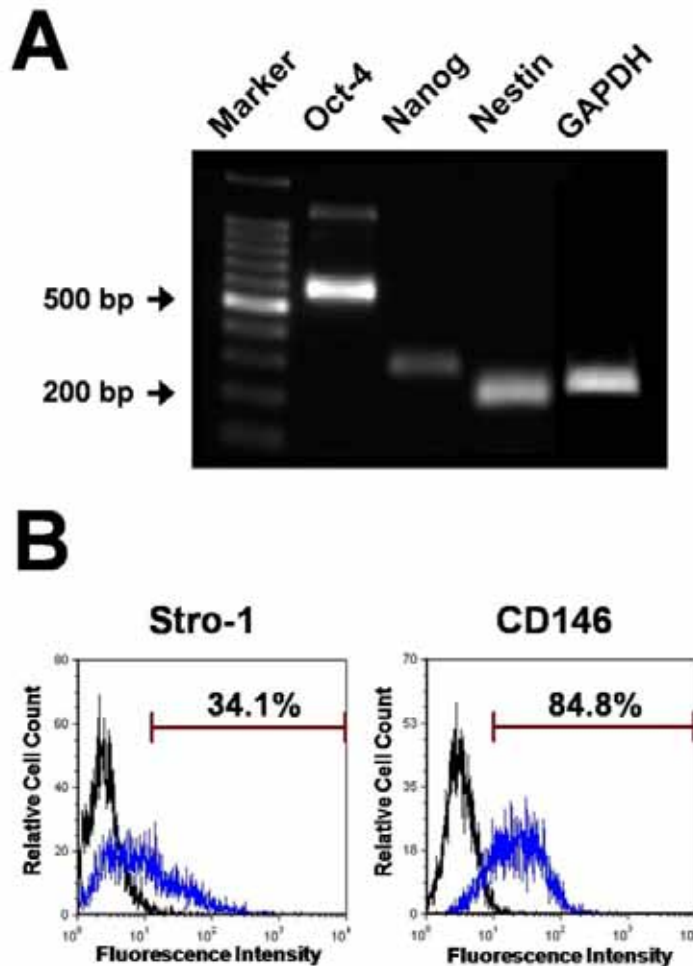
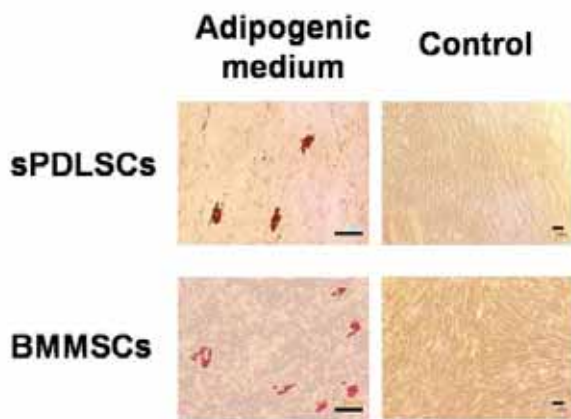


Figure 3. (A) Expression of genes associated with embryonic stem cells (Oct - 4 and Nanog) and ectomesenchymal stem cells (Nestin) in sPDLSCs. GAPDH was used as an internal control. (B) Flow cytometry analysis of mesenchymal stem - cell markers (Stro - 1 and CD146) in the sPDLSCs. Horizontal bars indicate 1% of control samples.

5. (Adipogenic Differentiation)

Oil red
(Fig 4A),
lipid vacuole
qRT - PCR PPAR 2 LPL
가가 (Fig 4B).
Oil red O
가가
PPAR 2 LPL

