

Expression of Wnt10b, β -catenin, Rankl and Runx2 in the Periodontal Tissues During Orthodontic Tooth Movement in Rats

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Research

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Abstract

Objective

to investigate the expression of Wnt10b, Rankl, β -catenin and Runx2 mRNA in periodontal tissues during orthodontic tooth movement (OTM) in rats;

Materials and methods

36 healthy male SD rats were selected and divided into six groups randomly according to the time of the force applied (0 hour, 12 hours, 24 hours, 5 days, 7 days, 14 days), with 6 rats in each group. 0hour group served as the control group. Except for control group, an orthodontic force of 50g was applied to the upper-left first molar with a nickel-titanium tension spring. The expression of Wnt10b, Rankl, β -catenin and Runx2 during OTM were detected by Real-time PCR (RT-PCR).

Results

Compared with the control group, Wnt10b mRNA expression in the pressure side was inhibited at the initial stage, and then increased, reached peak at day 5. A strong expression of Rankl mRNA in the pressure side can be seen from 12 hours to 14 days. The expression level was consistently higher than control group, and it peaked at day 7. The initial expression of β -catenin and Runx2 mRNA in the periodontal tissues of the experimental groups were small in the tension side, but still higher than the control group. The expression increased gradually with the extension of time, and reached the peak at day 7 after force treatment.

Conclusion

Wnt10b, β -catenin, Rankl and Runx2 are related to periodontal tissues remodeling during OTM in rats; Wnt10b/ β -catenin signaling pathways participate in periodontal tissue remodeling during OTM.

Introduction

OTM is achieved by applying suitable mechanical force to periodontal tissue to make it reconstruct. Orthodontic force causes local stress changes in periodontal tissue, cell deformation and the release of a variety of cytokines, chemokines and inflammatory factor, such as interleukin-1 beta (IL-1 beta), tumor necrosis factor alpha (TNF), receptor activator of nuclear factor kappa B ligand (Rankl), Runx2 and so on[1, 2]. These factors were found to play a major role in bone reconstruction[3]. Moreover, interaction between osteoblast and osteoclast in periodontal tissue causes alveolar bone absorption on pressure side and bone formation on tension side[4, 5]. Wnt/ β -catenin pathways were involved in osteoblasts differentiation and bone formation[6]. In recent years, studies found that the classic Wnt/ β -catenin signaling pathways were involved in regulating the physiological metabolism of periodontal membrane[7]. Wnt/ β -catenin signaling pathway in the periodontal ligament fibroblasts can be activated[8]. As an important factor in Wnt signaling pathways, β -catenin plays an important role in the early stages of teeth formation[9]. Kuniaki pointed out that in osteoclast, transforming growth factor- β 1 (TGF- β 1) can stimulate the generation and maturation of wnt10b, and then promote the osteoblast mineralization[10]. Research reports that the overexpression of β -catenin in WNT signaling pathways will inhibits Rankl expression and then affect the osteoclast activity[11]. However, it is still not clear that the influence on periodontal tissue reconstruction of pressure side by classic WNT signaling pathways under orthodontic force. Runx2 functions as an osteogenic differentiation factor in bone cells. As a control factor in the early stages of osteoblast formation, Runx2 participates in the skull reconstruction, teeth formation, cartilage formation, and angiogenesis[12]. OTM mechanism is unclear. Though the establishment of animal model of OTM,

using RT-PCR to detect wnt10b and Rankl expression in periodontal tissue of pressure side, β -catenin and Runx2 expression in periodontal tissue of tension side at different time point, this study aims to preliminarily discuss whether Wnt/ β -catenin signaling pathways are involved in periodontal tissue reconstruction during OTM.

Materials And Methods

Establishment of orthodontic tooth movement model in rats

36 healthy adult male SD rats (provided by the Animal Experimental Center of Harbin Medical University) with body mass (190 ± 20) g were selected. According to different loading time, they were randomly designed to 6 groups (0 hour, 12 hours, 24 hours, 5 days, 7 days, 14 days), with 6 rats in each group. 0hour group acted as control group, which were not subjected to mechanical stress. Adaptive feeding is applied for 1 week before experiment. This experiment was approved by the Ethics Committee of Harbin Medical University.

