



# The circadian expression of osteogenic factors in periodontal tissue loading mechanical force: new concepts of the personalized orthodontic care

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Received: 16 October 2018 / Accepted: 11 January 2019 / Published online: 18 February 2019  
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## Abstract

**Objective** The need for orthodontic treatment continues to increase. Strategies that shorten the treatment course and reduce discomfort are most welcome in clinic. Circadian rhythm plays important role in various physiological processes, including bone formation. This study intended to depict a possible circadian releasing property of the osteogenic factors within the periodontal tissue during orthodontic treatment, which may direct a more efficient and satisfactory orthodontic treatment to the patient.

**Methods** Primary periodontal ligament cells (PDLs) were obtained from the Sprague-Dawley (SD) rats. An equibiaxial strain value of 12% was applied on rat PDLs (rPDLs). After 2 h stimuli of  $10^{-7}$  M dexamethasone (DX), the osteogenic genes' expressions were detected by real-time polymerase chain reaction (RT-PCR) at Zeitgeber times 0, 4, 8, 12, 16, 20, and 24. An orthodontic appliance was placed on 45 SD rats. Animals were maintained under 12-h light/dark periods and euthanized at 9 time points over the diurnal cycle. The orthodontic sensitive tissues of the mesial root of the maxillary first molar were collected for RT-PCR and immunohistological assay.

**Results** The rPDLs displayed typical fibroblastic spindle shape, and subcultured steadily in vitro. Induced by DX, the mRNA expression of Col-1, OPN, and IBSP within the loaded/unloaded rPDLs oscillated as that of the main clock gene Per-1. The osteogenic genes' expressions as well as the protein releases sustained a circadian oscillation trend in vivo.

**Conclusions** This study indicates the existence of a circadian rhythm of the osteogenic factors within the orthodontic sensitive tissues, which highlights the importance of precise timing of force loading in further orthodontic treatment. Thus, a periodicity pattern of orthodontic traction at night may prove a more efficient tooth movement while minimizing the treatment window and discomfort complains.

**Keywords** Circadian rhythm · Orthodontic tooth movement · Osteogenesis · Animal study · Mechanical force · Predictive preventive personalised medicine · Prognosis · Biomarker panel · Expression pattern · Animal model · Personalized orthodontic care

## Introduction

Orthodontic treatment deals with the study and treatment of malocclusions. Nowadays, the need for efficient orthodontic

treatment continues to increase across the world [1]. Historically, orthodontic tooth movement (OTM) was described as a site-specific bone remodeling through the histological change of periodontal ligament (PDL) under mechanical loading [2]. PDL is a layer of soft tissue between the tooth and the alveolar bone that provides the necessary microenvironment for cells involved in the alveolar bone remodeling, and is now recognized as the cellular basis for OTM [3]. Orthodontic mechanical force acts on PDL cells (PDLs) and induces the secretion of a series of genes relating inflammation, osteogenesis, and osteoclastogenesis, which finally leads to the reestablishment of the involved periodontal apparatus and completes the realignment of the tooth [4].

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Circadian rhythms are self-sustained endogenous oscillations occurring over a 24-h period [5]. The master circadian clock, locating in the suprachiasmatic nuclei (SCN), mainly synchronizes by light and coordinates the peripheral circadian clocks in nearly every part of the human body. The genes of the molecular clocks, such as period (Per), cryptochrome (Cry), brain and muscle Arnt-like protein (BMAL), and circadian locomotor output cycles kaput (CLOCK), exist in the SCN and peripheral tissue clocks [6]. These genes regulate many physiological processes, including organ development, metabolic functions, and tissue remodeling/repair in mammals, and are believed to exert important functions [7]. Lately, studies unveiled a relationship between circadian rhythm and mineralized tissue development, including osteogenesis and osteoclastogenesis [6, 8, 9]. The findings elicited a periodic expression mode of both osteogenesis and bone resorption related genes under the manipulation of peripheral nerve system. More recent studies also indicated that Cry2 and Per2 affected distinct pathways in the regulation of bone volume [5, 10, 11]. Meanwhile, researchers have observed that rat maxillary first molar receiving orthodontic force only at night moved faster than those loading only at daytime as well as those loading all day long [12]. This result strongly suggested the effect of circadian rhythm on bone metabolism during OTM. However, direct evidence of the periodicity expression of relative genes remains unavailable.

