



Review

# Influence of the Periodontal Disease, the Most Prevalent Inflammatory Event, in Peroxisome Proliferator-Activated Receptors Linking Nutrition and Energy Metabolism

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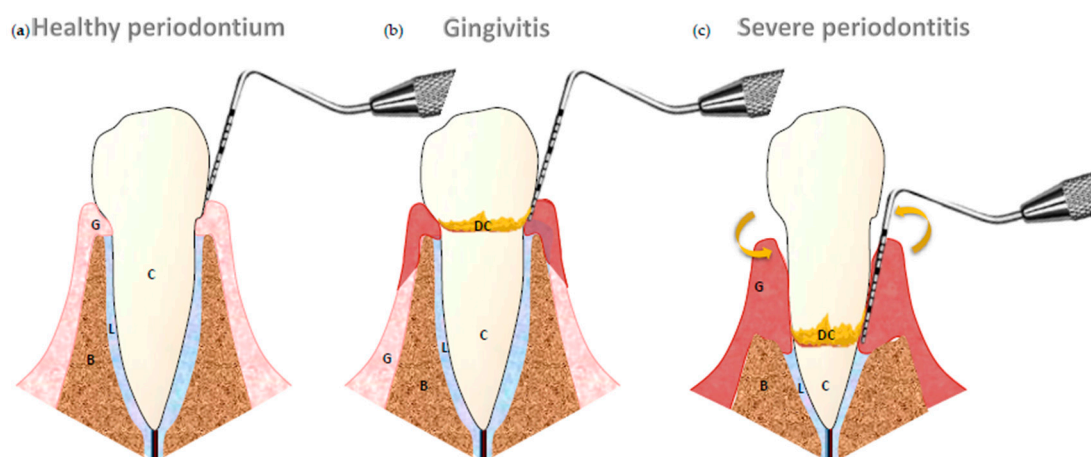
**Abstract:** Periodontal disease is considered one of the main pathologic diseases occurring in humans. Its pathologic process involves inflammatory reactions producing periodontal bone resorption and the tooth loss. But some patients do not present an evident clinical inflammation with bone resorption, and in others, the inflammation is prominent without bone resorption. A key question could be to investigate a different way of responding to aggression. Inflammation requires a complex intracellular metabolic process, starting with the harmful recognition and activation of the inflammasome, continues the energy supply with the alteration of oxidative stress conditions, and finishes with the elimination of the aggression with autophagy/apoptosis mechanisms, then concludes with recovery. Peroxisome proliferator-activated receptors (PPARs) are essential molecules produced in inflammation, and its genes and its activation have been related to periodontal disease. Also, an important aspect is the influence of PPARs in bone metabolism; the main periodontitis symptom is bone loss and PPAR $\gamma$  activation that can downregulate the bone resorption in experimental periodontitis, PPAR $\gamma$ -coated titanium dental implant surfaces could carry the antiinflammatory gene and restrain inflammation. PPARs could be one of the meeting background points with atherosclerosis/cardiovascular disease, diabetes and metabolic syndrome showing a modified proinflammatory statement such as it is described in periodontitis.

**Keywords:** periodontitis; inflammasome; mitochondria; autophagy; atherosclerosis; diabetes; metabolic syndrome

## 1. Introduction

Plaque-induced periodontal diseases are mixed infections which cause inflammation in the tissues around the teeth; this site is known as the periodontium and compromises all of its components. We divided it into two general categories based on whether illness to periodontium is reversible or not. So, respectively, we can differentiate between gingivitis and periodontitis [1].

Gingivitis supposes the presence of gingival inflammation without loss of gingival attachment; the site affected is limited to the gingiva. On the other hand, periodontitis is an inflammatory disease of the periodontal tissues with a more complex pathogenic process. Main symptoms in periodontitis are an apical migration of gingival attachment, a progressive periodontal bone loss and the formation of a periodontal pocket. The endpoint in the progression of this process is the tooth loss. This disease does not compromise the life of the patient, but produces a decrease in their quality of life. Severe periodontitis may result in discomfort for the patient, tooth mobility, abscess formation, occasional pain, difficulty in chewing and finally, tooth loss [2] (Figure 1).



**Figure 1.** (a) Healthy periodontium: gingiva (G), periodontal ligament (L), root cementum (C) and alveolar bone (B); (b) Gingivitis; there is presence of calculus (DC). The inflammation is reversible; and (c) Severe form of chronic periodontitis: gingival inflammation, depth pocket, subgingival calculus and mobility.