The rats were anesthetized by intraperitoneal injection of chloral hydrates (10%, 3ml/kg). Except for control group, the nickel-titanium tension spring was ligated to the upper-left first molar and central incisors with ligation wire (0.08mm in diameter and 9mm in length) to generate 50g force to move the first molar in the mesial direction. Grind a groove of about 0.5mm at the cervical margin of the central incisors, fix the ligation wire on the central incisors with fluid resin to prevent it from falling off, and complete the rat OTM model (Fig. 1).

Periodontal tissue acquisition and RT-PCR experiment

Animals in each group were sacrificed separately after 0 hour, 12 hours, 24 hours, 5 days, 7 days and 14 days of force application, and the maxillae were separated. 2mm wide periodontal tissues were taken from the pressure area and tension area respectively, weighing about 100mg, and then grind the periodontal tissues in liquid nitrogen. Total RNA of samples was extracted by using TriZol Total RNA Isolation Reagent (Life Technologies BRL) according to the manufacturer's instructions. Ultraviolet spectrophotometer was used to detect the concentration and purity of RNA. Reverse transcription was performed according to 1000ng mRNA in each reaction system, which consists of 20ul stuff. The first strand of Complementary DNA (cDNA) (TAKARA kit) is synthesized and stored at -20°C .

ABI 7500 Fast Real-Time PCR system is as follows: SYBR Premix Ex Taq II 10ul, 0.8ul for each primer, cDNA template 2ul, ROX Reference Dye II 0.4ul, dH_2O 6u, totally 20 ul. Every sample is provided with three auxiliary holes, with β -actin as internal reference. The expression of Wnt10b, β -catenin, Rankl and Runx2 were detected with the 0hour group as the control group. The primers were synthesized by Sangon Biotech(shanghai), and the primer sequences were as follows (Table 1 and Table 2).

Results

Statistical analysis is performed by using SPSS 17.0 software, all data are presented as mean \pm standard deviation, $P < 0.05$ was statistically significant.

Expression of Wnt10b mRNA in the pressure side of periodontal tissue

Compared with the control group, the expression level of wnt10b mRNA was inhibited at the primary stage, increased after 24 hours, reached the peak at day 5, then decreased to the normal level at day 14 (Fig. 2).

Expression of Rankl mRNA in the pressure side of periodontal tissue

Compared with the control group, the expression level of Rankl mRNA in experimental groups was higher and gradually increased with the extension of the loading time ($P < 0.05$), reached the peak at day 7, and decreased to the normal level at day 14(Fig. 3).

Expression of β -catenin mRNA in the tension side of periodontal tissue

Compared with the control group, the expression amount of β -catenin mRNA increased after the force treatment ($P < 0.05$). The expression showed a gradual upward trend with the extension of loading time, and reached the peak value at day 7 after loading, decreased after 14 days ($P < 0.05$) (Fig. 4).

Expression of Runx2 mRNA in the tension side of periodontal tissue

Compared with the control group, the expression level of Runx2 mRNA increased after applying force ($P < 0.05$). The expression level of Runx2 mRNA showed an upward trend with the extension of the loading time, which increased slowly at the beginning, reached the highest level at day 7 after loading, and decreased after 14 days. Compared with the control group, there were statistically significant differences at all time points except for day 14 ($P < 0.05$) (Fig. 5).