Hormonal signaling is one of the two routes through which the central circadian clock coordinates the peripheral circadian clocks [13]. In current study, we adopted dexamethasone (DX) to mimic the dictation of hormonal signaling from SCN to arouse the circadian variation of rat PDLs (rPDLs). Real-time polymerase chain reaction (RT-PCR) was applied to evaluate Q1, whether osteogenic gene expression of rPDLs exhibit periodicity; Q2, whether the addition of mechanical force changes the periodicity; Q3, whether the addition of mechanical force influences the gene expression level. Furthermore, the *in vivo* expression of the osteogenic factors was also tested by RT-PCR and immunohistological staining on an orthodontic model of rats. This study detected the existence of circadian rhythm in orthodontic sensitive tissue and laid foundation for chromobiological design of force timing design in a more effective personalized orthodontic care.

## Materials and methods

### Rat PDLs isolation and culture

Eight-week-old male Sprague-Dawley (SD) rats were obtained from the Tongji Medical College Animal Center. (Wuhan, China). All procedures concerning animal use were conformed to the guidelines of the Animal Ethics Committee of Huazhong University of Science and Technology (Wuhan, China) [14, 15]. rPDLs of the maxillary first molars were isolated and

cultured according to a modified way of the previous study [16]. Briefly, maxillary first molars were thoroughly washed with phosphate buffered saline (PBS) after careful extraction, and the periodontal ligament within the mid-third portion of the mesial root of the maxillary upper molar was scraped by a surgical steel scalpel (no. 11 blade, Junbei Ltd., Shanghai, China). All the explants were then placed onto the bottom of a six-well culture plates. Dulbecco's modified Eagle's medium (DMEM, Gibco, Grand Island, NY, USA) containing 10% fetal bovine serum (FBS, Gibco), 100 U/ml penicillin, and 100 mg/L streptomycin (Invitrogen, Carlsbad, CA, USA), was applied to sustain the primary rPDLs. Cells were cultured in a humidified 37 °C, 5% CO<sub>2</sub> incubator. In the present study, cells after passage 3 were used.

### Application of tensile strain to PDL cells *in vitro*

Primary rPDLs ( $5 \times 10^4$  per ml) were seeding into six-well, 35-mm flexible-bottomed Uniflex culture plates coated with type-I collagen. One day after seeding, the supplied DMEM containing 2% FBS were afforded for a further 24 h. After that, the seeded cells were subjected to an intermittent deformation of 12% for 6 s every 90 s with a FX-5000™ Tension System (Flexcell Corporation, Hillsborough, NC, USA), as described previously [17].

### Treatment with dexamethasone on rPDLs

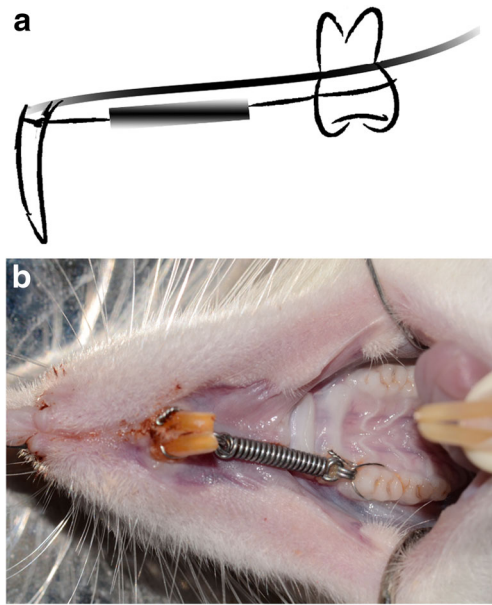
After 4 h loading of PDLs,  $10^{-7}$  M DX (Sigma-Aldrich) was added to the culture medium to mimic the dictation of hormonal signaling from SCN. The culture medium was replaced 2 h later with fresh supplied DMEM containing 2% FBS, and cells were collected every 3 h. All time interval calculations are based at the indicated Zeitgeber (ZT—is an event that provides the sets of a biological clock) and ZT-0 is considered 2 h after medium changing. Cells are first harvested at ZT-0, and then are collected every 3 h for a total of 9 time points. RNAs were isolated and analyzed by RT-PCR [18].