Nowadays, periodontal diseases take real importance in our society on account of their high prevalence. The prevalence of gingivitis is over 80% with the peak being reached during puberty [3]. Chronic periodontitis affects about 50% of the European population, and over 10% may suffer from an aggressive form, with this prevalence increasing up to 70–85% at 60–65 years of age [4]. Also, it is expected that this prevalence data is maintaining during the next decade in Europe [5]. Therefore, periodontal diseases are considered one of the most common diseases.

The key questions are the etiological and pathogenic processes involved, but many aspects are still unclear. Authors try to explain why in some patients, the disease produced by bacterial infection is limited to the gingiva and in others ones produces bone loss, and also the degree of bone resorption could be more or less severe, and even in aggressive periodontitis a lack of plaque and inflammation is described.

To explain it, researches have been focused on individual factors like the host response or microbial infection. Traditionally, the main cause of periodontal diseases is attributed to some bacterial species that colonize subgingival sites or pockets, which are the responsible for the inflammatory reaction. Nowadays, our treatment consists of reducing and eliminating the subgingival microorganisms. But, we have to recognize some significant limitations: we do not obtain success in all the patients, prevention is not feasible, we do not have any ways to identify the susceptible population, and a complete regeneration of periodontal tissues is presently impracticable. Besides, plaque is not essential in aggressive forms of periodontitis, and the control of the periodontal disease becomes more difficult [6]. Certain microorganisms are related with the pathogenesis of periodontal disease, but the evidence does not support the invasion of a periodontal pathogen as a key step in this process [7]. Therefore, periodontitis is produced by an imbalance among microbiota/dental biofilm and the inflammatory response that have to play a key role.

Inflammation is the main common backgrounds, with others systemic diseases such as atherosclerosis and atherosclerosis, diabetes and metabolic syndrome. These have been correlated with periodontitis in epidemiological studies. But, while periodontitis does not kill people, these others pathologies jeopardize the life of patients and are the main causes of death in our countries. All have a complex etiology, with a combination of non-modifiable risk factors (age or genetic), and modifiable risk factors: tobacco, alcohol, malnutrition, and chronic infection [8].

Inflammation is a physiological response to stress which involves a multitude of processes. Firstly, it is necessary that innate immune system recognizes the aggression made by microorganisms, toxins and chemical compounds. These are identified as the pathogen-associated molecular patterns

(PAMPs) or danger-associated molecular patterns (DAMPs) that are recognized via the pattern recognition receptors (PRRs) [9]. The cytoplasmic PAMPs and endogenous DAMPs activate PRRs and induce an intracellular signaling leading to the activation of inflammasome-driven inflammation [10]. Inflammasomes are large multiprotein complexes localized in the cytoplasm of the cell; it is essential in initiating innate immune responses. They produce the maturation of pro-inflammatory cytokines and the activation of inflammatory cell death [11]. Several inflammasomes are well-characterized for their role in recognition of PAMPs and DAMPs such as NLRP3 (nucleotide-binding leucine-rich repeat containing family, pyrin domain containing 3) [12]. Inflammasome activity is central to appropriate host defense and protection from sepsis, but excessive inflammasome activation could be detrimental to health [13].

Secondly, to maintain this inflammatory reaction, a sufficient supply of energy obtained from nutrients of the diet is necessary. But nowadays, humans have an excessive offer of nutrients that produce metabolic disorders. In the origin of these is a “proinflammatory” state produced by excessive caloric intake and over-nutrition and other innate conditions related with inflammation [14]. These nutrients are metabolized via the Krebs cycle, fatty-acid oxidation and amino acid oxidation to obtain adenosine 5'-triphosphate (ATP). This ATP is the fuel in cells of living organisms and the mitochondrion is the major source of ATP via cell respiration [15]. During respiration, oxygen is used to oxidize molecules rich in carbon and hydrogen, and a reduction of oxygen to water. This reaction realizes free radicals and/or reactive oxygen species (ROS); both are capable of damage different substrates. The antioxidants are responsible for preventing the oxidative damage, but any defect in the respiratory chain, mitochondrial diseases and age could alter this balance and produce oxidative stress [16]. Also, we cannot forget free radicals and ROS are essential in the physiological processes, but when an antioxidant system is unable to counteract their action efficiently, tissue and cell damage result [17]. So, the role of oxidative stress is essential in the mechanisms of inflammatory condition [14].