Discussion

During OTM, when appropriate orthodontic force is applied, the osteoclast activity in the pressure area of the periodontal tissue is enhanced, the osteogenesis activity in the tension area increased, and the periodontal tissue reconstruction is active[13]. Wnt/ β -catenin classical signaling pathway and Rankl/Rank/OPG system are indispensable for osteoblast differentiation and maturation[14]. RANKL is one of the family members in RANKL/RANK/OPG system. It is a key cytokine that regulates osteoclast differentiation and bone resorption by binding to receptor activator of nuclear factor- κ B (RANK) and sending signals from cytoplasm to nucleus[15]. Patricia et al. confirmed that RANK positive staining cells were pre-osteoclasts at the initial stage of OTM, and RANKL promoted the maturation and fusion of pre-osteoclasts into activated osteoclasts[16]. Studies have demonstrated that the expression of RANKL mRNA increased in compression side as a consequence of application of orthodontic force[17]. In this experiment, compared with the control group, the expression level of Rankl mRNA in the pressure side increased with the extension of the time of force treatment, and reached its peak at day 7 after force treatment. The expression decreased but was still higher than that in the control group after 14 days of force treatment. The results are consistent with previous studies.

Runx2, also known as core binding factor A1 (Cbfa1), is an important transcription factor in bone formation and plays an pivotal role in bone formation, cartilage formation, and bone metabolism-related diseases[18]. Experiments show that in Runx2-deficient mice, the formation of intramembranous and cartilage bone were completely interrupted, and finally the bone and cartilage-like bone were completely lost[19]. Runx2 gene mutation will lead to Cleidocranial dysplasia (CCD)[12]. W et al. showed that Runx2 promoted osteoblast differentiation in periodontal tissues of tension side during the initial stage of rat tooth movement[20]. Study has shown that it had a significantly higher Runx2 mRNA after being loaded with mechanical stress in osteoblasts.[21] Similar to the results of the study, the expression of Runx2 in the tension side presented to be higher in experimental groups compared with the control group, and peaked at day 7 and began to decline after 14 days in this experiment.

The Wnt pathways include the classical Wnt / β -catenin signaling pathway, planar polar cell pathway, Wnt /Ca2 + pathway, and intracellular pathways that regulate spindle orientation and asymmetric cell division. Wnt protein is concentrated in the areas with more active differentiation in embryonic tissue, such as periodontal membrane of incisor and the cement-periodontal membrane interface[22]. Wnt/ β -catenin classical signaling pathways expressed at

different stages of osteoblast differentiation. When classical pathways are inhibited, osteogenesis imperfectness and osteoporosis will occur in the early stages of osteoblast differentiation[23, 24]. R et al. also found that Wnt/ β -catenin signaling pathway can promote bone repair of injured growth plate. Inhibition of β -catenin with β -catenin inhibitor can lead to reduced bone formation, but can promote the formation of cartilage in injure repair areas through animal experiments[25]. C et al. showed that Wnt10b^{-/-} mice showed reduced bone trabecular formation, and confirmed that Wnt10b promoted bone formation by affecting the differentiation of mesenchymal cells, which was considered as an important endogenous factor regulating bone formation[26]. P et al. demonstrated that mechanical loading could stimulate the transient accumulation of dephosphorylated β -catenin in the cytoplasm and it translocated into the nucleus through the cellular experiment in vitro, which suggested that mechanical force stimulation to the periodontal membrane cells could activate the expression of β -catenin[8]. Juan Lu et al. found that DKK1, Wnt3a, Wnt10b and β -catenin were involved in periodontal tissue remodeling during OTM in rats by immunohistochemistry[27]. In this experiment, it was found by RT-PCR method that Wnt10b mRNA expression tended to be weak at the pressure side in the initial stage of mechanical loading, and after 24 hours, the expression first increased, reached its peak at day 5 and then decreased, returned to the normal level at day 14 after force loading. This result is similar to the expression trend of Rankl mRNA, which indicated that 5–7 days after force treatment is the most active period of bone remodeling, and Wnt10b is an important factor involved in osteoblast differentiation in the classical signaling pathway. This result suggests that the Wnt classical signaling pathway is involved in the reconstruction of periodontal tissues on the pressure side, and there is a tendency of mesenchymal cell differentiation and bone formation during the active period of bone reconstruction. On the tension side, the expression of β -catenin mRNA increased gradually with the extension of loading time and reached the maximum at day 7. These changes were consistent with our previous experimental results and were similar to the expression trend of Runx2 mRNA, which indicated that bone remodeling was most active during 5–7 days of mechanical loading. In this experiment, the relative quantitative data of Wnt10b, β -catenin, Rankl and Runx2 mRNA were analyzed. It was found that the expression changes trend of wnt10b and Rankl mRNA on the pressure side were closely related in different periods, and both of the expression of β -catenin and Runx2 mRNA on the tension side gradually increased in parallel with time of mechanical stress.

