# Relationships between Caries Bacteria, Host Responses, and Clinical Signs and Symptoms of Pulpitis

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## Abstract

Knowledge of caries bacteria and the inflammatory responses they elicit in the dental pulp is prerequisite to our understanding of the pathogenesis of pulpitis. Recent advances in immunology and neurophysiology can now explain some of the clinical manifestations of pulpitis. The purpose of this review is twofold. The first purpose is to review the literature of the caries microflora, the host immune responses they elicit, and how they do so. The relationship between both proinflammatory and anti-inflammatory cytokines and pulpitis is discussed. The proinflammatory properties of lipoteichoic acid, which is a common virulence factor among Gram-positive bacteria such as those found among the caries bacteria, are reviewed. The second purpose is to review how bacteria and their metabolites, as well as pulpal immune and inflammatory reactions to them, modify the pain sensation in pulpitis. (J Endod 2007;33: 213-219)

**Key Words** 

Caries bacteria, cytokines, pain, pulpitis

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Caries bacteria are the major cause of pulpal inflammation and infection. The outcome of pulpal insult is a dynamic process that depends on both the invading microorganisms and host responses to them, which include inflammation and immunity. Evidence from experimental pulpitis models has clearly demonstrated that bacterial antigens and/or metabolic by-products can diffuse through dentinal tubules to elicit immune responses in the dental pulp (1–5). Immune complexes and by-products from immune responses, such as extracellular proteolytic enzymes released by phagocytosis, can further aggravate pulpal inflammation (6, 7), making the problem worse. A case in point was a patient suffering from impaired cellular immunity (athymic dysplasia) with IgA deficiency in whom only a mild inflammatory response with little tissue destruction was observed in a caries-exposed pulp (8). Thus an immunopathological mechanism of pulpitis may be operational.

Pulpal pain is usually the first clinical sign of pathology if the insult is not removed to resolve the edema from inflammation. Persistent inflammation in a low-compliance environment such as the dental pulp elicits pain and eventually leads to total pulp destruction (9, 10) and periapical pathosis (11). Inflammation is the outcome of complex interactions among various cell types. Because of the limited scope of this review, the roles of odontoblasts, fibroblasts, nerve fibers, mast cells, and endothelial cells will be reviewed in a future paper. The goals of this review are: (1) to summarize our current knowledge of caries bacteria and their elicited host immune responses and (2) to discuss the correlation between caries bacteria, their role in pulpal immune responses and inflammation, and the symptoms of pulpitis.

# **Caries Pathogens and Caries Progression**

The microflora in dental caries is highly complex and vary between individual lesions. The composition of the dominant groups may depend on diet, saliva, and the chronicity of the lesion. Mutans group streptococci, such as *Streptococcus mutans* and *Streptococcus sobrinus*, and lactobacilli are important in the initiation and progression of caries (12-14). These microorganisms are acidogenic (produce acid) by fermenting dietary carbohydrates, which results in the demineralization of enamel and dentin. They are also aciduric (acid tolerant), which gives them a competitive survival advantage. Among acidogenic bacteria, arginolytic strains of oral streptococci and lactobacilli (bacteria that can produce base from arginine and thus produce both a decline and increase in pH rather than a decline only) are considered less cariogenic. Their ratio to that of nonarginolytic bacteria may determine the level of caries activity (15, 16). Furthermore, oral streptococci and lactobacilli are capable of intratubular invasion by binding to collagen type I in dentin (17, 18). After demineralization the exposed collagen is further degraded by host-derived matrix metalloproteinases that promote the advance of the caries (19).

As the lesion progresses deeper into the dentin, a transition from predominantly facultative, Gram-positive bacteria in shallow caries to deep dentinal caries dominated with lactobacilli and/or anaerobic bacteria takes place (20-22). This transition is probably influenced by a change in the ecosystem (i.e., nutrients,  $O_2$  etc.). The availability of serum-like nutrients diffusing from the pulp into deep caries favors the growth of proteolytic over saccharolytic bacteria (23, 24) as does blood hemin for Prevotella species (25). Streptococci that require salivary glycoproteins and carbohydrates as an energy source are unlikely to thrive without saliva in a deep carious lesion.