Finally, to maintain and control inflammation, cells need to function properly. The degradation of organelles and the elimination of invading pathogens are essential for the cell's homeostasis. In this sense, one of the main catabolic pathways is autophagy. Autophagy allows for an appropriate response to stress and extrinsic or intrinsic insults [18]. Impaired autophagy produce various pathologies, including cardiovascular disease and periodontitis [19,20], and is directly related to a regulatory role for ROS originated in mitochondria [21]. When autophagy is blocked, this leads to activation the NLRP3 inflammasome, which explains the frequent association of mitochondrial damage with inflammatory diseases [22]. Additionally, autophagy is regulated by the mammalian target of rapamycin (mTOR), which shuts off autophagy in the presence of growth factors and abundant nutrients [23]. In vitro surveys suggest that periodontal pathogens can induce mTOR degradation in epithelial cells [24]. Furthermore, the mTOR activates peroxisome proliferator-activated receptors (PPARs) [25], which have been increasingly recognized as key players in the pathogenesis of metabolic diseases and primary metabolic regulators [26]. In addition, PPARs are in direct relation with the NLRP3 inflammasome [27].

It is well-known that periodontitis shares pathological features with the mentioned systemic pathologies [7]. The evidence appears to indicate that the common link between periodontitis and systemic diseases seems to be inflammation [9]. Although inflammation is a protective physiological response, it can sometimes lead to the destruction of tissue, starting a disease. There are a multitude of molecular routes related with inflammation which have been studied to explain the physiopathological processes taking place in periodontitis. Routes related with inflammasomes and autophagy have been widely investigated [9,19]. Also, there is evidence for a decreased antioxidant capacity and mitochondrial dysfunction in patients with periodontitis [28,29]. In fact, animals studies have shown that antioxidant treatment with coenzyme Q is able to reduce the progression of alveolar bone destruction [30].

Nowadays, the key questions in periodontitis remain unanswered. Why in some patients does the inflammatory response cause a pathology, and why not in another? With the recent knowledge we

have gained, we have begun to look for new keys in this process and to try to obtain an answer within this new family of receptors known as PPARs.

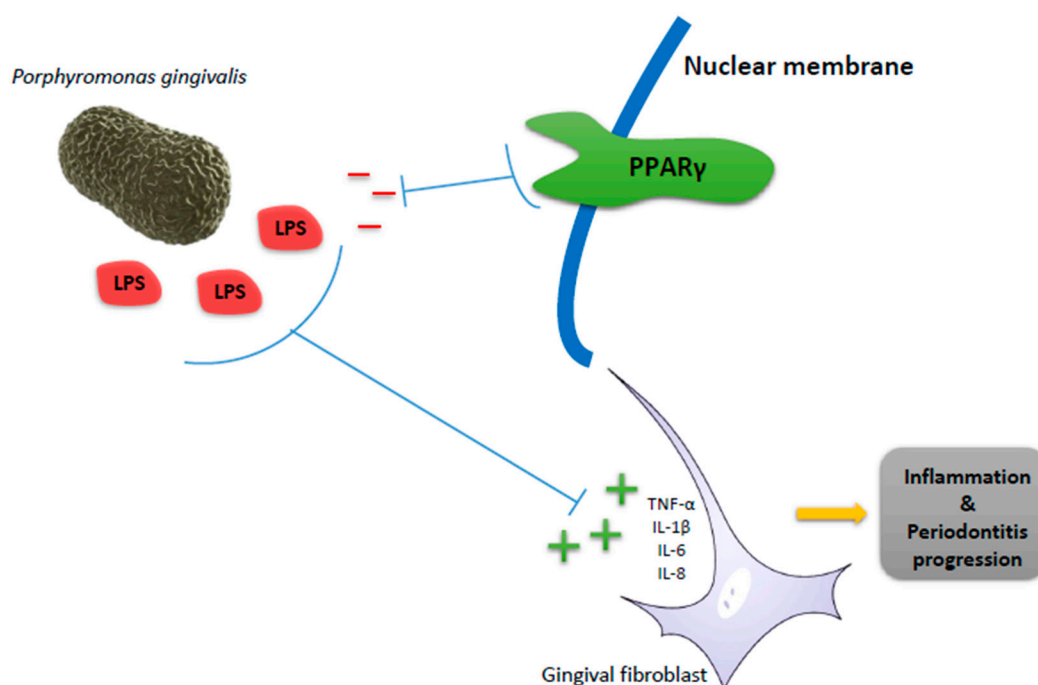
## 2. Peroxisome Proliferator-Activated Receptors (PPARs) in Periodontitis as Inflammatory Disease

