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The role of the Wnt/ β -catenin signaling pathway in the proliferation of gold nanoparticle-treated human periodontal ligament stem cells

Chen Li¹, Zhuoquan Li², Yan Zhang³, Abdel Hamid Fathy⁴ and Min Zhou^{5*}

Abstract

Background: Several studies have confirmed that gold nanoparticles (AuNPs) of specific concentration and size exert a boosting effect on cell proliferation; however, the mechanism through which this effect occurs remains unknown. This study explores the canonical Wnt signaling pathway in AuNP promotion of human periodontal ligament stem cell (hPDLSC) proliferation.

Methods: MTS was employed to evaluate hPDLSC proliferation. The interference of *LRP5* and β -catenin was steered by shRNA plasmids and siRNA, respectively, at which point the expression of *MYC*, *CCND1*, *AXIN2*, and *POU5F1* had been estimated via real-time PCR. The expressions of *LRP5* and β -catenin were detected via western blot assay.

Results: The proliferation of hPDLSCs treated with 60 nm AuNPs at 56 μ M was clearly elevated. In contrast, β -catenin siRNA significantly decreased cell viability. The *LRP5* shRNA plasmid did not consistently impact cells. The expressions of these four genes downstream of the Wnt/ β -catenin signaling pathway were not significantly overexpressed in response to the interference of shRNA plasmid/siRNA with the treatment of AuNPs.

Conclusions: These results suggest that the Wnt/ β -catenin signaling pathway plays a significant role in the process of AuNP promotion of hPDLSC proliferation.

Keywords: Wnt/ β -catenin signaling pathway, Human periodontal ligament stem cell proliferation, Gold nanoparticle, shRNA, siRNA

Background

As a result of the rapid development of tissue engineering, proliferation of human periodontal ligament stem cells (hPDLSCs) [1] as seed cells both in vitro and in vivo is a newly emerging technique. Gold nanoparticles (AuNPs) have been used for many scientific applications during the last two decades. Their specific chemical and physical-dependent properties highlight them as very promising agents in many fields such as drug delivery, targeting, imaging [2], and even vaccination [3]. Recently, multiple ex-vivo studies have shown that a

specific size and concentration of AuNPs have a promoting effect on cell proliferation [4, 5], and it has been confirmed that the cytotoxicity of AuNPs is a size-dependent, concentration-dependent, and time-dependent effect [6–9]. Consequently, gold nanoparticles have an extensive application prospect in tissue engineering.

Our previous experiments confirmed that 60 nm AuNPs at a concentration of 56 μ M significantly promoted the proliferation of hPDLSCs, while exerting no apparent side effects on the differentiation potential of cells. However, another study reported that AuNPs mainly promote the differentiation of human adipose-derived mesenchymal stem cells (MSCs) in osteogenesis via various signaling pathways [10]. Moreover, AuNPs of different sizes can affect the physical processes of cells via diverse signaling

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pathways. However, the mechanism by which AuNPs promote the proliferation or differentiation of MSCs remains unclear. Therefore, the molecular mechanism by which AuNPs promote MSC proliferation remains to be explored.

Wnt/ β -catenin signaling is crucial for regulating embryonic development and cell proliferation as well as for the determination of the cell fate [11]. Proliferation and growth of stem cells are premises of the mass differentiation of cells needed to form tissues and ultimately organisms. This process is established on canonical Wnt signaling, which increases nuclear and cytoplasmic β -catenin and initiates transcriptional activators such as *MYC* and *CCND1* [12]. Recently, Wnt signaling has been implicated in the control of MSC differentiation, including osteogenic, chondrogenic, adipogenic, and myogenic differentiation [10, 13].

Here, an in-vitro study was conducted to investigate the role of the Wnt/ β -catenin signaling pathway for the proliferation of hPDLSCs treated with AuNPs (60 nm at 56 μ M). LRP5 and β -catenin were interfered with by shRNA and siRNA, respectively. Changes in the expression of Wnt-related genes (*MYC*, *CCND1*, *AXIN2*, and *POU5F1*) were evaluated via real-time PCR. The results indicate that AuNPs promote the proliferation of hPDLSCs, which is likely a result of the Wnt/ β -catenin signaling pathway.

Methods

hPDLSC culture

Premolars from teenagers 12 and 13 years old, which had been extracted for orthodontic reasons, were freshly obtained from the Oral Surgery Department, Hospital of Stomatology, Tongji University with informed consent. The teeth were rinsed three times with PBS complemented with Antibiotic–Antimycotic solution. The periodontal ligament tissue in the middle third of the root was carefully removed with a surgical scalpel. The tissue was minced and enzymatically digested using type I collagenase (1%; Gibco) and Dispase II (2.5%; Roche) for 1 h at 37 °C. A suspended cell solution was passed through a 70-mm cell strainer to prepare a single cell suspension. The cells were centrifuged and rinsed with PBS and the obtained cell pellet was resuspended and cultured in alpha-modified minimum essential medium (α -MEM; Gibco) with 10% FBS (Gibco) at 37 °C in 5% CO₂ in humidified air, until the cells proliferated and reached confluence. Human periodontal ligament cells (hPDLSCs) were supplemented with an immunofluorescence identification using vimentin and cytokeratin. STRO-1⁺ hPDLSCs were sorted via flow cytometry (FCM) and preliminary identification of the stem cell characteristics was achieved via a plate clone formation assay, multidirectional induction as osteoblasts, and adipocytic differentiation.

Characterization of 60 nm AuNPs

The 60 nm AuNPs were acquired from British Biocell International Company (BBI, UK). The final concentration of AuNPs was adjusted with α -MEM medium. The AuNPs were characterized using both a UV–Vis spectrophotometer (Cary 50; Varian) and a Zetasizer (Nano-ZS90; Malvern).

shRNA plasmid and siRNA interference

For both shRNA plasmid and siRNA interference, cells were seeded at a density of 2×10^6 cells per well in six-well culture plates and were cultured until 50–70% confluence. Cells were transfected according to an established shRNA transfection protocol [14]. After incubation for 6 h at 37 °C in 5% CO₂ in humidified air, cells were cultured overnight with 1 ml α -MEM (Gibco) containing 20% FBS and 1% Antibiotic–Antimycotic. The medium was carefully removed and cells were washed with PBS. After culture for 3 h with AuNPs (60 nm at 56 μ M), hPDLSCs were washed with PBS. The culture medium was replaced by α -MEM, and the cells were cultured for 48 h.