A



B

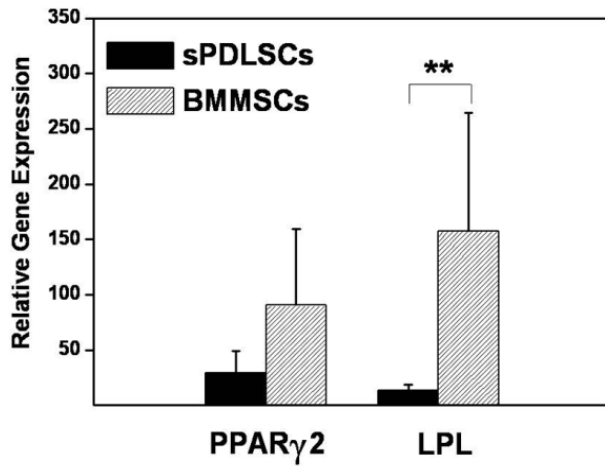


Figure 4. Adipogenic differentiation of the sPDLSCs and BMMSCs. (A) Oil Red O staining after 20 days of culture in adipogenic differentiation medium and control medium. Scale bar: 100 μ m. (B) Quantitative RT - PCR analysis revealed up - regulation of peroxisome - proliferator - activated receptor 2 (PPAR 2) and lipoprotein lipase (LPL) gene expressions at 20 days of culture under adipogenic stimuli. ** p<0.01 (Mann - Whitney U test, n=3).

6. (Osteogenic Differentiation)

Alizarin red S

가

(Fig 5A). qRT - PCR

ALP BSP 가 , ALP 1 -

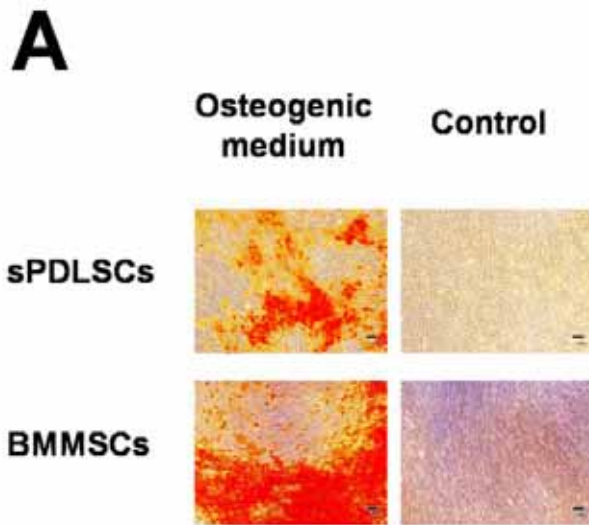
2 가 가 BSP

가 가 .

ALP 가가

BSP 가 (Fig

5B).



B

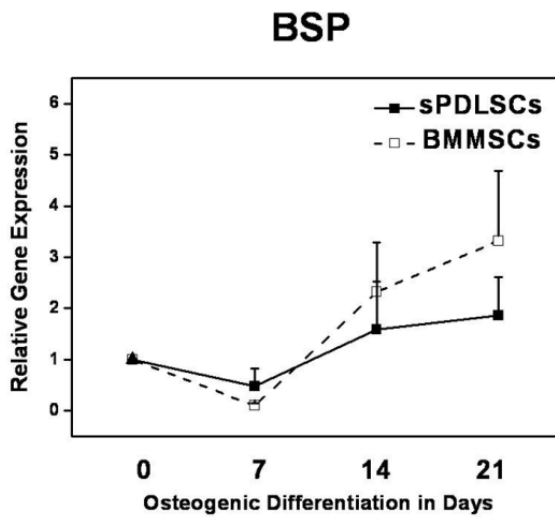
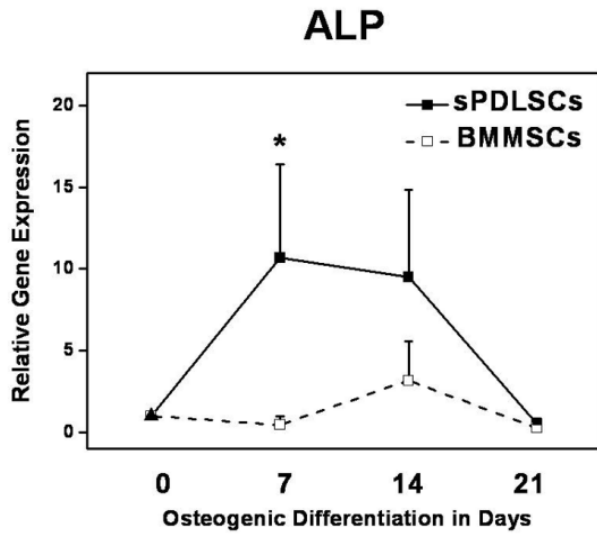