### Animals for OTM

Forty-five 12-week-old male SD rats with a weight of  $250 \text{ g} \pm 15 \text{ g}$  were obtained from the Tongji Medical College Animal Center. Five rats were kept in one cage under identical condition: room temperature at  $24 \text{ }^\circ\text{C} \pm 1 \text{ }^\circ\text{C}$ , humidity at 55~65%, standardized laboratory rat diet, free access to water, and a controlled light condition (12 h:12-h light: dark, 330 lx).

### *In vivo* orthodontic treatment

After a 2-week-adapting period, all animals received an orthodontic appliance that was installed before (Fig. 1) [19, 20]. Generally, the maxillary right first molar was connected with



**Fig. 1** Established orthodontic tooth movement (OTM) model on Sprague-Dawley (SD) rats. **a** The schematic illustration of the animal model. **b** Photograph of the orthodontic device in the mouth of SD rats

both maxillary central incisors with a nickel-titanium closed-coil spring (3M Unitek, Monrovia, California, USA) to perform the mesial movement. The orthodontic force was continuous and maintained about 30 g for 2 weeks.

### Sample harvest

Two weeks after orthodontic appliance placement, animals were euthanized. All time interval calculations are based at the indicated Zeitgeber. At the designed time point, orthodontic sensitive tissue of each animal, including the cervical half of the mesial root of the moved molar, its adjacent alveolar and the periodontal ligament, was immediately dissected and bisected into the buccal and palatal halves along the maximal sagittal plane. One half was directly cryoconserved in liquid nitrogen for RT-PCR test ( $n = 5$  for each group); the other half was quickly immersed into 4% paraformaldehyde for histomorphometric analysis ( $n = 5$  for each group). The cryoconserved samples were stored at  $-80\text{ }^{\circ}\text{C}$  before tissue homogenizations (mixer mill MM 200, 30 s/sample at 25 Hz, Retsch® GmbH, Haan, Germany). The total RNAs were then extracted by Trizol reagent (Invitrogen, USA).

### Immunohistological evaluation

The fixed pieces were immersed in 4% paraformaldehyde solution for 24 h at  $4\text{ }^{\circ}\text{C}$  followed by decalcification in EDTA for 15 days at room temperature. After embedding in paraffin,  $5\text{ }\mu\text{m}$  sections were prepared for immunohistochemistry (IHC) [21, 22]. Briefly, sections were deparaffinized, rehydrated, and incubated with primary antibodies against OPN (Abcam, USA, 1:200 dilution) overnight at  $4\text{ }^{\circ}\text{C}$ . The

slides were then incubated with HRP-conjugated secondary antibody before staining in diaminobenzidine (DAB) kit. Finally, all sections were counterstained with hematoxylin. The average optical density (AOD) of the immunohistochemistry assay was measured using Image-Pro Plus 6.0 software.

### Real-time quantitative PCR

cDNA was synthesized using a Prime-Script™ RT reagent kit (Takara Bio, Shiga, Japan) according to the manufacturer's instructions. The related genes, including Period 1 (Per-1), collagen-1 (Col-1), osteopontin (OPN), and integrin-binding sialoprotein (IBSP) were measured, and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as the house-keeping gene for the normalization of RNA expression levels. The PCR primer sequences are displayed in Table 1.

### Statistical analysis

To determine the periodicity, the data of RT-PCR was tested with Time Series Analysis Single Cosinor v.6.0. The period was determined using the chronobiometric ellipse test. The mean  $\pm$  standard deviation of the gene expression levels were compared to ZT-0 within a group using the Student's *t* test. The gene expression levels at per time point were compared between the static and the dynamic group using the ANOVA at the 0.950 probability level by an SAS 6.12 software package (SAS, Cary, NC, USA). [9]

## Results

### Primary rPDLCs culture

Five days after tissue explant placed on the bottom of the dish, sporadic long-shuttle rPDLCs were seen around the edge of it. After 2 weeks, 80~90% confluence was reached as shown in Fig. 2.