PPARs were originally identified in *Xenopus* frogs as receptors that induce the proliferation of peroxisomes in cells. They are a group of nuclear receptor proteins regulating the expression of genes. Three types of PPARs have been identified:  $\alpha$ ,  $\gamma$ ,  $\beta/\delta$ . PPAR $\alpha$  is expressed in liver, kidney, heart, muscle, and adipose tissue. PPAR $\gamma$  is expressed in all tissues. PPAR $\beta/\delta$  is strongly expressed in brain, adipose tissue and skin. They are key players in the regulation of cellular differentiation, development, and metabolism (carbohydrate, lipid, protein), and tumor production [31]. Basically, PPARs are considered as lipid sensors, that according to its availability leads to fatty acid metabolism or lipid storage. Also, PPARs play a major role in inflammation. For instance, they are involved in age-related inflammation, caloric restriction, and longevity [32]. Activation of PPAR $\gamma$  reduces the inflammation and can be activated by some non-steroidal anti-inflammatory and oral antidiabetic drugs. Many of PPAR $\gamma$ 's established anti-inflammatory effects have been shown to occur through innate immune signaling, particularly in monocytes and macrophages, but new data is investigating the role of PPAR $\gamma$  and its ligands (including thiazolidinediones, prostaglandins, and oleanolic acids) in the resolution of inflammation [33]. Genetically modified mice showed that PPAR $\alpha$ -impaired activity results in a modified inflammatory reaction, which causes a temporary delay in epidermal healing [34]. The upregulation of PPAR $\beta/\delta$  expression and endogenous agonist is triggered by tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), a proinflammatory cytokine [35]. The antiinflammatory role of PPAR $\beta/\delta$  has been related to the utilization of agonists in different models of neurodegenerative diseases [36]. In inflammatory bowel diseases, the activation of PPAR $\alpha$  or PPAR $\gamma$  has antiinflammatory effects in the intestine, decreasing the production of inflammatory markers and slows the progression of colitis [37]. PPAR $\alpha$  is the molecular target of the fibrates class of drugs, such as fenofibrate, which act as agonistic ligands of PPAR $\alpha$  [38]. Also, PPAR $\alpha$  ligands can inhibit the expression of various proinflammatory genes, such as interleukin (IL)-6, vascular cell adhesion molecule-1, platelet-activating factor (PAF) receptor and cyclooxygenase (COX)-2 generating prostaglandin E2 (PGE2) and thromboxane B2 (TxB2), in response to cytokine activation [39].

On the whole, PPARs ligands activate PPARs to express their target genes. These agonists can be synthetic molecules, such as anti-hypertriglyceridemia drugs and insulin resistance, or fatty acids and eicosanoids. The first synthetic high-affinity ligand for PPAR $\beta/\delta$  is the GW0742; it can modulate the inflammatory process and could limit the development of periodontitis. It has been demonstrated in rats to result in a substantial reduction of ligature-induced periodontitis, reducing plasma extravasation and the degree of bone resorption. Therefore, it could provide a promising approach for the treatment of periodontitis, reducing plasma extravasation and the bone resorption during periodontitis [40].

At present, therapy with mesenchymal stem cells (MSC) is one of the most promising options to treat some diseases. Periodontal ligament stem cells have emerged as readily available therapeutic sources. They share core MSC properties such as colony formation and multipotential differentiation abilities, and may be ideal for repairing periodontal defects and achieving tooth regeneration with fewer drawbacks. Real-time PCR shows a highly elevated level of PPAR $\gamma$  in these cells [41]. One of the main bacteria involved in periodontitis is *Porphyromonas gingivalis*, and its lipopolysaccharide (LPS) induce the expression of inflammatory cytokines TNF- $\alpha$ , IL-1 $\beta$ , and IL-6 and chemokines IL-8 in human gingival fibroblasts (HGF) [42]. The activation of PPAR $\gamma$  could inhibit LPS-induced inflammatory response [43] (Figure 2). A major metabolite of anthocyanins, the protocatechic acid, inhibit LPS-induced inflammation, suppressed IL-6 and IL-8 production in LPS-stimulated HGF and can be reversed by the PPAR $\gamma$  antagonist GW9662 [44]. The evidence that rosiglitazone (RGZ), a PPAR $\gamma$  agonist, reduces acute and chronic inflammation has been tested in a rat model of periodontitis. It demonstrated a reduction in the development of periodontitis, the presence in the mucogingival tissues with polymorphonuclear cells, the degree of nitrotyrosine formation and the