Cell viability assay

To measure cell viability, hPDLSCs were interfered with by either shRNA plasmid or siRNA, and were then plated in triplicate in 96-well tissue culture-treated plates at a density of 1000 cells per well and incubated overnight. The medium was carefully removed and cells were washed with PBS. The 60 nm AuNPs were diluted with fresh serum-free α -MEM medium at a concentration of 56 μ M and were added to the hPDLSCs. After 2 h, the medium was aspirated and replaced with full medium. Cells without AuNPs were used as control; wells without AuNPs and cells were used as blanks for spectrophotometric readings. MTS (20 μ l; Promega) was added to each well at 0, 24, 48, 72, and 96 h. After incubation at 37 °C for 2 h, the optical density (OD) value for each well was measured spectrophotometrically at 490 nm.

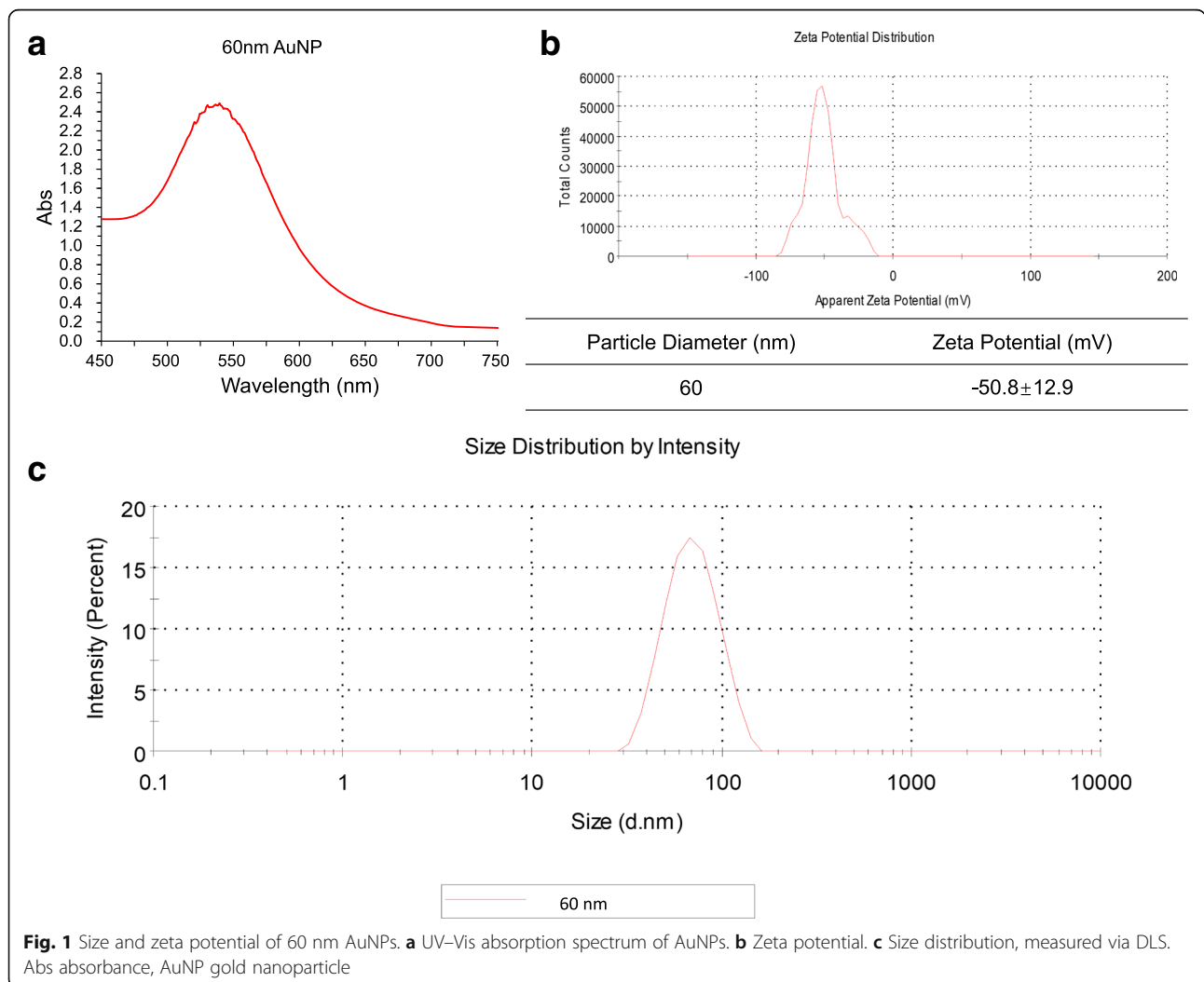
The viability assay (Promega) was conducted according to the manufacturer's instructions. Results were reported as percentage relative fold changes for each group and were internally normalized to the OD reading at 0 h.