### Per-1 and osteogenesis-related genes expression of loaded/unloaded PDLCs

The gene expression levels of Col-1, OPN, and IBSP were assessed every 3 h over a 24-h period to determine whether these osteogenic genes expression of rPDLCs coincided with the circadian cycle-related gene under loaded/unloaded condition. The results showed that, in the static condition, the osteogenic genes' levels fluctuated in a manner largely coincident with the light/dark cycle indicated by Per-1. Specifically, the levels of osteogenic related genes initially began to rise at 0:00 h, peaking between 03:00 and 06:00 h, and then declined starting at 06:00 h (Col-1) and 09:00 h (OPN and IBSP). To be noticed, levels of Col-1 have two

**Table 1** Nucleotide sequences for real-time polymerase chain reaction primers (rat)

Genes	Forward primer	Reverse primer	Accession no.
PER-1	ACATCTGAATACACTCTCCGCAAC	GCAGGCGAGATGGTGTAGTAGAG	NM_001034125.1
COL1-1	CAGATTGAGAACATCCGCAGC	CGGAACCTTCGTTCCATACTC	NM_053304.1
OPN	GATGAACAGTATCCCGATGCC	CCCTCTGCTTATACTCCTTGGAC	NM_012881.2
IBSP	GAAAGAGCAGCACGGTTGAGTAT	CGTCATAGGTTTCATACGCAGTG	NM_012587.2
$\beta$ -actin	TGCTATGTTGCCCTAGACTTCG	GTTGGCATAGAGGTCTTTACGG	NM_031144.3

peaks as that of Per-1, and they (03:00 h and 21:00 h) were significantly higher than levels at 00:00 h ( $p < 0.01$ ). Otherwise, levels of OPN and IBSP at 06:00 h~09:00 h were statistically higher as that of the 00:00 h ( $p < 0.01$ ). On the other hand, dexamethasone also induced the oscillation of the osteogenic-related gene expressions of rPDLcs under loaded situation. Interestingly, the fluctuation mode kept identically when compared to unloaded situation. The differences of Col-1, OPN, and IBSP expression between static/tensile situations mainly appears in the elevated levels instead of oscillating pattern, while the expression level of Per-1 remains steady under both culture conditions (Fig. 3). A synergistic effect was also noticed to increase the level of Col-1 (03:00 h and 21:00 h) and IBSP (03:00 h and 06:00 h) at the highest expression points.

### Per-1 and osteogenesis-related genes expression during OTM

To determine whether or not the expression of osteogenic genes fluctuated over a 24-h time period during orthodontic treatment, real-time RT-PCR analyses were conducted on total RNA isolated from the periodontal tissue of the moving teeth in the orthodontic model on rats. As shown in Fig. 4, Per-1, Col-1, and OPN mRNA levels showed two troughs within a 24-h observation window. For Per-1, the highest expressions appeared at 03:00 h and 15:00 h, while Col-1 and OPN appeared at 03:00 h~06:00 h and 18:00 h. The trough of IBSP

delayed 12 h as compared to in vitro test and increased to  $(13.22 \pm 1.68)$  folds to IBSP at 00:00 h in vivo.

