Orthodontic treatment requires moving and aligning the teeth into a favorable position aesthetically and functionally. To achieve that, the teeth are subjected to forces to push them to their new position. Loading of force on the crown causes tipping of tooth, subsequently the periodontal ligament is compressed adjacent to the alveolar bone on the side toward which the force is directed. On the opposite side of the root, the periodontal ligament is stretched and undergoing tension. Blood vessels are compressed and blood flow is decreased on the compression side, hence less oxygen supply is received on that side. Compression and tension are triggering specific signaling factors and mediators, which create local environment and gradients to regulate remodeling of the periodontal ligament and bone for tooth movement. In this review, we highlight the structures and mediators released in the very narrow zone periodontal ligament during orthodontic tooth loading.
1. INTRODUCTION

The periodontal ligament (PDL) is a specialised connective tissue apparatus, located between the cementum of the tooth root and the bone forming the alveolar socket (alveolo-dental ligament) [1]. It acts as a shock absorber which transmits the biting forces to the alveolar bone. Cementum of the tooth root has intrinsic fibres (Ebner fibrils) run horizontally in a circle around the root and extrinsic PDL fibers (Sharpey’s fibers) that are inserted into the cementum in one side and to the alveolar bone of the socket in the other side [2]. The extracellular matrix of PDL is composed of a network of fibrous structural proteins embedded in amorphous ground substance surrounding cells [3].

The ground substance of PDL is 70% water and is thought to have a role in absorbing stress loads applied on the tooth. Tissue fluids tend to increase within the matrix of the PDL ground substance in case of injury or inflammation. PDL matrix contains several noncollagenous matrix proteins released from the blood vessels passing through the ligament or produced locally by resident cells, these include alkaline phosphatase [4], proteoglycans [5], glycosaminoglycan, glycolipids and glycoproteins such as undulin, tenascin, and fibronectin [6]. The cells of periodontal ligament include fibroblasts, osteoblasts, osteoclasts, epithelial rests of Malassez, macrophages, monocytes, mesenchymal or stem-like cells, cementoblasts and odontoclasts, therefore, PDL considered as a cell reservoir for tissue proliferation and repair, homeostasis or regeneration [7-9]. Periodontal ligament also contains blood and lymphatic vessels, and nerve fibres. The width PDL ranges from 0.15 in the middle third of the root to 0.38 mm in the coronal and apical third of the root. It has been reported that PDL space decreases progressively with age or in certain bone disease conditions [1].

Regarding the aging of PDL, it has been found that the space between the alveolar bone and the cementum is gradually reduced over lifespan. Though bone formation decreases and bone resorption increases with aging, yet the PDL region tends to be thinner. This is probably attributed to the increased cementum thickness with age on the expense of PDL width. In addition, the reduced or loss of occlusal forces on aging result in osteoporotic alveolar bone and atrophy of PDL fibres, hence a narrower PDL width [10,11].

Orthodontic forces result in structural changes of the dental tissues features. Cementum, PDL, bone, gingival tissues and bioactive substances (chemical mediators, enzymes, growth factors, neuropeptides and ligands) are collectively reacting and responding to the forces, subsequently periodontal tissues are re-organised and movement of teeth can be achieved [12,13].

The levels of these bioactive molecules may be affected (either increased or decreased) by certain medical conditions or medications taken. Unwanted disruption of the levels these molecules will aggravate or suppress the inflammation induced upon loading. If the normal scenario sequence of aseptic inflammation is disrupted, the orthodontic tooth movement may either accelerated, delayed or even halted. Hence, clinically these molecules can be used as targets to control or adjust the teeth movement to achieve the most favorable aesthetic outcome of orthodontic treatment.

In this article, we provide insights into PDL components and bioactive substances released within the PDL region during orthodontic tooth movement.