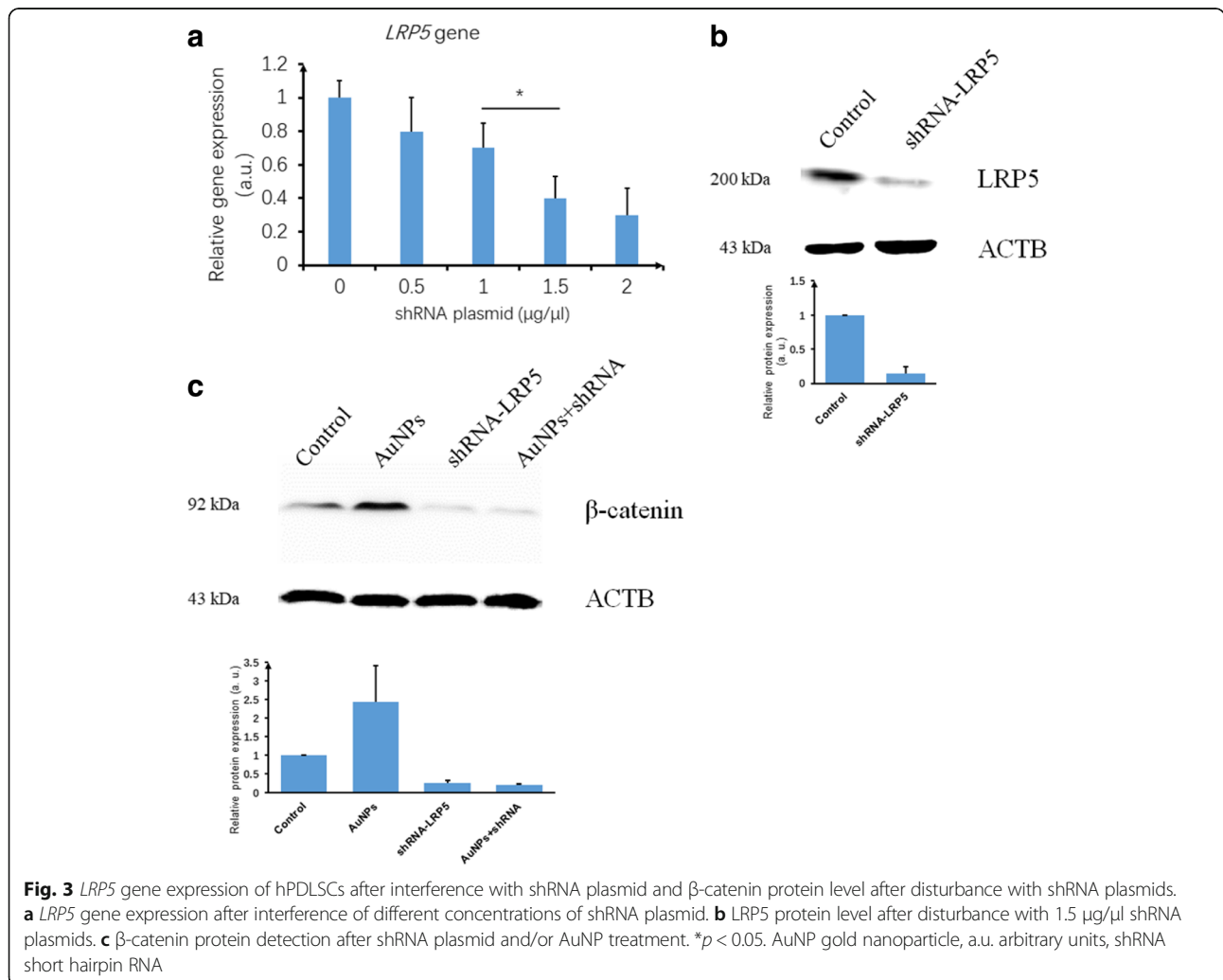
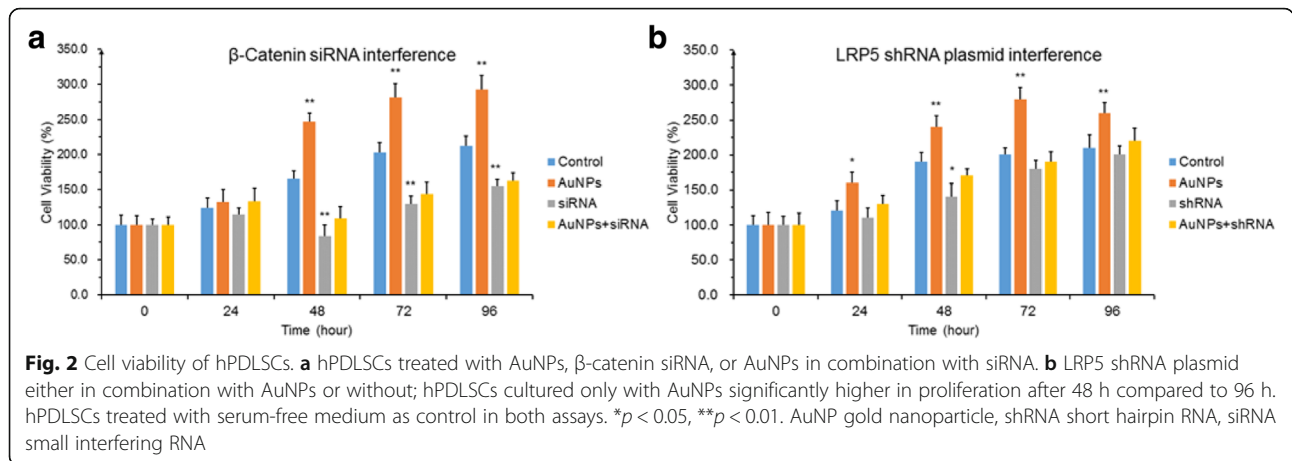
Real-time PCR analysis

After inducing interference with siRNA or shRNA (with or without AuNPs), hPDLSCs were washed with PBS. Total RNA was isolated using Trizol reagent following the manufacturer's instructions and the RNA concentration was measured spectrophotometrically (Eppendorf Biophotometer, Germany). First-strand cDNA was then synthesized using a PrimeScript First Strand cDNA Synthesis Kit (Takara, China). PCR was performed using 1 μ l of the RT products as a template. For