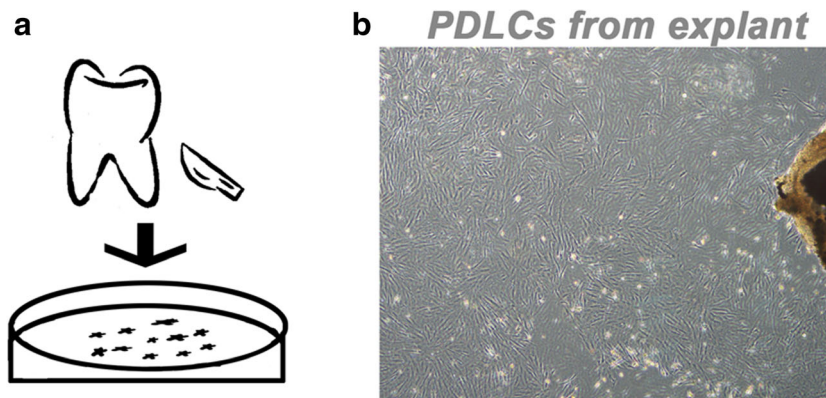
### OPN expression during OTM

Immunohistochemical analyses were conducted to detect the expression of OPN in periodontal tissue during OTM [22]. OPN was released in cells of the periodontal ligament region between alveolar bone and root surface. Expression of protein was upregulated in the 06:00 h and 18:00 h time points compared with the 00:00 h and 12:00 h (Fig. 5). The AOD was also measured. We noted that the 06:00 h and 18:00 h points presented an increase in the content of OPN. Compared to 00:00 h, 12:00 h, and 18:00 h time points, the 06:00 h showed significantly enhanced levels of OPN (Fig. 5e).

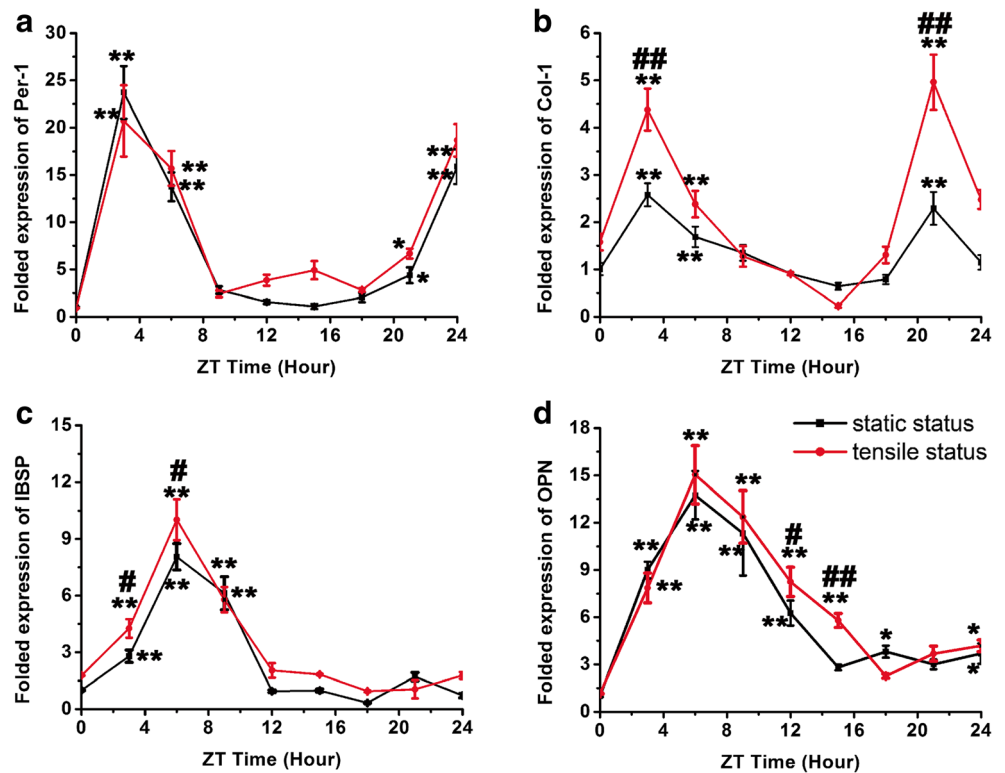
### Discussion

Biological clocks have evolved as an organic evolution and an adaptation to life on a rhythmic planet, synchronizing physiological processes to the environmental light-dark [23]. Although accumulating evidences implied the significance of circadian rhythm in bone metabolism, there are few studies focusing on that within the periodontal tissue. The current study showed that the expression of osteogenic-related genes oscillated along with that of main clock gene in rPDLcs. Also, the in vivo expressions of circadian gene and osteogenic genes/proteins in periodontal tissue were detected on the

**Fig. 2** Primary PDLcs culture. **a** The schematic illustration of the harvestment of PDLcs from the upper first molar of SD rat. **b** Spindle-shaped PDLcs migrated from explant 14 days after implantation