We identified two types of deep carious lesions: those with high Lactobacillus and those with low Lactobacillus counts (26). In the low Lactobacillus lesions, a great

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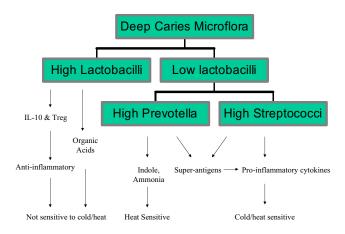


Figure 1. The metabolites of deep caries bacteria and their contributions to pulpal thermal sensitivity.

variation of the dominant microbes was noted and Gram-positive non-Lactobacilli rods, Gram-positive cocci, or *Prevotella intermedia* were dominant in certain samples. Results of recent molecular studies assessing the predominant flora in deep carious dentin support our findings (27, 28). Furthermore, Chhour et al. (29) used real-time polymerase chain reaction (PCR) to assess the total bacterial load of deep carious lesions and categorized the lesions into high-Lactobacillus, mid-Lactobacillus/Prevotella, high-Prevotella, and low-Lactobacillus/ Prevotella lesions. However, neither Lactobacilli nor Prevotella were the dominant isolates reported by Hoshino (21). They observed predominantly anaerobic Gram-positive bacteria and Gram-negative rods in deep dentinal caries, among which *Pseudoramibacter alactolyticus* (was *Eubacterium alactolyticum*) was the predominant isolate.

When caries invade pulpally, the pulpal inflammation manifests itself by pain or hypersensitivity. In the following section, we will focus on the bacterial metabolites that can be the modifiers of pain symptoms associated with deep caries. The relationship between fermentation end products of various bacteria in deep caries and clinical manifestations to thermal tests is presented in Fig. 1.

## Acid, Bacterial By-products, and Pain Modification

Bacterial metabolic by-products and cell wall components can diffuse through dentinal tubules to elicit pulpal inflammation (2, 5, 30-32). Although we have not identified all of the molecules that come through the dentinal tubules, the main metabolites in caries have been examined. Lactic acid is the predominant microbial by-product in active carious dentin (88%), which exhibits a low pH (mean 4.9) (33). Aciduric bacteria such as streptococci and lactobacilli produce more lactate under acidic conditions than at neutral pH (34, 35). In contrast, arrested dentinal caries exhibits an acetate-dominant profile with a higher pH (mean 5.7) (33). Organic acid from bacterial fermentation of carbohydrates, including lactic, acetic, and propionic acids (1 mM to 1 M), not only fail to excite the intratubular A- $\delta$  nerves but also reversibly suppress nerve impulses elicited by other stimuli (36). These findings contradict the action of these acids in other tissues, which is to cause severe pain. Two possible explanations have been proposed. First, the dental pulp may lack the chemosensitive pain fibers found in other tissues (37, 38). Second, increased concentrations of H<sup>+</sup> ions (39, 40)and/or  $Ca^{++}$  ions (41) liberated from dentin decalcified by the acids may decrease the excitability of the dentinal nerve fibers. The direct suppressive effect of acids on intratubular nerves could partially explain why carious teeth with high Lactobacillus counts usually are not sensitive to thermal stimuli (42). This finding also may explain why a higher neural density found beneath caries is not related to a worse pain experience (43).