level of injury in mucogingival tissues. All these findings support an attenuation of the damage produced in experimental periodontitis [45]. The interaction between PPAR $\gamma$  gene polymorphism with periodontal disease has been studied in some papers without a clear conclusion. The association of five polymorphisms (rs10865710, rs2067819, rs3892175, rs1801282, rs3856806) within the PPAR $\gamma$  gene with chronic periodontitis was studied in a group of patients aged from eight to 85 years-old. The results show that none of the haplotypes was significantly different between the periodontitis and healthy controls. Therefore, it could not demonstrate a significant association between PPAR $\gamma$  gene variants and chronic periodontitis [46]. In a group of pregnant women (the average age of the subjects being approximately 31 years-old), it could not be demonstrated an association between the PPAR $\gamma$  Pro12Ala polymorphism and the presence of periodontitis because they did not reach statistical significance [47]. Later the same group of authors showed a significant association with periodontal disease in obese elderly females, indicating a gene–gene or gene–environmental interaction among periodontal disease and each gene polymorphism [48]. Another study made in a group of postmenopausal women with slight periodontitis do not demonstrate any independent associations between the PPAR $\gamma$  Pro12Ala polymorphism and any of the periodontal clinical parameters [49].



**Figure 2.** Anti-inflammatory effect of peroxisome proliferator-activated receptors  $\gamma$  (PPAR $\gamma$ ). A rat model of ligature-induced periodontitis shows the effects of WY14643, a potent peroxisome proliferator activator receptor  $\alpha$  (PPAR $\alpha$ ) agonist. The treatment with WY14643 attenuates the degree of inflammatory response in the gingival tissues. All of these data support that PPAR $\alpha$  has a pernicious role in the development of damage associated with periodontitis in rats [50]. TNF- $\alpha$ , tumor necrosis factor- $\alpha$ ; IL-1 $\beta$ , interleukin 1  $\beta$ ; IL-6, interleukin 6; IL-8, interleukin 8; and LPS, lipopolysaccharide.

### 3. PPARs in Periodontitis as Bone Disease

The balance of bone metabolism is supported by osteoclast-induced bone resorption and osteoblast-induced bone formation. Osteoblastic lineage cells and activated T lymphocytes produce a receptor activator of nuclear factor (NF- $\kappa$ B) ligand (RANKL). It is the essential factor for osteoclast formation, fusion, activation, and survival, resulting in bone resorption and bone loss. RANKL has a specific receptor located on osteoclast and dendritic cells: RANK. The activation of RANK promotes the signaling cascade including stimulation of the c-jun, NF- $\kappa$ B, and protein serine/threonine



kinase (PKB/Akt) pathways. The effects of RANKL are counteracted by osteoprotegerin (OPG) which acts as a soluble neutralizing receptor. The regulation of RANKL and OPG are mediated by cytokines (tumor necrosis factor  $\alpha$ , interleukins 1, 4, 6, 11, and 17), various hormones (glucocorticoids, vitamin D, estrogen), and several mesenchymal transcription factors (such as Cbfa-1, peroxisome proliferator-activated receptor  $\gamma$ , and Indian hedgehog).

Stimulation of PPAR $\gamma$  induces the development of mesenchymal stem cells into adipocytes instead of osteoblasts and results in the decrease of the number of osteoblasts and bone mineral density (BMD). PPAR $\gamma$  gene polymorphisms have been reported to be associated with bone loss and osteoporosis [51]. A study showed that PPAR $\gamma$  also functions as an essential modulator of osteoclast differentiation. Mice with PPAR $\gamma$ -deficiency in their hematopoietic lineage cells are osteopetrotic as a result of impaired c-Fos expression. These data suggest that PPAR $\gamma$  acts as an important regulator of c-Fos expression [52]. Another study demonstrates that the activation of liver X receptor (LXR) attenuated the PPAR $\gamma$  agonist-induced c-Fos expression, and the ectopic expression of LXR suppressed PPAR $\gamma$ -induced Activator protein 1 (AP-1) promoter activity. LXR can be identified as a negative regulator of PPAR-mediated induction of c-Fos and transactivation of AP-1 in the context of osteoclast development. The authors propose that LXR agonists may be of value in the treatment of bone-erosive diseases characterized by increased osteoclast number [53].