**Fig. 3** Osteogenic genes' expression of loaded/unloaded rPDLCs stimulated by dexamethasone. **a** Per-1 expression of rPDLCs. **b** Col-1 expression of rPDLCs. **c** OPN expression of rPDLCs. **d** IBSP expression of rPDLCs. (\* $p < 0.05$  and \*\* $p < 0.01$  as compared to 00:00 h gene expression in identical culture environment; # $p < 0.05$  and ## $p < 0.01$  as compared to the static culture environment group at identical time point)



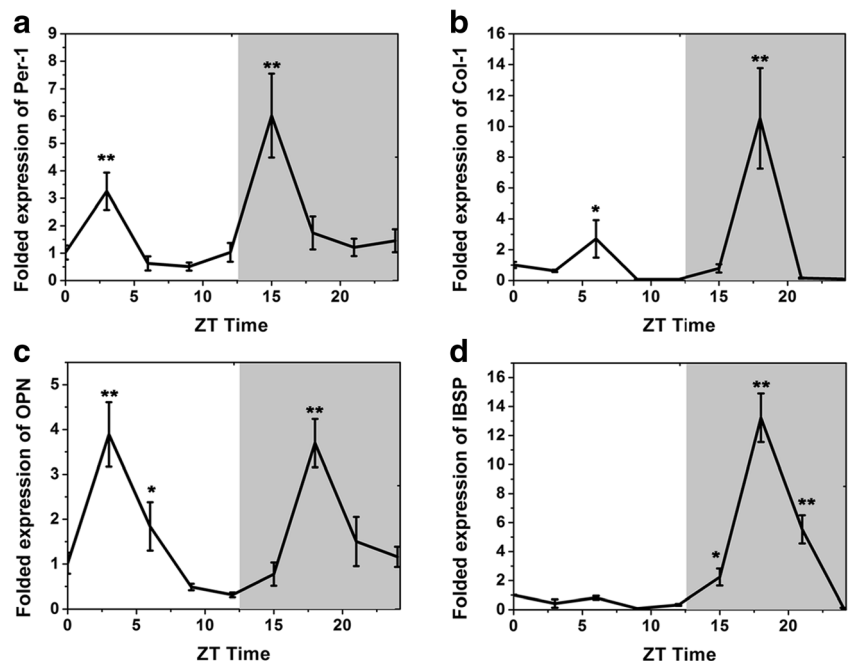
OTM model for the first time to demonstrate the potential function of circadian cycle in orthodontic moving tooth.

PDLCs play essential role in periodontal metabolism. Previously, researchers found that cyclic tension promotes osteogenic differentiation in human PDLCs, which indicated the main role of PDLCs during OTM on the molecular level [24]. Similarly, we chose PDLCs to reflect the status of periodontal