Among the algogenic (pain producing) molecules produced by bacterial metabolism, ammonia is the most potent pain inducer, followed by urea and indole, which are by-products of amino acid fermentation (44). Many anaerobic bacteria are asaccharolytic (i.e., do not gain energy from conversion of sugars to acidic fermentation products) but are proteolytic, their growth depending on their ability to metabolize proteins or peptides. Asaccharolytic bacteria such as Fusobacterium nucleatum, Prevotella intermedia, and Porphyromonas gingivalis can incorporate and ferment amino acids such as glutamic and aspartic acids into organic acids and ammonia (45, 46). Bacteria associated with deep caries and infected canals, such as Porphyromonas spp, P. intermedia, F. nucleatum, Propionobacterium acnes, and a few isolates of Actinomyces, Peptostreptococci, Bacteroides, and Eubacteria are indole-positive (a test used for bacterial taxonomy that determines the ability of the bacterium to split indole from the amino acid tryptophan) (47). Many studies have demonstrated a close association between pain and the recovery of Prevotella, Porphyromonas, and Fusobacterium from caries (26, 48) and infected canals (49, 50). The algogenic metabolites from anaerobic Gram-negative bacteria in deep caries could partially explain why Bacteroides spp, P. intermedia, and the amount of lipopolysaccharide (LPS) in caries are positively related to heat sensitivity or pain (42, 51). LPS activates the Hageman factor, leading to bradykinin production, a potent pain inducer (52, 53).

#### **Lipoteichoic Acid**

LTA is an amphiphilic molecule consisting of a polyglycerolphosphate with a complex glycolipid group attached (54, 55). It is anchored by hydrophobic forces to the cell membranes of Gram-positive bacteria, including most streptococcal strains (56). LTA is produced in large quantities by cariogenic bacteria when sucrose is available (57) and can be exported extracellularly when bacteria are grown at low pH (58). LTA released extracellularly by Gram-positive, acidogenic bacteria could diffuse pulpally and elicit immune responses.

LPS and LTA activate the innate immune system by similar mechanisms. Both LPS and LTA bind to CD14, activate signaling by Toll-like receptors (TLRs) (59, 60), and induce proinflammatory cytokines such as tumor necrosis factor-alpha (TNF- $\alpha$ ), interleukin-1 (IL-1), interlukin-8 (IL-8), interleukin-12 (IL-12), and anti-inflammatory cytokine interleukin-10 (IL-10) (61, 62). A recent study demonstrated that Bacillus subtilis LTA stimulates odontoblasts by TLR2 and induces the secretion of chemokines (CCL2 and CXCL2). CCL2 attracts immature dendritic cells (DCs) and CXCL2 is angiogenic (63). Lactobacillus LTA induces TNF- $\alpha$  production by TLR2 (64). Furthermore, LTA from S. mutans induced apoptosis of cultured pulp cells (mainly fibroblasts) in vitro, which could contribute to the initiation of and/or progression of pulpitis (65). A recent study demonstrated that TGF- $\beta$  (transforming growth factor) gene expression was down-regulated when odontoblasts were challenged with LTA, which promotes immune defense rather than mineralization (63). A summary of the proinflammatory and antiinflammatory effects of LTA is illustrated in Fig. 2.

Although LTA is much less potent than LPS in inducing proinflammatory cytokine production by macrophages (62), it exhibits a similar potency in the induction of macrophage vascular endothelial cell growth factor (VEGF) expression (66, 67). VEGF can also be produced by LTA-stimulated odontoblast-like cells and pulpal cells (67). VEGF is a potent inducer of angiogenesis and vascular permeability (68, 69). The ability of VEGF to enhance vascular permeability is estimated to be 50,000 times higher than that of histamine (70). Furthermore, VEGF is expressed in dentin matrix and the rate of its release from the matrix after injury closely relates to the healing capability of the pulpal tissue

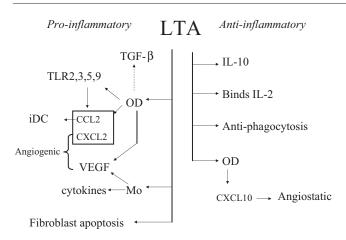


Figure 2. The pro-inflammatory and anti-inflammatory properties of LTA. Negative regulation is expressed by a dashed line (- - - - -). Abbreviations: iDC, immature dendritic cells; Mo, monocytes; OD, odontoblasts; VEGF, vascular endothelial cell growth factor.