Conclusion

Wnt10b, β -catenin, Rankl and Runx2 are expressed when appropriate orthodontic force is applied to periodontal tissue. Wnt/ β -catenin signaling pathway might play a pivotal role in periodontal tissue remodeling.

Declarations

Ethics approval and consent to participate

This study was approved by Ethics Committee of Harbin Medical University.

Consent for publication

Not applicable.

Competing interests

The authors declare no potential conflicts of interest.

Availability of data and material

All data generated or analyzed during the current study are included in this article.

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Authors' contributions

Author(First name, Last name)	Criteria 1(and/or)		Criteria 2(and/or)		Criteria 3	Criteria 4
	Substantially contributed to conception or design	contributed to acquisition,analysis,or interpretation of data	drafted the manuscript	critically revised the manuscript for important intellectual content	Gave final approval	Agree to be accountable for all aspects of the work in ensuring that questions relating to the accuracy or integrity of any part of the work are appropriately investigated and resolved
He Wang	contributed to conception and design	contributed to acquisition,analysis,and interpretation of data	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Siyan Chen	contributed to conception	contributed to acquisition	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Kaixin Ji	contributed to design	contributed to acquisition	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Minjie Zhang	contributed to design	contributed to acquisition and analysis	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Man Zhang	contributed to design	contributed to acquisition	drafted manuscript	critically revised the manuscript	Gave final approval	agree to be accountable for all aspects of the work ensuring accuracy and integrity

Yi Li	contributed to design	contributed to acquisition	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Yanyun Zhang	contributed to design	contributed to analysis	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity
Miaomiao Zhang	contributed to conception and design	contributed to acquisition, analysis, and interpretation	drafted manuscript	critically revised the manuscript	Gave final approval	agrees to be accountable for all aspects of the work ensuring accuracy and integrity

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Tables

Table 1: Pressure side

primer sequence		
RANKL	sense primer	5-AGGCTGGGCCAAGATCTCT-3´
	antisense primer	5-GATAGTCCGCAGGTACGCT-3´
Wnt10b	sense primer	5-TCAGGCTCTTAGGGAGGCT-3´
	antisense primer	5-ATTATCCATTCCCACCCGCC-3´
β-actin	sense primer	5-CACCCGCGAGTACAACCTT-3´
	antisense primer	5-CCCATACCCACCATCACACC-3´

Table 2: Tension side

primer sequence		
RUNX2	sense primer	5´-CGCCTCACAAACAACCACAG-3´
	antisense primer	5´-AATGACTCGGTTGGTCTCGG-3´
β-catenin	sense primer	5´-ATCATTCTGGCCAGTGGTGG-3´
	antisense primer	5´-GACAGCACCTTCAGCACTCT-3´
β-actin	sense primer	5´-TCAGGTCATCACTATCGGCAAT-3´
	antisense primer	5´-AAAGAAAGGGTGTAACGCA-3

Figures



Figure 1

Establishment of orthodontic tooth movement model in rats. The rats were anesthetized by intraperitoneal injection of chloral hydrates (10%, 3ml/kg). The limbs of the rats were extended and tied to the board. The mouth of the rats was propped open with a self-made mouth opener. the nickel-titanium tension spring was ligated to the left maxillary first molar and central incisors with ligation wire to generate 50g force to move the first molar in the mesial direction. Grind a groove of about 0.5mm at the cervical margin of the central incisors, fix the ligation wire on the central incisors with fluid resin.