2. MAINTENANCE OF PERIODONTAL SPACE

PDL is occupying a space surrounded by two mineralised tissues (root cementum from one side and alveolar bone from the other side). This narrow space is maintained by molecules secreted by PDL cells that control calcification and avoid the ankylosis of tooth root with the adjacent alveolar bone throughout the life. The activities of these cells in secreting molecules of various types of proteins are increased during orthodontic movement. Certain mineralisation inhibitors such as matrix gamma-carboxyglutamic acid protein, a vitamin-K-dependent molecule; osteogenic transcription factor Msx2 and glycosaminoglycans or tripeptide–cementum attachment protein may have a pivotal role in maintaining the PDL space [14,15]. It has been found that balanced and reciprocal activities between bone sialoprotein and osteopontin may play a significant role in preserving and maintaining an unmineralized
PDL space. Osteopontin is expressed in the periodontium since tooth root development and has a potential role in periodontal tissue formation and maintaining the PDL - alveolar bone interface [16]. Due to its chemotactic activity in attracting osteoclasts precursor, osteopontin has been implicated in hastening tooth movement and root resorption during orthodontic treatment [17-19]. It has been reported that ERM has a role in the osteoclastogenesis of the alveolar socket, a process that subsequently leads to maintain the periodontal space during orthodontic treatment [20-22].

3. ADAPTATION OF PDL TO FUNCTION

Teeth are subjected to two types of forces: continuous, light and horizontal forces represented by soft tissues pressure from the tongue, cheeks and lips; and intermittent, heavy and vertical forces produced by chewing [23]. The PDL has the capacity to adapt to load changes. Increasing the loads for long term can increase markedly the width and thickness of the PDL fiber bundles. Contrariwise, a decrease in the load subjected on teeth leads to thinning of periodontal ligament and a reduction in thickness and number of the fiber bundles [24]. These load changes of the PDL are taking place concurrently with adaptive changes at the interface with cementum and alveolar bone [25].

4. PERIODONTAL LIGAMENT COMPONENTS IMPLICATED IN ORTHODONTIC MOVEMENT

Epithelial rests of Malassez (ERM) cells are remnants of the disintegrated Hertwig's epithelial root sheath, are found in the PDL space. TALIC et al. [26] conducted an animal study and found that ERM cells tend to proliferate and increase in size during experimental tooth movement. Orthodontic tooth movement causes an increased level of IL-6 released by human PDL fibroblast [27]. IL-6, in turn, promotes the proliferation and migration of ERM [28]. ERM proliferation is reflecting the potential role for these epithelial cells in accelerating the collagen turnover in the periodontal ligament during tooth movement [26-28].

It has been shown that ERM and PDL fibroblast have a reciprocal activity based on the singling and stimuli they receive. Periodontal fibroblasts produce collagen and a collagenase inhibitor [29], while ERM produce latent collagenase which subsequently transforms to active collagenase by enzymatic cleavage [30]. ERM plays a role in the rejuvenation of collagen in the periodontal ligament. During PDL remodelling, collagenase of ERM degrades the collagen molecule into three-fourth and one-fourth peptide fragments, which then phagocytize by fibroblasts. Consequently, fibroblasts synthesize and secret collagenase inhibitor which inhibits the active collagenase enzyme to form an inactive enzyme inhibitor complex [29,30].

It has been reported that the ERM contributes to the homeostasis of periodontium. ERM is thought to be involved in the induced tooth movement by increasing epidermal growth factor (EGF) production in PDL, preventing ankyloses and helping to repair root resorption areas by inducing cementogenesis [30,31]. EGF in return, stimulates ERM proliferation and maintains their growth and integrity [32]. EGF receptors are composed of transmembrane proteins that trigger tyrosine kinase intracellularly and initiate cellular events that activate cell division and remodelling/regeneration of periodontal tissue [33]. EGF secreted by the ERM is directly involved in osteoclastogenesis of alveolar bone through the suppression of osteoprotegerin, an important decoy receptor for RANKL [34]. Continuous release of EGF results in the upregulation of expression of monocyte chemoattractant protein-1 (MCP1), and hence, promoting the resorption of the adjacent alveolar bone, that ultimately leads to tooth movement, prevents ankylosis, as well as, maintaining the PDL space [20,30].