Table 1 Primers for quantitative real-time polymerase chain reaction analysis

Gene	Product size (base pairs)	Sequences
MYC	189	Forward: 5'-TCCTTTATCACACCCCCTACC-3' Reverse: 5'-TTAGCCAAATTGTTCCCTGC-3'
CCND1	94	Forward: 5'-CCGAGAAGCTGTGCATCTACAC-3' Reverse: 5'-AGGTTCCACTTGAGCTTGTTTAC-3'
AXIN2	112	Forward: 5'-GGCTCCAGAAGATCACAAG-3' Reverse: 5'-TATGGAATTTCTCCACACA-3'
POU5F1	115	Forward: 5'-AGCAAAACCCGGAGGAGT-3' Reverse: 5'-CCACATCGGCCTGTGTATATC-3'
LRP5	138	Forward: 5'-GACCAGCCCTTTGTTTTGAC-3' Reverse: 5'-TGTGGACGTTGATGGTATTGGT-3'
β-catenin	77	Forward: 5'-AAGTGGTGGTATAGAGGCTCTTG-3' Reverse: 5'-GATGGCAGGCTCAGTGATGC-3'
ACTB	108	Forward: 5'-TTGCTGACAGGATGCAGAAG-3' Reverse: 5'-TAGAGCCACCAATCCACACA-3'





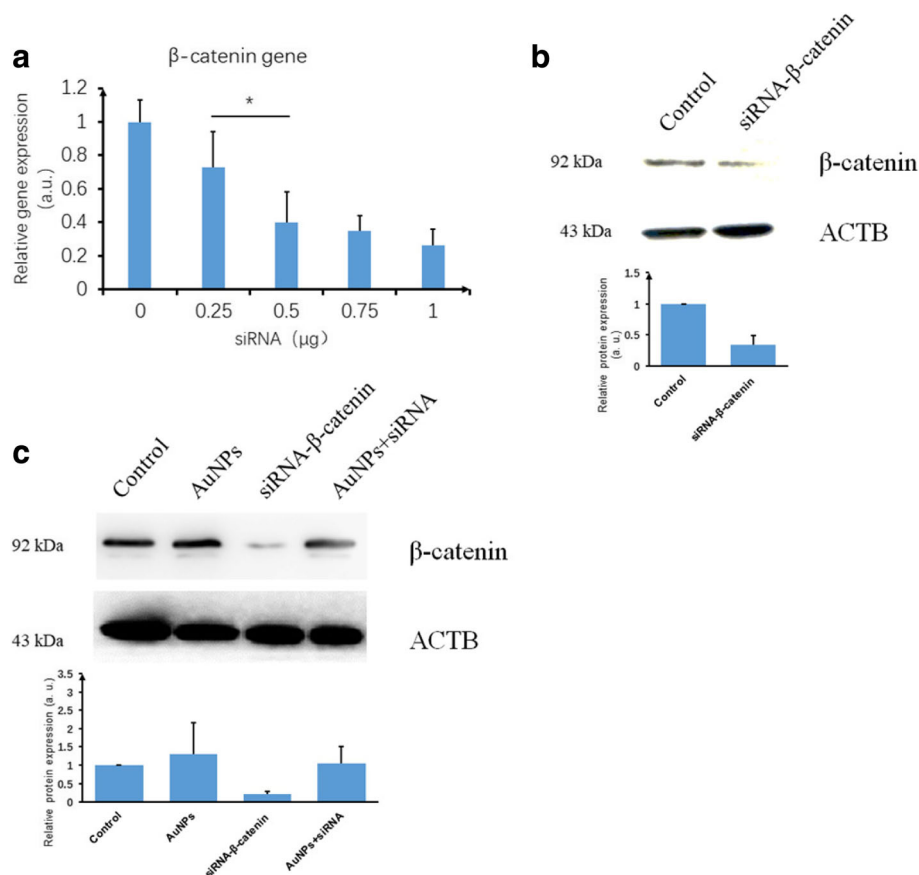


Fig. 4 β -catenin gene expression after interference with siRNA. **a** Different concentrations of siRNA interfere with β -catenin gene expression in hPDLSCs. **b** β -catenin protein expression after disturbance with 0.5 μ g/ μ l siRNA. **c** β -catenin protein detection after siRNA and/or AuNP treatment. * $p < 0.05$. AuNP gold nanoparticle, a.u. arbitrary units, siRNA small interfering RNA

the amplification step, 40 cycles were conducted under the following conditions: denaturation at 94 °C for 2 min, annealing at 58 °C for 30 s, and extension at 72 °C for 45 s. The specific gene primers used for real-time PCR are presented in Table 1. Real-time PCR was used to measure the expression of the genes of Wnt (*MYC*, *CCND1*, *AXIN2*, *POU5F1*, *LRP5*) and β -catenin. All target gene expression results have been normalized to *ACTB*.

Western blot analysis

Cells were washed with ice-cold PBS twice and lysed in high-salt lysis buffer (45 mM HEPES (pH 7.5), 400 mM NaCl, 1 mM EDTA, 10% Glycerol, 0.5% NP40, 6.25 mM NaF, 20 mM β -glycerophosphate, 1 mM DTT, 20 mM sodium butyrate, and 1 \times protease inhibitor cocktail; Roche). Equal amounts of protein extracts were resolved via 12% SDS-PAGE, and then transferred to a polyvinylidene difluoride membrane (BIO-RAD). The membranes were blocked with 5% (wt/vol) dry milk in PBS-Tween-20 (0.5% vol/vol) and probed with primary antibodies overnight. Anti- β -catenin (#9562) and anti-LRP5 (#5731)

antibodies were purchased from Cell Signaling Technology (CST, USA). Secondary antibodies were conjugated to horseradish peroxidase (#7074; CST, USA) for 1 h at room temperature. The immunoreactivity was detected with an ECL kit (#6883; CST). Anti-*ACTB* antibodies (sc-47,778; Santa Cruz Biotechnology) were used as loading control. The quantification of western blot analysis was ushered by ImageJ software.

Statistical analysis

Data were collected from three separate experiments and are shown as the mean \pm standard deviation (SD). Statistical differences were analyzed with one-way analysis of variance (ANOVA) using SPSS 17.0. $p < 0.05$ was considered statistically significant.