Figure 5. Osteogenic differentiation of the sPDLSCs and BMMSCs. (A) Alizarin Red S staining after 5 weeks of culture in osteogenic

differentiation medium and control medium. Scale bar: 100 μm . (B) Quantitative RT-PCR analysis revealed the changes in alkaline phosphatase (ALP) and bone sialoprotein (BSP) gene expression at 0, 7, 14, and 21 days of culture under osteogenic stimuli. * $p < 0.05$ (Mann-Whitney U test, $n=3$).

가

가

6.0%

Nagatomo ¹² 1% Tanaka ³⁰ 15~35%

가

31, 32

(pluripotent stem cells) 가

(ES cells) (pluripotency)

(Parkinson's disease), (Alzheimer's disease),

11. Oct4 Nanog

(pluripotency)

33, 34, 가

가

Oct - 4

Nanog

STRO - 1 CD146

STRO - 1

CD146

^{25, 35} Nagatomo ¹²

STRO - 1,

CD105, CD166

Xu ¹⁸

STRO - 1 CD146

가 가 2.6%

Gay ⁹ 3

27% STRO - 1 가

STRO - 1 가 34.1%

가

가 가

가

(multipotentiality)

^{13, 36}

가

가

Oil red O

37

PPAR 2

38

LPL

가

,
9, 13, 14

Song 25

가

(Alizarin Red S)

ALP

가

39

가

가

9

PPAR 2,

Wnt , hedgehogs, Bone morphogenic proteins(BMPs)

IGF,

FGF

40 - 44

가

가

.

가

가

.

,
 가 ,
 Oct - 4 Nanog
 Nestin STRO - 1
 CD146 ,
 가 가 가
 ,
 가
 가 (apical papilla) (dental follicle)
 가 가

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Abstract

Characteristics of stem cells derived from the periodontal ligament
of supernumerary teeth

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(Directed by Professor Jae-ho Lee)

Human teeth have been identified as a new source of postnatal stem cells that have the capacities of self - renewal and multilineage differentiation. In this study we compared the characteristics of stem cells obtained from the periodontal ligament of supernumerary teeth (sPDLSCs) with those of bone - marrow - derived mesenchymal stem cells (BMMSCs), with the aim of extending the sources of stem cells.

We performed a colony forming unit–fibroblast (CFU - F) assay to evaluate the self - renewal ability of the cells. Reverse transcription - polymerase chain reaction (RT - PCR) and flow cytometry analyses were used to detect the expressions of various stem - cell markers: Oct - 4, Nanog, Nestin, Stro - 1, and CD146. The abilities of adipogenic and osteogenic differentiation were monitored by histochemical staining and quantitative RT - PCR.

The colony - forming efficiency was slightly higher for sPDLSCs than for BMMSCs (6.0% vs. 5.1%, respectively). The sPDLSCs expressed mRNA of Oct - 4, Nanog, and Nestin, and presented surface proteins of Stro - 1 and CD146 (34.1% and 84.8%, respectively). The sPDLSCs could differentiate to adipogenic lineage; however, the up - regulation of peroxisome - proliferator - activated receptor 2 (PPAR 2) and lipoprotein lipase (LPL) gene expressions was less than for the BMMSCs. Osteogenic differentiation of the sPDLSCs was confirmed by up - regulation of alkaline phosphatase (ALP) and bone sialoprotein (BSP) gene expressions, but elevation of ALP gene expression was observed earlier than for the BMMSCs.

We have shown that sPDLSCs exhibit stem - cell properties such as self - renewal ability, expression of stem - cell markers, and differentiation into adipogenic and osteogenic lineages. This cell type could represent a good stem - cell candidate for use in regenerative medicine.

Key Words : supernumerary teeth, periodontal ligament, stem cells, differentiation, regenerative medicine