Mechanical loading is essential in the maintenance of bone mass and structures around the teeth and bone physiology. Some papers indicate that the mechanical loading of osteoblasts produces the expression of transcription factor PPAR $\gamma$ -1 mRNA. The coordinated synthesis of  $\Delta^{12}\text{PGJ}_2$ , a natural ligand for PPAR $\gamma$ -1, with the increased expression of PPAR $\gamma$ -1, point out that the biomechanical transduction pathways that initially involve the activation of cyclooxygenases may also require the activation of the  $\Delta^{12}\text{PGJ}_2$ -PPAR pathway [54]. These data also suggest that prostaglandin (PGD<sub>2</sub>) biosynthesis induced in bone cells by mechanical stress may be one of the major inducers of osteogenesis. The same group of authors investigated the molecular events produced in osteogenesis secondary to mechanical loading and  $\Delta^{12}\text{PGJ}_2$ , namely the induction of bone morphogenetic proteins and peroxisome proliferator-activated receptor  $\gamma$ -1, a nuclear receptor for  $\Delta^{12}\text{PGJ}_2$ . Mechanical strain increased the mRNA expression of peroxisome proliferator-activated receptor  $\gamma$ -1. In stretched cells, bone peroxisome proliferator-activated receptor  $\gamma$ -1 expression was blocked by cyclooxygenase inhibitors, but improved by exogenous  $\Delta^{12}\text{PGJ}_2$ . These data suggest that the  $\Delta^{12}\text{PGJ}_2$ /peroxisome proliferator-activated receptor  $\gamma$ -1 is essential in osteogenesis [55]. The effect of  $\Delta^{12}\text{PGJ}_2$  on periodontitis inhibited the production of interleukin-6 (IL-6) in osteoblast-like cells MC3T3E-1 incubated with LPS. Data showed that it is an effective inhibitor of the LPS-stimulated IL-6 production, in a dose-dependent manner, through the blockade of NF- $\kappa$ B and the Akt pathway. These data suggest a possible mechanism responsible for its anti-inflammatory action and the use as a therapeutic drug for treating LPS-stimulated periodontal diseases [56].

A relationship between osteoporosis and periodontal conditions has been studied. Systemic bone density and oral infection, analyzed in the presence of a periodontal bacteria such as *Tannerella forsythensis*, was found to modify oral bone loss in post-menopausal women aged <70 years [57]. In a group of 400 postmenopausal women, osteoporosis was significantly associated with severe alveolar crestal bone loss and the prevalence of periodontitis cases [58]. Another paper showed data from 90 women aged 45–70 years; the osteoporotic women presented severe periodontitis with greater gingival inflammation, greater clinical attachment level and greater gingival recession than the women with normal BMD [59]. A group of 179 male and female patients were divided into an osteopenia, and a non-osteopenia group and a follow-up clinical surveys were done by measuring periodontal conditions after 3 years. BMD was related to the number of progressive sites which had additional attachment loss during the observational period [60].

Pathogenic bone resorption occurring in periodontal disease is known to be produced by inflammation, PPAR $\gamma$  activation can downregulate the bone resorption mediated by RANKL-dependent osteoclastogenesis. RGZ, a high-affinity synthetic agonist for PPAR $\gamma$ , influences the

bone resorption induced in rats by experimental periodontitis caused by ligature placement around the tooth. Histological and immunohistochemical analysis showed that the treatment with RGZ decreased bone resorption, alongside reduced RANKL expression, in contrast to those animals treated with empty vehicles. Also, along with these results obtained from *in vivo* experiments, RGZ also suppressed *in vitro* osteoclast differentiation in the presence of RANKL in mouse monocyte/macrophage (MOCP-5) osteoclast precursor cells, and the downregulation of the expression of RANKL-induced TRAP mRNA. So, RGZ might suppress the bone resorption by inhibiting RANKL-mediated osteoclastogenesis obtained during experimental periodontitis in rats [61].

A study investigated the potential associations between the PPAR $\gamma$  Pro12Ala polymorphism, periodontitis and bone mineral density (BMD) in 674 women between the ages of 55 and 74 years-old. The association between BMD or periodontitis and PPAR $\gamma$  Pro12Ala polymorphism was not independent. The common point could be in the modulator role of the polymorphism in this relationship [62].