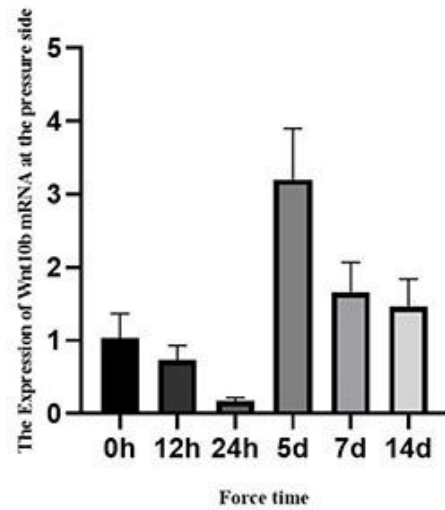


Figure 2

The expression of Wnt10b mRNA at the different period. The expression was inhibited initially. Over the next 24 hours, the expression level gradually decreased. It peaked on day 5, then decreased, and by day 14, returned to normal levels.

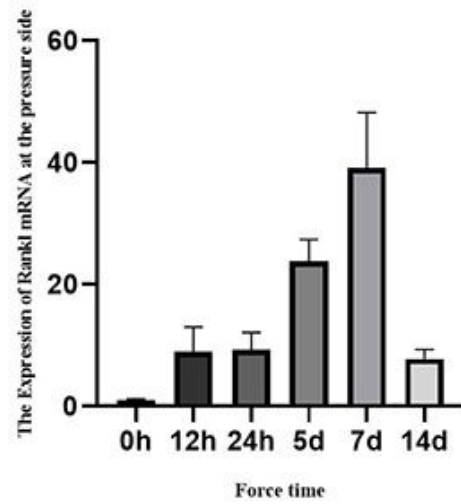


Figure 3

The expression of Rankl mRNA at the different period. The expression level of Rankl mRNA gradually increased with the extension of the loading time ($P < 0.05$), reached the peak on day 7, and decreased to the normal level after 14 days.

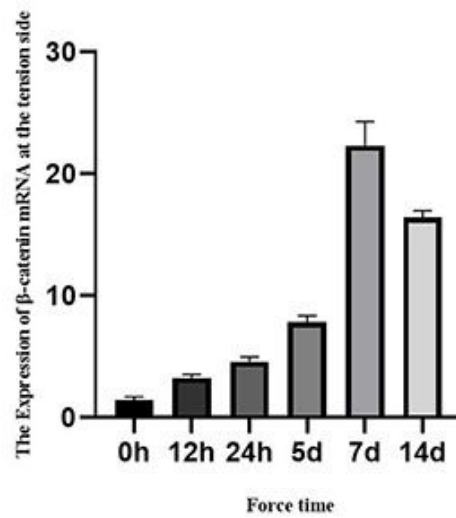


Figure 4

The expression of β -catenin mRNA at the different period. The expression increased gradually after force loading, and it peaked on day 7, decreased at day 14.

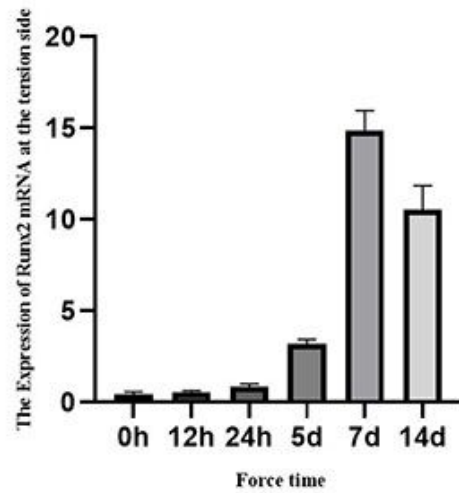


Figure 5

The expression of RunX2 mRNA at the different period. The expression level of Runx2 mRNA increased after applying force. It showed an upward trend with the extension of the loading time, which increased slowly at the beginning, reached the highest level on day 7, and decreased after day 14.