Results

Characterization of 60 nm AuNPs

The 60 nm AuNPs peaked at 538 ± 14 nm (Fig. 1a). The Zetasizer showed that the size of AuNPs was 58.71 ± 22.33 nm (Fig. 1c) and the zeta potential was -50.8 ± 12.9 mV (Fig. 1b).

Cell viability of hPDLSCs treated with β -catenin siRNA or LRP5 shRNA plasmid and cultured with AuNPs

To evaluate the proliferation of hPDLSCs in the cell viability assay, hPDLSCs were cultured with AuNPs, β -catenin siRNA, or AuNPs and β -catenin siRNA (Fig. 2a), or with LRP5 shRNA plasmid or AuNPs and LRP5 shRNA plasmid (Fig. 2b). hPDLSCs treated by AuNPs profoundly increased in proliferation after 48 h of incubation and reached the plateau phase after 96 h of incubation. In addition, β -catenin siRNA significantly decreased the cell viability after 48 h of treatment regardless of whether AuNPs were added. The LRP5 shRNA plasmid did not profoundly impact hPDLSCs compared to the control, except in the 48-h treatment.

β -Catenin in the Wnt signaling pathway after blocking LRP5 by treating hPDLSCs with shRNA plasmid

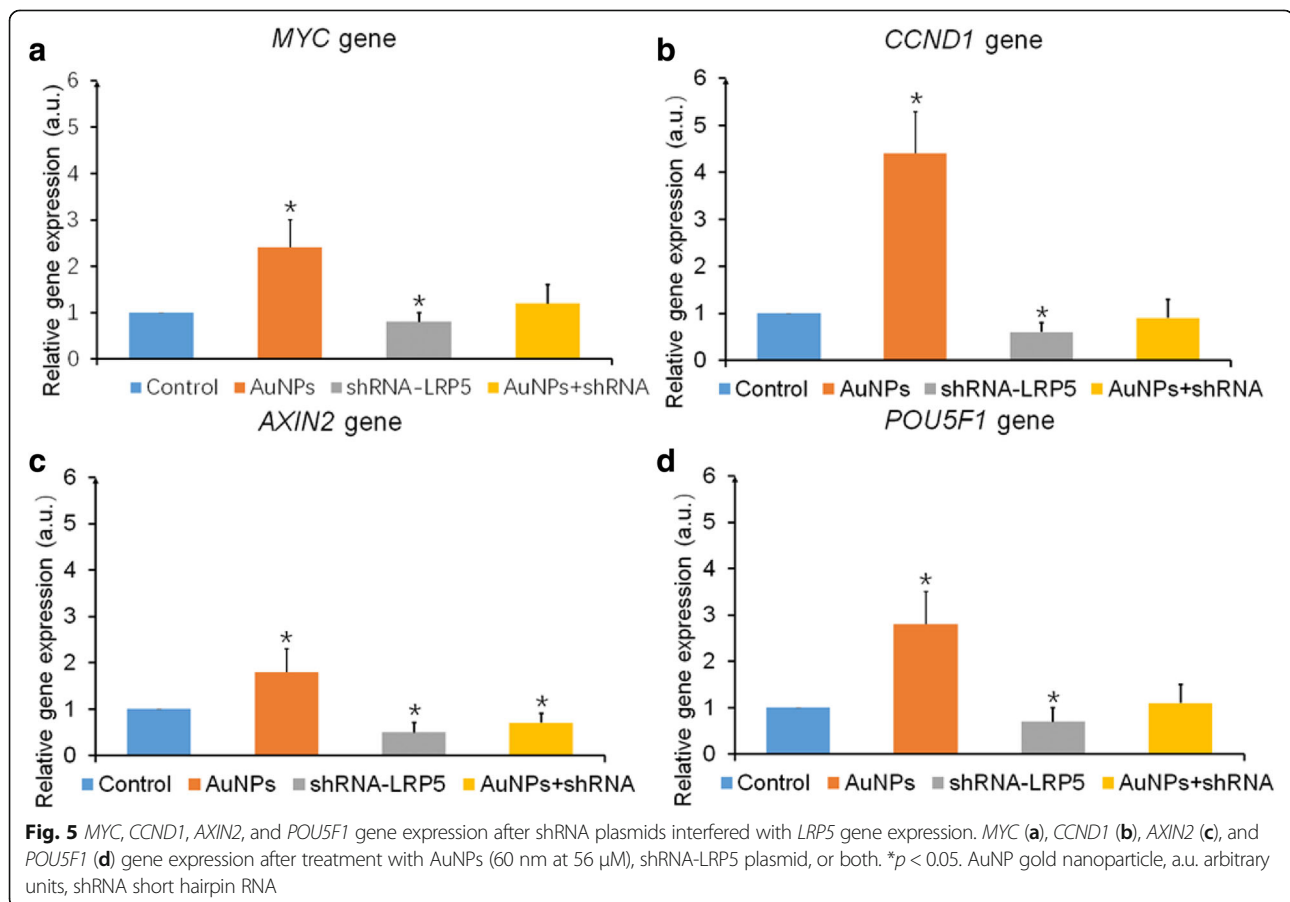
The *LRP5* gene expression was detected via real-time PCR after interference with shRNA plasmids. Both 1.5 and 2.0 $\mu\text{g}/\mu\text{l}$ concentration of the plasmid shRNA significantly downregulated the expression of the *LRP5* gene ($p < 0.05$). No significant difference was found between 1.5 and 2.0 $\mu\text{g}/\mu\text{l}$ (Fig. 3a). However, compared to 1.0 $\mu\text{g}/\mu\text{l}$ shRNA, 1.5 $\mu\text{g}/\mu\text{l}$ had a profound effect. Therefore, 1.5 $\mu\text{g}/\mu\text{l}$ shRNA plasmid was used for further experiments. Western

blot assay results (Fig. 3b) showed that, compared to the control group, 1.5 $\mu\text{g}/\mu\text{l}$ shRNA plasmid significantly downregulated the expression of LRP5 protein.