As part of its multifunction, ERM secret inflammatory mediators such as prostaglandins and enamel proteins such as amelogenin and amelin. Amelin, in turn, promotes the formation of bone matrix proteins, osteopontin, osteoprotegerin, BMP-2 and sialoprotein. These proteins participate in the regeneration and repair of periodontium [35-38]. Furthermore, it has been reported that disruption of periodontal integrity by trauma or orthodontic movement results in the expression of amyloid enamel protein APIN in ERM, which indicates that this protein may play a role in the initial phases of periodontal regeneration [39]. Mechanical forces induced by orthodontic appliance enhance the expression of bone matrix proteins, such as osteopontin, which stimulates resorption and repair of root and helps to prevent ankylosis [36,38,40].
4.1 Inflammatory Mediators, Cytokine and Transcription Factors

The orthodontic treatment induces an acute inflammatory response at its initial phase. In the early phase of tooth movement, leukocytes tend to migrate out of blood capillaries of periodontal ligament and producing cytokines which subsequently promote the excretion of prostaglandins and other mediators [41,42]. Prostaglandins, which are also released by ERM, specifically prostaglandin E2, induce the recruitment and activation of osteoclasts and stimulate bone resorption/remodeling [20,30]. Application of orthodontic force stresses the extracellular matrix and deforms the nearby osteocytes of alveolar bone. Subsequently, this deformation opens the hemi-channels of the stressed osteocytes to release prostaglandins and recruit osteoclasts, which enhance orthodontic tooth movement [43,44]. Therefore, amount of prostaglandins produced plays a key role in orthodontic movement and also affects the rate of tooth movement [45,46].

Orthodontic forces enhance the production of various cytokines within the PDL space. Previous studies highlighted the role of IL-1β, IL-6, and TNF-α in bone remodelling [47,48,49]. The concentration of these cytokines is increased in the gingival crevicular fluid at the beginning stage of orthodontic treatment (12th and 24th hour) [50]. The cytokines are released by the infiltrating polymorphonuclear leukocytes and PDL cells in response to the applied forces [51]. The maximum level of the released molecules is attained three days after the application of orthodontic loading [52]. Cytokines induce local stimulation of osteoclasts which perform bone resorption [53,54]. The process involves nuclear transcription factor Kappa B (NF-kB) (receptor activator of NF-kB ligand, RANKL) which activates receptors of TNF-α (TNF–R1) at osteoblasts [55,56]. RANKL binds on a receptor at osteoclast named RANK. Subsequently RANKL activates osteoclasts to perform osteoclastogenesis [57]. Resorption is suppressed through prevention of RANKL binding for RANK by the decoy receptor osteoprotegerin, followed by the inhibition of osteoclasts differentiation [58,59].

This phase of alveolar bone resorption is shortly followed by decreasing the levels of secreted pro-inflammatory cytokines and reduction of blood vessels permeability [60]. As cytokines release is diminished in the tissues, the levels of these cytokines in gingival crevicular fluid and the number of inflammatory cells is reduced after 7–10 days since the commencement of the orthodontic forces application [52,61]. The later events overlap with the beginning stage of regeneration/repairing of tooth supporting tissues, which lasts for around 9 days [62]. During the restoration of periodontal tissue, stimulation of osteoblasts takes place through the overexpression of various growth factors and interleukins which enhance the osteoid formation and suppress bone resorption [63,64].

During the early stage of orthodontic loading and after the initiation of early stage periodontal remodelling, there is a second wave of other cytokines such as IL-8 which was shown to have an increased level in the gingival crevicular fluid [6,45]. IL-1β, IL-6, and TNF-α stimulate the production of the other cytokines in monocytes, macrophages, epithelial cells, and fibroblasts of periodontium, this in turn, provoke the release of another wave of IL-1β, IL-6, IL-2 and TNF-α [65,66].