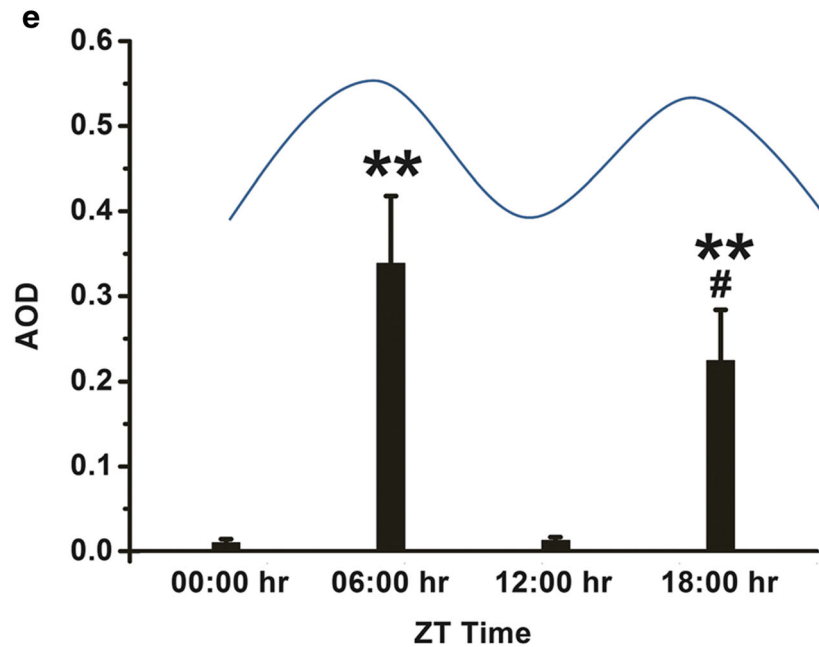
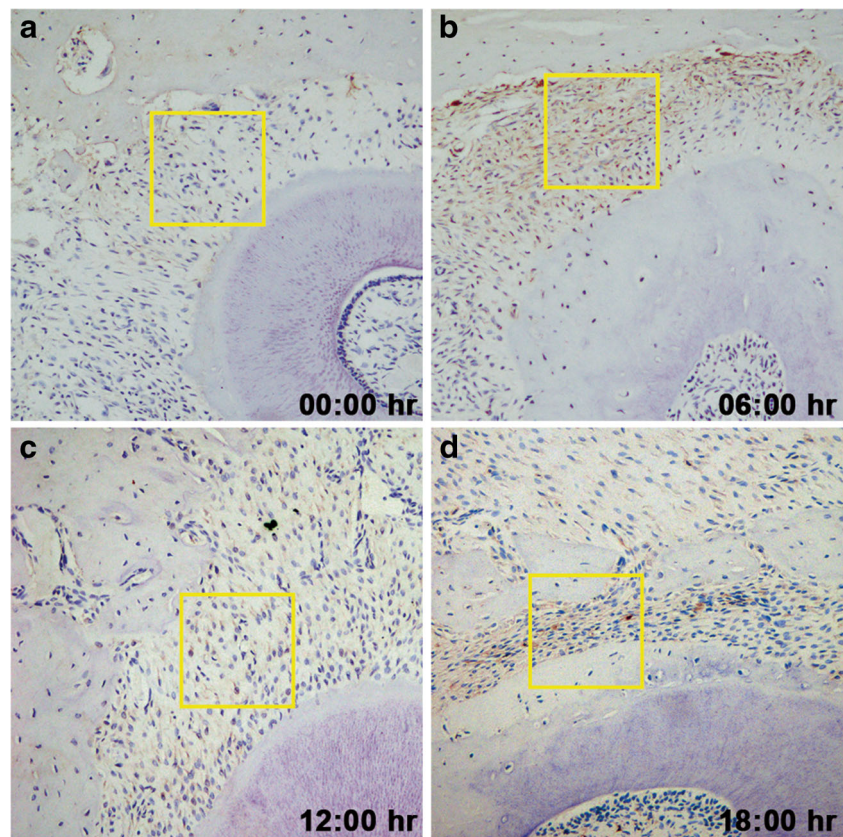
tissue during OTM. On the other hand, an equibiaxial strain value of 12% was adopted on rPDLCs to represent the in vivo deformation of the cells during occlusal loading and OTM [17, 25].

To mimic the hormonal signal, one of the main routes from SCN to transmit central circadian time to peripheral tissue, we introduce dexamethasone to induce the clock gene Per-1

**Fig. 4** Osteogenic genes' expression of periodontal tissue during OTM. **a** Per-1 expression. **b** Col-1 expression. **c** OPN expression. **d** IBSP expression. (\* $p < 0.05$  and \*\* $p < 0.01$  as compared to 00:00 h gene expression)



**Fig. 5** Immunohistological staining of OPN in the periodontal tissue during OTM. **a** OPN staining at the 00:00 h time point. **b** OPN staining at the 06:00 h time point. **c** OPN staining in the 12:00 h time point. **d** OPN staining in the 18:00 h time point. **e** Average optical density (AOD) of OPN immunohistological staining at 00:00 h, 06:00 h, 12:00 h, and 18:00 h (\* $p < 0.05$  and \*\* $p < 0.01$  as compared to 00:00 h AOD; # $p < 0.05$  as compared to 06:00 h AOD)



expression. Consistent with previous reports,  $10^{-7}$  M DX stimulation was effective to initiate the oscillatory expression of Per1 in rPDLCs as shown in the human osteoblast line by Komoto et al. [6]. The maximum expression emerged at 03:00 h and 24:00 h, and exhibited a sinusoidal wave-like pattern. This indicated the existence of a molecular expression