Results showed that β -catenin significantly increased after incubation with AuNPs (60 nm at 56 μM) compared to the control group. Protein β -catenin of hPDLSCs did not increase after shRNA plasmid interference with LRP5 and even incubation with AuNPs, compared to the control group (Fig. 3c).

Blocking of β -catenin via treating hPDLSCs with siRNA

siRNA 0, 0.25, 0.5, 0.75, or 1 μg was added to a 100 μl culture system. According to the results (Fig. 4a), 0.25, 0.5, 0.75, and 1 μg siRNA interfered with β -catenin expression, of which 0.5 μg siRNA significantly lowered β -catenin expression ($p < 0.05$) compared to 0.25 μg siRNA, while 0.5, 0.75, and 1 μg siRNA did not significantly downregulate β -catenin expression. Therefore, we chose 0.5 μg siRNA for further experiments. According to the results of western blot analysis (Fig. 4b) and compared to the control group, 0.5 μg siRNA significantly lowered β -catenin protein levels. Compared to the control, 0.5 μg siRNA profoundly decreased the β -catenin protein level, even under



AuNP addition. The AuNP group had no significant impact on the β -catenin protein level (Fig. 4c).

Expression of *MYC*, *CCND1*, *AXIN2*, and *POU5F1*

AuNPs (60 nm at 56 μ M) and/or 1.5 μ g shRNA plasmid were added to the hPDLSC culture system and the expressions of Wnt-related genes *MYC* (Fig. 5a), *CCND1* (Fig. 5b), *AXIN2* (Fig. 5c), and *POU5F1* (Fig. 5d) were detected via real-time PCR. AuNPs significantly increased the expression of *MYC*, *CCND1*, *AXIN2*, and *POU5F1* genes compared to control ($p < 0.05$). Interfering in the *LRP5* gene expression with shRNA plasmids downregulated expression of these four genes ($p < 0.05$). Under combined AuNP and shRNA plasmid treatment, only *AXIN2* gene expression was downregulated compared to control ($p < 0.05$).

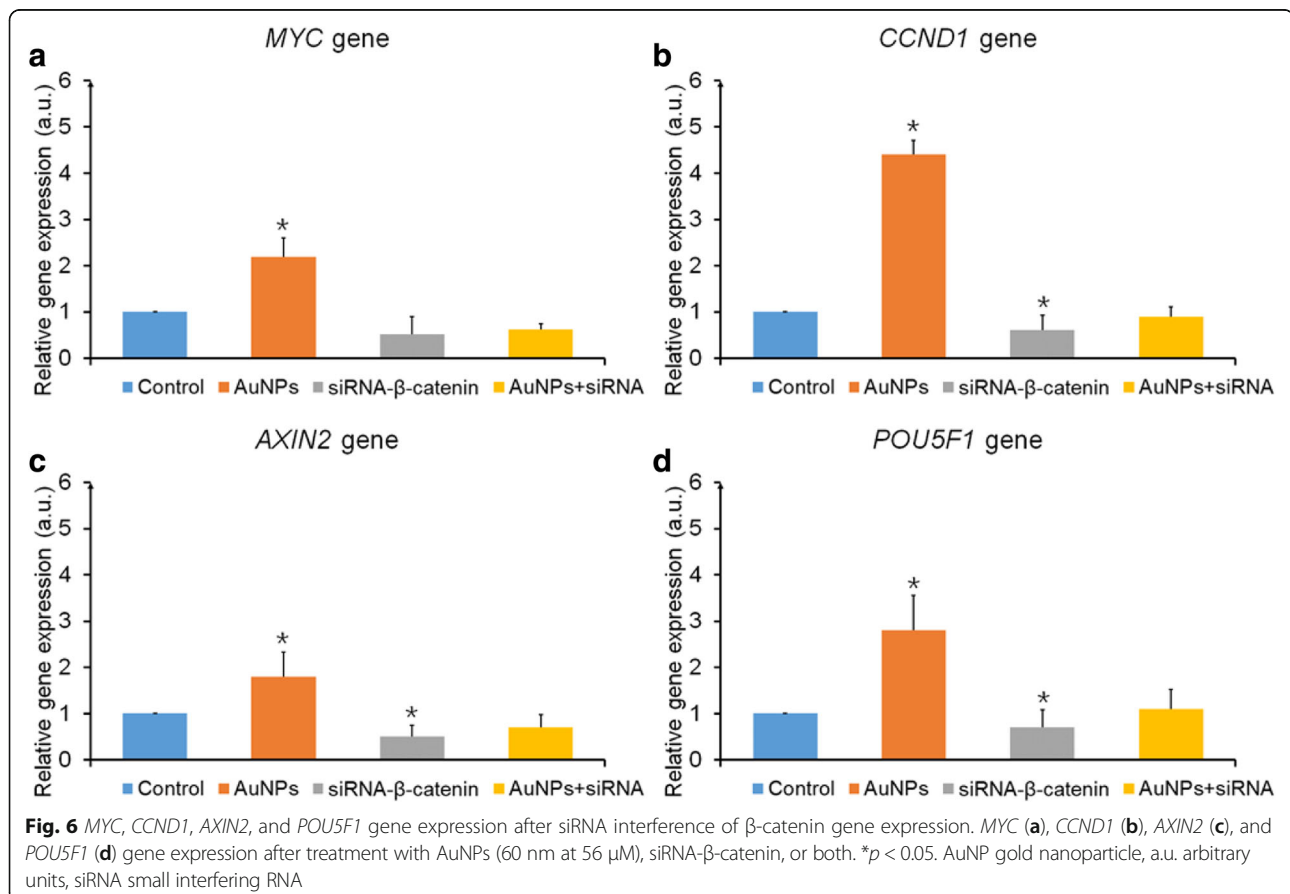
AuNPs (60 nm at 56 μ M) and/or 0.5 μ g siRNA to β -catenin were employed to investigate the resulting effects on hPDLSCs in 100 μ l medium per well and the Wnt/ β -catenin signaling pathway-related genes *MYC* (Fig. 6a), *CCND1* (Fig. 6b), *AXIN2* (Fig. 6c), and *POU5F1* (Fig. 6d) were also detected via real-time PCR. AuNPs upregulated the expression of four genes significantly ($p < 0.05$). β -catenin was interfered with by siRNA and, except for *MYC*, the other three genes were

downregulated ($p < 0.05$). Although β -catenin was part of the interference of siRNA, addition of AuNPs decreased the interference compared to the control.