High mobility group box 1 (HMGB1), a late inflammatory cytokine, is secreted by the periodontal ligament cells in response to mechanical stimuli [67]. It is also produced by stimulated dendritic cells, necrotic cells, macrophages and monocytes as a cytokine mediator of Inflammation [68,69]. On compression, HMGB1 provokes local inflammatory responses by stimulating the secretion of chemokines and cytokines from stressed cells [70,71]. The released HMGB1, would, in turn, stimulates human monocytes to produce and release TNF, IL-1α, IL-1β, IL-1RA, IL-6, IL-8, IL-10, macrophage inflammatory protein (MIP)–1alpha, and MIP-1beta in cultured periodontal ligament cells [70,71]. Furthermore, HMGB1 induces monocytes chemotaxis, macrophage migration and osteoclastogenesis [69,72]. HMGB1 is produced in the tension zone of PDL during orthodontic tooth movement, where it plays a role in tissues remodelling through stimulation of cells proliferation and formation of mineralized nodule in PDL cells [73,74].

Previous study has reported that periostin, a 90 kDa extracellular matrix protein, has an inhibitory effect on HMGB1 [75]. Periostin has a potential role in preserving the integrity of PDL collagen fibrils during orthodontic tooth movement, and its deficiency causes impairment of collagen
fibers degradation at the compression side of periodontal ligament, which obstructs orthodontic tooth movement [67].

Periostin is primarily expressed in connective tissues subjected to mechanical stress and stretch, such as skin, heart valves, PDL, bones and tendons [76-78]. Xu et al. [79] found that tensile stress load upregulates the levels of periostin in animals and human periodontal ligament fibroblasts during orthodontic tooth movement. Previous studies have also found that the periostin expression is regulated by transforming growth factor β, and that periostin promotes fibrillogenesis in PDL; and enhances migration of fibroblasts and osteoblasts, hence, it is essential for the PDL and bone remodelling during orthodontic loading [80,81].

4.2 Heat Shock Protein

Orthodontic force disturbs the blood supply of periodontal ligament and leads to hypoxia and ischemia in the tension zone during the early stages of tooth movement [82]. Consequently, these disturbances lead to the activation of cellular self-defense activities to reduce the stress and preserve the structure of periodontal ligament. It has been reported that heat shock protein (HSP) is released to serve this mission. HSP is upregulated in periodontal ligament during the early stages of orthodontic tooth movement [83-85].

Previous studies revealed that HSP70 has a regulatory role in the reduction of cytokines and RANKL expression and controlling the inflammatory periodontal ligament cell response to compression forces [84,86]. Inhibition of HSP70 resulted in significant reduction in cell division and proliferation and wound healing; while necrosis and apoptosis, osteoclastic differentiation and monocyte adhesion were highly increased in order to limit the tissue damage during orthodontic tooth movement [87].

4.3 Periodontal Ligament Blood Vessels

During orthodontic tooth movement, periodontal ligament blood vessels are significantly involved in the renovation of tooth surrounding tissues [3,88,89]. Several signals are generated after the compression of extracellular matrix which surrounds the cells of endothelia of blood vessels. These signals activate the restructuring of existing vessels and also formation of new blood vessels in the periodontal ligament [90].

Mechanical forces cause distortion of the nerve terminals which in response they release vasoactive neurotransmitters [91]. In the periodontal ligament, most terminals are near blood-vessel walls. Subsequently, the released neurotransmitters activate the capillary endothelial cells receptors to bind circulating leukocytes, enhancing their migration out of the capillaries [92]. The migrating leukocytes secrete many signal molecules, including cytokines and growth factors, which play important role in the aseptic inflammatory reaction that stimulates periodontal ligament and alveolar bone remodeling and facilitates the movement of teeth [43].