In dentistry, the treatment with implants involved the bone integration of the titanium surface without any inflammation. Some papers have been proposed to deliver antiinflammatory molecules PPAR $\gamma$  on titanium surfaces using chitosan gold nanoparticles (Ch-GNPs). They evaluated the protein levels of TNF- $\alpha$  and IL-1 $\beta$  in RAW 264.7 cells. Expression of these cytokines was not observed in cells cultured on Ch-GNPs/PPAR $\gamma$ -coated titanium surfaces. But, upregulation of pro-inflammatory cytokines was detected on non-coated and Ch-GNPs/LacZ-coated titanium surfaces. These data suggested that the Ch-GNPs/PPAR $\gamma$  coated titanium surfaces could carry the anti-inflammatory gene and inhibit inflammation [63].

#### 4. PPARs in Periodontitis and Its Related Systemic Diseases

We have shown that periodontitis is related to some systemic diseases such as cardiovascular disease produced by atherosclerosis, diabetes mellitus, and metabolic syndrome. The pathophysiology of these diseases share a common background of low-grade modified inflammation. Periodontitis results from an imbalance of the bacteria and the host inflammatory response that produce an irreversible alveolar bone resorption, leading to tooth loss. Nowadays, the key question is why in some patients, the inflammation is limited to gingivitis as a reversible disease, and in others produces periodontitis with an irreversible alveolar bone loss. One of the possible mechanisms is an impaired inflammatory response. We have proposed that the interrelationship could lie at a cellular level, namely oxidative stress and mitochondrial dysfunction, as a meeting background among such chronic diseases and periodontitis [64]. Also, all the periodontitis-related systemic diseases show in its pathogenic mechanism an altered inflammation associated with an excessive caloric intake and increased or inadequate free fatty acid intakes [65]. Even the origins of susceptibility for these diseases in the adult can be identified to the early life, early-life nutrition plays an essential role in placental and fetal growth, organogenesis, and development [66]. PPARs contribute to the plasticity of the placenta and are under epigenetic control; it has been proposed there is a relationship between environmental factors, such as maternal nutrition, and the epigenetic regulation of PPARs [67]. Glucose and related sugars, lipids, and amino acids are essential nutrients. The control of the above environmental nutrient levels is essential for life; it is controlled by nutrient-sensing mechanisms and pathways. One of them is PPARs; they are nutrient-sensing nuclear receptors with integrators activity in metabolic responses [68].

Metabolic syndrome is defined as the combination of multiple metabolic disorders, including obesity, dyslipidemia, glucose intolerance, inflammation, and hypertension. Low-grade inflammation is a common background between insulin resistance, obesity and type-2 diabetes [65]. In recent years, some studies showed that PPARs could improve several of the metabolic abnormalities associated with the metabolic syndrome [69].

Specific genetic polymorphisms have been associated with a susceptibility to obesity. Expression of these susceptibility genes can be modified by epigenetic and environmental factors. Diet and

nutritional status can affect the transcription factors and microRNAs (miRNAs) regulatory systems. Evidence from *in vitro* studies and from human studies has identified specific miRNAs associated with obesity. miRNAs are small, non-coding RNA molecules that bind and form complexes with mRNA species, namely to the 3'UTR of the mRNA [70]. miRNAs regulate adipocyte development and lipid metabolism. Expression of miR-27 suppresses the translation of peroxisome proliferator-activated receptor- $\gamma$  (PPAR $\gamma$ ) and is an essential early regulator of adipogenesis [71]. miR-143 expression is upregulated 3.3-fold in obese mice, and increases the adipocyte differentiation markers PPAR $\gamma$  and the adipocyte fatty-acid-binding protein [72]. PPAR $\gamma$  also regulates adipogenesis, and its polymorphisms have been suggested to be a risk for obesity. Body mass index (BMI) has been associated with Pro12Ala polymorphism [73]. Also, it was studied whether the Pro12Ala polymorphism of peroxisome proliferator-activated receptor  $\gamma$  (PPAR $\gamma$ ) is related to insulin resistance, obesity, and weight loss and the potential interactions between fat intake and PPAR $\gamma$  polymorphism was explored. Data showed a protective role for the Ala12 allele against insulin resistance, and the interaction between dietary monounsaturated fatty acids (MUFA) and PPAR $\gamma$ 2 for BMI [74]. PPAR $\gamma$  increased the plasma adiponectin concentration and is suggested as a therapeutic tool through the dietary regulation by derivatives of its active compounds, such as vitamins, polyunsaturated fatty acids (PUFA) or carotenoids [75].