Discussion

Recently, many in-vitro and in-vivo studies have indicated that AuNPs are nontoxic; however, other publications contradict these results [7, 8, 15]. In spite of this, most studies focus on the pathophysiology and not on the physiology of AuNPs. Multiple in-vitro studies have shown that AuNPs at a certain particle size and concentration promote cell proliferation [4, 10]. Our previous experiments confirmed that 60 nm AuNPs at a concentration of 56 μ M significantly promote the proliferation of human periodontal ligament cells and hPDLSCs [5]. However, the precise mechanism controlling the proliferation process still remains poorly understood.

AuNPs (60 nm, NIST standard reference material) in murine macrophages indicated no cytotoxicity under the exposure conditions tested, and neither do these elicit pro-inflammatory responses; however, the localization of AuNPs in intracellular vacuoles suggests endosomal containment and an uptake mechanism involving endocytosis [6]. The suggested size scale of AuNPs of 30–74 nm was

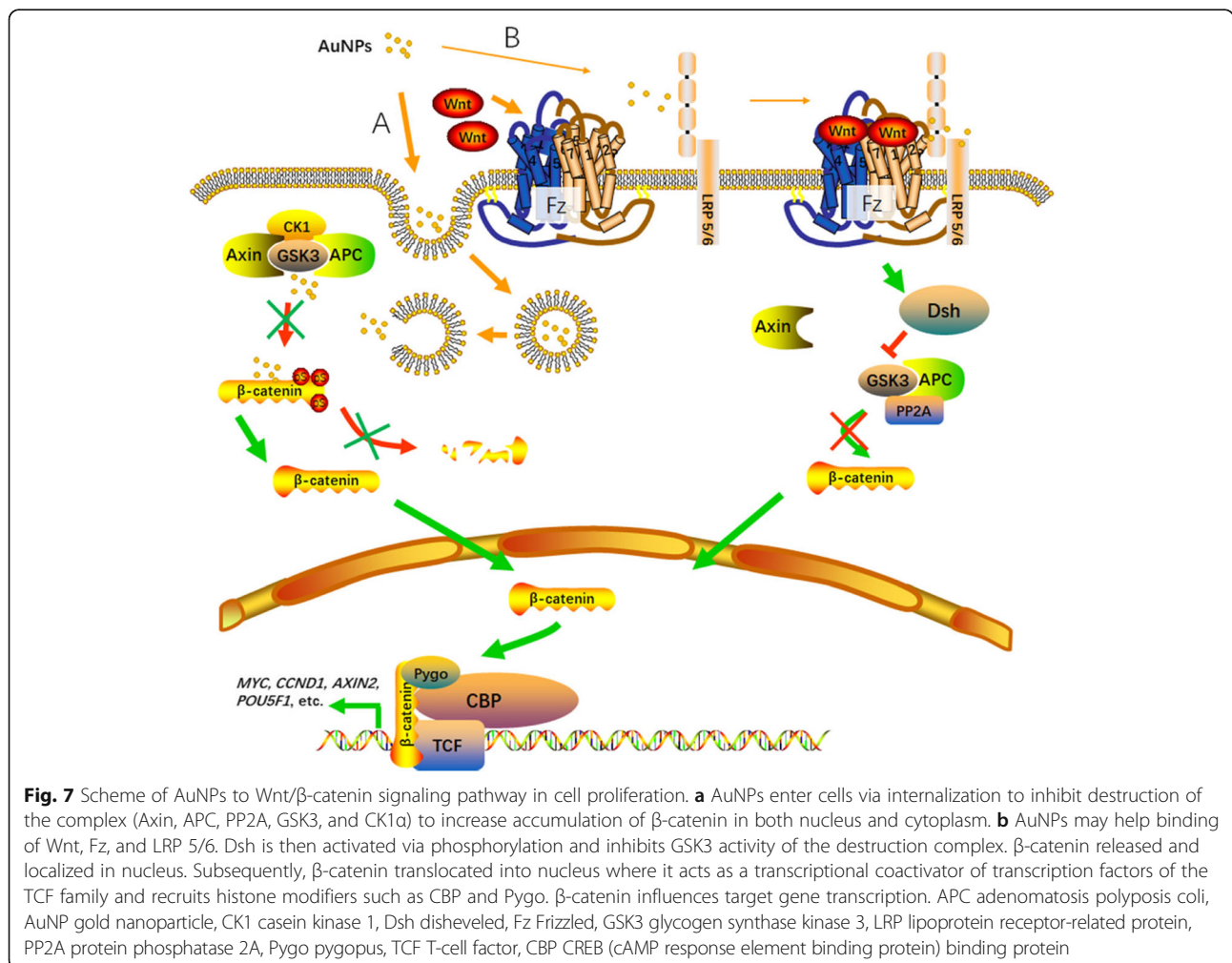


presumed to facilitate internalization. AuNP sizes via DLS were 6.3 ± 2.5 nm in GNP-S, 18.1 ± 2.1 nm in GNP-M, and 23.0 ± 10.2 nm in GNP-L, respectively. The GNP-S showed significant cytotoxicity by ROS; however, the GNP-L exerted little toxicity. Therefore, AuNPs sizes exceeding 40 nm tend to show no cytotoxicity [16]. Considering the uptake efficacy was not increasing with AuNP size, AuNP diameters not exceeding 70 nm were chosen accordingly.