4.4 Periodontal Ligament Cells Role in Osteoclastogenesis

It has been shown that periodontal ligament fibroblasts are adapted to the bacterial invasion and mechanical loading by secreting high amounts osteoclastogenesis-inducing factors. Hence, they possibly contribute to the escalated osteoclast recruitment observed during periodontitis and to orthodontic tooth movement [54]. It is believed that periodontal ligament cells regulate osteoclast differentiation through RANKL stimulation and osteoprotegerin inhibition, and also support osteoclastogenesis through cell-to-cell contact [93]. Cell-cell adhesion between periodontal ligament cells and osteoclast precursors significantly enhances the upregulation of genes for osteoclast differentiation and the eventual formation of osteoclasts [94].

4.5 Periodontal Ligament Cells Role in Osteoblastogenesis (Osteogenesis)

Periodontal ligament fibroblasts subjected to tensile strain were induced to express ephrin-B2 protein which in turn stimulate the osteoblasts of the alveolar bone and increase their osteoblastogenic gene expression at the tension sites during orthodontic tooth movement [95]. At the compression site of root, compressive forces induce periodontal ligament fibroblasts to produce ephrin-A2, while the expression of ephrin-B2 fet is down-regulated. Ephrin-A2 suppresses osteoblastogenic gene expression (RUNX2, ALPL) of osteoblasts and decreases the signs of osteoblastic differentiation at the compression sites [95,96].
4.6 Growth Factors

Orthodontic loading alters the flow of blood in the periodontal ligament and leads to the production of the angiogenic regulator, vascular endothelial growth factor (VEGF), and tissue growth factor β (TGF-β) by the periodontal ligament fibroblasts and circulating leukocytes [97,98]. It has been shown that the level of anti-inflammatory cytokine (IL-4, IL-10, TGF-β) is higher at the tension side compared to the compression side of the root, which is probably attributed to its role in the process of osteogenesis and tissue formation at the tension side [3,41].

4.7 Enzymes

The inducible nitric oxide synthase (iNOS), endothelial nitric oxide synthase (eNOS) and neuronal nitric oxide synthase (nNOS) are three important enzymes released in the periodontal space during the early phase of orthodontic loading [99,100]. These enzymes are necessary for the production of nitric oxide (NO) which is an important regulator of bone remodelling [101,102]. Previous studies revealed that stimulation of NOS increases tooth movement, while inhibition of NOS reduces that movement [103,104]. The gene expression of synthase enzymes is controlled by pro-inflammatory cytokines (IL-1β, TNF-α) and anti-inflammatory cytokines (IL-4, IL-10, TGF-β), which are secreted during tissues remodelling [105]. Orthodontic loading on animals have shown that iNOS is associated with bone resorption at the compression zone while eNOS is associated with the bone formation at the tension zone [100].

4.8 Regulating Occlusal Forces

The periodontal ligament has a pivotal role in controlling occlusal force associated with muscle spindles in jaw-closing muscles [106,107]. There are multiple receptors mediating the somatosensation in the periodontal ligament. These receptors include Ruffini endings and Merkel cells for mechanosensation processing. Nociception and itching are processed by the free nerve endings of Aδ- and C-fibers [108-110]. Merkel cells of the epithelial rest of Malassez release neuropeptides which control the mechanosensation function [108,111]. Application of orthodontic forces leads to the constriction of the blood capillaries of the periodontal ligament in the pressure area that result in focal necrosis, with histological features of hyalinization [112]. During this process, collagen fibers formation is regulated by ERM in order to promote adaptation to the orthodontic forces, homeostasis and maintenance of the periodontal tissues [12,30,38].

5. CONCLUSION

The periodontal ligament plays a significant role in the orthodontic tooth movement for being the field and source where many bioactive molecules are released in response to the orthodontic forces. Periodontal ligament is the first periodontal structure receives and reacts to the impact of loading. It is also important for being a pivotal part of theater for many molecular events which take place in order to achieve the orthodontic treatment. Clinically, several molecules produced by the periodontal cells upon orthodontic loading can be used as targets to accelerate or improve the movement of teeth.

CONSENT

It is not applicable.

ETHICAL APPROVAL

It is not applicable.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

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