The relationship between cardiovascular disease and PPAR has been reported. PPAR $\beta/\delta$  ligands and the overexpression of this nuclear receptor inhibits myocardial inflammatory responses, such as the lipopolysaccharide-mediated production of TNF- $\alpha$ . This had positive effects on animals that had undergone an ischemia/reperfusion injury or cardiac hypertrophy [76]. PPAR $\beta/\delta$  ligands are capable of improving cardiac function, and can even enhance some cardiac diseases such as heart failure, oxidative damage, hypertrophy, ischemia-reperfusion injury, lipotoxic cardiac dysfunction and lipid-induced cardiac inflammation [77].

The pathophysiology of diabetes includes an important inflammatory aspect. PPAR $\alpha$  is essential in the dysregulation of inflammation. PPAR $\alpha$  protein level was examined by immunohistochemistry in type-2 diabetic Goto-Kakizaki rats with induced periodontitis. The results showed the number of gingival fibroblast with PPAR- $\alpha$  protein was low at baseline and then in both, normoglycemic and diabetic rats with ligature-induced periodontitis. When the ligature was removed and the periodontitis solved, the normoglycemic rats presented a threefold increase in the relative expression of PPAR $\alpha$ . On the contrary, no change was observed in the diabetic animals. However, diabetic rats treated with TNF-inhibitor, showed a significant increase, so that the diabetic rats with TNF-inhibitor function similarly to the normal animals. Thus, PPAR $\alpha$  could participate in the resolution of periodontitis through its antiinflammatory properties, but under diabetic conditions, its upregulation is restrained [78].

The role of monocytes and macrophages in diabetic periodontal diseases has been investigated; the results indicated that monocytes and macrophages are important players in periodontal tissue inflammation and destruction in diabetic patients. Monocytes isolated from diabetic patients produce a greater amount of TNF- $\alpha$  and interleukin-1 $\beta$  *in vitro* than do nondiabetic controls [79]. A high glucose content markedly increased lipopolysaccharide-induced inflammatory cytokine and inducible nitric oxide synthase (iNOS) expression in U937 macrophages. These data indicated that hyperglycemia in diabetic patients may boost the inflammatory response leading to increased cytokine release. An antidyslipidemic drug, fenofibrate, increased lipopolysaccharide, induced TNF- $\alpha$  secretion and an antidiabetic drug, pioglitazone; increased lipopolysaccharide-induced TNF- $\alpha$ , interleukin-1 $\beta$ , and interleukin-6 secretion. But, pioglitazone and fenofibrate do not inhibit the stimulatory effect of high glucose and lipopolysaccharide on the secretion of pro-inflammatory cytokines effectively, but only if it suppressed by statin [80].



## 5. Conclusions

In the complex process involving inflammatory diseases such as periodontal disease, cardiovascular disease, diabetes and metabolic syndrome, PPARs could play an important role and give new clues to a better therapeutic approach. PPARs are the molecular targets of a number of marketed drugs. We should use those to prevent or treat periodontal diseases. This is our new challenge.

**Conflicts of Interest:** The authors declare no conflict of interest.

## Abbreviations

PAMPs	Pathogen-Associated Molecular Patterns
DAMPs	Danger-Associated Molecular Patterns
PRRs	Pattern Recognition Receptors
NLRP3	Nod Like Receptors Family, Pyrin Domain Containing 3
ATP	Adenosine 5'-Triphosphate
ROS	Reactive Oxygen Species
mTOR	Mammalian Target of Rapamycin
IL	Interleukin
COX	Cyclooxygenase
MSC	Mesenchymal Stem Cells
HGF	Human Gingival Fibroblasts
RGZ	Rosiglitazone
RANKL	Receptor Activator for Nuclear Factor $\kappa$ B Ligand
NF- $\kappa$ B	nuclear factor kappa-light-chain-enhancer of activated B cells
OPG	Osteoprotegerin
BMD	Bone Mineral Density
Ch-GNPs	Chitosan Gold Nanoparticles
miRNAs	MicroRNAs
BMI	Body Mass Index
MUFA	Monounsaturated Fatty Acids
PUFA	Polyunsaturated Fatty Acids
TNF	Tumor Necrosis Factor

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