AuNPs with different surface charge were employed to treat human bone marrow-derived mesenchymal stem cells (hMSCs). Surface-functionalized AuNPs were without acute toxicity. Positively charged AuNPs showed higher cellular uptake. AuNPs did not inhibit osteogenesis but ALP activity and calcium deposition were remarkably reduced in response to negatively charged AuNP treatment. The proteins in the biological medium were likely coating AuNPs, leading to a change of the initial surface charge of nanoparticles [7]. Therefore, the initial surface charge was not sufficiently convincing as a simple predictor of nanoparticle uptake and cytotoxicity.

Therefore, internalization was considered a reasonable route by which AuNPs enter cells [17] (Fig. 7).

Multiple signaling pathways, including SMAD, Notch, FGF, Hedgehog, and Wnt, have been closely associated with cell proliferation. Many related signaling pathways of stem cell proliferation have an intrusive cross-talk with Wnt signaling. Hence, the variation of the canonical Wnt signal pathway for the proliferation of hPDLSCs treated with AuNPs was investigated in this experimental study. Furthermore, AuNPs of size 30–74 nm had an uptake half-time of about 2 h, while the removal half-time was about 0.5 min, depending on cell types. Global gene expression analysis [18] and proteomic analysis [19] of AuNPs to human dermal fibroblasts indicated that the signaling pathways were inferred to respond during a short time. The Wnt signaling pathway was a robust candidate for the mechanism of AuNPs for promoting cell proliferation [11, 13]. This study showed that the proliferation of hPDLSCs boosted by AuNPs was significantly interfered with by β -catenin siRNA and slightly by LRP5 shRNA plasmid (Fig. 2). The Wnt signaling pathway was thus



suggested as an important mechanism with which AuNPs promote the growth of hPDLSCs.

Our experimental results showed that the AuNPs could activate LRP5 and/or β -catenin, and then activate downstream stem cell proliferation-related genes such as *MYC*, *CCND1*, *AXIN2*, and *POU5F1*. Interfering with the gene expressions of *LRP5* and β -catenin with shRNA plasmid and siRNA respectively, the expressions of the aforesaid genes were downregulated. After interfering with the gene expressions of *LRP5* and β -catenin with shRNA plasmid and siRNA, AuNPs were added and β -catenin protein did not increase. The expressions of the four tested genes were downregulated. The possible mechanisms with which AuNPs promote proliferation of human periodontal ligament stem cells were as follows: AuNPs outside the cells could promote LRP5, Wnt, and Frizzled protein aggregation and act on β -catenin in the cytoplasm. Phosphorylated β -catenin was dephosphorylated, transferred into the nuclei, and the expression levels of intracellular hPDLSC proliferation-related genes such as *MYC*, *CCND1*, *AXIN2*, and *POU5F1* increased, thus promoting hPDLSC proliferation.

When Wnt/ β -catenin signaling pathways were activated, β -catenin accumulated in the cytoplasm, entered the nucleus, and activated the downstream gene *CCND1*. A protein of the cell cycle, *CCND1* regulates cells via G1 to S phase transformation, cell proliferation, and differentiation. Studies have shown that *CCND1* expression and its upregulation have a close relationship with activated Wnt signaling pathways and increase the β -catenin expression level [20]; however, this is still disputed [21]. β -catenin proteins and TCF-4 activate *MYC* after combination [22], thus regulating the proliferation and differentiation of stem cells.

POU5F1 (also called *Oct4*) belongs to the family of POU and plays an important role in the maintenance of self-renewal of stem cells. In addition to the differentiation of mature stem cells, the expression of *POU5F1* gradually declines [23]. Therefore, its relative expression is very low in mature somatic cells in vivo and in vitro. The activation of Wnt signaling pathways promotes the expression of *POU5F1*. *AXIN2* participates in a β -catenin degradation compound. The *AXIN* family has two members: *AXIN1* and *AXIN2*. *AXIN2* has a very sensitive negative feedback effect on the Wnt signaling pathway. Whether *AXIN2* expression increases is an important indicator of whether the Wnt signaling pathway is involved in the regulation of cellular physiological and pathological physiology change. *AXIN2* as a Wnt signaling pathway inhibitory factor in the cell may regulate Wnt signal activation, where it leads to cell proliferation, prevents excessive cellular proliferation, and prevents malignant transformation [24, 25].

Cell proliferation is a complex process in which many factors interact. Our results suggest that AuNPs promote hPDLSC proliferation effectively, which is closely related to the Wnt/ β -catenin signaling pathway; however, their relationship and the underlying mechanism have yet to be further revealed. Further studies of AuNPs on the hPDLSC proliferation effect and the underlying mechanism will not only allow us to better understand AuNPs and their interaction with hPDLSCs, but can also provide excellent advancements of periodontal tissue engineering and thus lead to good seed cells.

Conclusion

In a previous experiment, we suggested that AuNPs (60 nm at 56 μ M) efficiently promote the proliferation of hPDLSCs/hPDLSCs in vitro. However, the AuNP-induced proliferation-boosting effect on hPDLSCs was diminished as β -catenin siRNA or LRP5 shRNA plasmid was added. The AuNP treatment equalized the accumulation of β -catenin in hPDLSCs. The proliferation of hPDLSCs was initiated by the Wnt/ β -catenin signaling pathway. Nevertheless, more details of the interaction between AuNP proliferation and the Wnt/ β -catenin signaling pathway are required to suggest more potential applications.

Abbreviations

AuNP: Gold nanoparticle; BBI: British Biocell International; FCM: Flow cytometry; hMSC: Human bone marrow-derived mesenchymal stem cell; hPDLSC: Human periodontal ligament cell; hPDLSC: Human periodontal ligament stem cell

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Authors' contributions

CL managed the samples, cell culture, and manuscript writing. ZL performed the AuNP quality and biomolecular detections using real-time PCR and western blot analysis. YZ analyzed and interpreted all of the data. AHF reviewed and modified writing of the manuscript. MZ contributed the ideas and financed the project. All authors read and approved the final manuscript.

Ethics approval and consent to participate

This work received approval from the Tongji University affiliated Stomatological Hospital Ethics Committee and informed consent was obtained from the LAR of the participant and the participant's guardian as premolar samples were obtained from minors in the study.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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