periodicity in periodontal apparatus. The oscillatory expressions of osteogenic genes Col-1, OPN, and IBSP in rPDLCs were also stimulated, with the Col-1 expression pattern being closely similar to that of Per1. As Col-1, OPN, and IBSP are important markers of osteoblastogenesis and cementogenesis [26], their circadian expression specifically indicated an

oscillatory property of the periodontal formation. We also noticed that the intervention of mechanical force did not interfere with the circadian expression of these osteogenic genes. Interestingly, a synergistic effect between hormonal signal (DX) and orthodontic force (mechanical loading) was first noted in the expression of rPDLcs' osteogenic genes on the mRNA level, which coincided with previous findings that there are considerable variations in tooth movement to orthodontic force when the force is applied at different times of the day in rats [12, 27].

To further clarify the circadian rhythm of bone metabolism during OTM, the *in vivo* study was conducted on rats. A nickel-titanium closed-coil spring appliance was applied to establish an orthodontic model on the right maxillary first molar. The force applied was 30 g, which moderately affects the normal tooth of rats and has been shown to be comparable to the force used in human tooth movement [28–30]. The maxillary first molar of rat has five roots. The compressed zone appeared on the cervical half of the middle and mesial roots (mesial sides) [31, 32]. Since the mesial root is the biggest and relatively easy to expose, we took it and its neighboring alveolar as the orthodontic sensitive sample for the histological analysis, including RT-PCR and IHC. Also, the observation level was taken within the cervical half of the mesial root. Meanwhile, to avoid the interference of tension or compression side differences, we divided the orthodontic sensitive tissue along the mesio-distal direction. Thus, the images were taken at the buccal/palatal side of the mesial root of the moved molar. The stress distribution could be considered identically based on previous report [32]. The RT-PCR results showed that the expression of osteogenic genes as Col-1, OPN, and IBSP all demonstrated a circadian cycle pattern. A synchronizing change could also be noticed in these genes, as they simultaneously reached their highest expression level at 18:00 h. This time point was 3 h later compared to *Per1*, which suggested the controlling role of clock gene in osteogenesis [33, 34]. The immunohistological staining of OPN further proved the fluctuated secretion pattern on the protein expression level. This trend was basically consistent with the circadian rhythm of osteocalcin in the maxillomandibular complex [33], which strongly implied the circadian oscillation state of osteogenic-related factors in periodontal tissue during OTM. This finding first unveiled the relationship between OMT and circadian rhythm on a protein level. It is worth mentioning that, our previous clinical observation found 21:00 h the acrophase of the rapid palate expansion [35]. In this study, we further proved that the acrophase of the osteogenic genes were mainly appeared around 17:00 h~18:00 h (2 or 3 h earlier), which may explained the human change from a mRNA level and indicate new strategy of activation time for a more efficient orthodontic treatment.

## Outlook and expert recommendations

This research is especially relevant to the objectives of predictive, preventive, and personalized medicine (PPPM) given that, circadian rhythm affects bone formation, in turn, contribute to a periodicity-regulated OTM [36]. The results presented herein are used to guide further research, like accelerating the tooth movement rate by controlling peripheral oscillation. Also, depending on individual genetic and environmental conditions, different activation time should be recognized to give personalized care. In a broader interest for maxillofacial surgery, rehabilitation, and plastic surgery, the role played by circadian rhythm and its underlying potential in provoking more efficient soft/hard tissue reconstruction worth further consideration [37].

In conclusion, our research highlights the circadian rhythm of osteogenic factors within the periodontal tissue during OTM. This study may benefit the prognosis and personalized treatment design of the current orthodontic therapy through the manipulation of circadian rhythm genes. Furthermore, the understanding of the physiological adjustment in OMT from the perspective of light-dark cycle would be more comprehensive and may suggest revolutionary change to its current concept and therapy.

**Acknowledgements** The authors would like to thank the Tongji Medical College Animal Center, Anyi Li and the Stomatology faculty of Tongji Medical College Zuoqiao Yin for technical support.

**Funding** The study was funded by the National Science Foundation of China (No. 81170986 and No.81800891).

## Compliance with ethical standards

All procedures concerning animal use were conformed to the guidelines of the Animal Ethics Committee of Huazhong University of Science and Technology (Wuhan, China)

**Competing interests** The authors declare that they have no competing interests.

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