(71). A rapid increase in VEGF expression may result in an acute increase in interstitial tissue pressure in the noncompliant pulp space, leading to pulpal necrosis.

LTA also exhibits anti-inflammatory effects including anti-phagocytosis and direct binding to interleukin-2 (IL-2) (72, 73). LTA from hemolytic streptococci inhibits the uptake of streptococci by epithelial cells. LTA binding to IL-2 inhibits the function and measurability of IL-2 (73, 74). An increase of IL-2 titer in irreversible pulpitis was observed by Rauschenberger et al. (75) but not by Anderson et al. (76). This discrepancy could be the result of different amounts of LTA in their samples, which was not measured. Because IL-2 performs many functions critical to the successful elimination of pathogens, LTA release could significantly dampen the immune responses in general. Thus, LTA may not only provide a selective advantage to Gram-positive bacteria but also interfere with ongoing responses to other infectious agents.

#### **Pro-inflammatory Cytokine Induction by Caries Bacteria**

Bacterial antigens induce proinflammatory cytokines including IL-12, IL-1, TNF- $\alpha$ , and interferon-gamma (IFN- $\gamma$ ) (61, 77–79). IL-12 is mainly secreted by DCs and monocytes/macrophages. Its chief biologic function is to stimulate IFN- $\gamma$  production by activated T cells and natural killer (NK) cells. IFN- $\gamma$  in turn activates macrophages to kill phagocytosed microbes. IL-1 and TNF- $\alpha$  are rapidly produced by activated monocytes/macrophages to recruit neutrophils and monocytes to the site of infection. In general, Gram-positive and Gram-negative bacteria are comparable in their IL-1 induction but Gram-positive are more potent IL-12 and TNF- $\alpha$  inducers than Gram-negative bacteria (78, 79).

IL-6 is secreted by various cell types in response to microbes or cytokines such as IL-1 and TNF- $\alpha$  (80–83). IL-6 stimulates hepatocytes to synthesize two major acute-phase proteins: C-reactive protein (CRP), which increases the rate of bacterial phagocytosis, and serum amyloid A (SAA), which influences cell adhesion, migration, proliferation, and aggregation. These proteins also produce a systemic reaction that includes fever, increased erythrocyte sedimentation rate, increased secretion of glucocorticoids, activation of the complement, and clotting cascades (84, 85). IL-6 also stimulates the production of neutrophils from bone marrow in innate immunity (86, 87). IL-6 is involved in adaptive immunity by inducing the permanent differentiation of B-cells into plasma cells that produce antibodies and is therefore considered to be a type-2 cytokine (one that stimulates antibody production) by some researchers (88). IL-6, along with IL-1 $\beta$ , is secreted when pulpal cells

are challenged with peptidoglycan preparations of Gram-positive bacteria (89, 90). Thus, IL-6 may be important in the later stage of pulpitis when the number of B cells increases.

Gram-positive cell walls stimulate the synthesis of TNF- $\alpha$  and IL-6 by human monocytes (91). Peptidoglycan is the major cell wall component of Gram-positive bacteria and a thin layer is also present in Gram-negative bacteria. It has been increasingly recognized as a potent proinflammatory molecule that plays an important role in rheumatoid arthritis and other autoimmune diseases (92–95). Because peptidoglycan is released upon lysis of the cell, its role in the progression of pulpitis needs further investigation.

The LPS of Gram-negative bacterial cells induces potent proinflammatory cytokines (96). Various preparations of Gram-negative bacteria induce dental pulp cells to secrete IL-6, IL-1 $\beta$ , and IL-8 (which recruits neutrophils to the site of inflammation, and thus is also known as the Neutrophil Chemotactic Factor) (81, 97–101). Although pulpal fibroblasts can produce pro-inflammatory cytokines and chemokines in vitro, the majority of IL-1 or IL-8 positive cells in inflamed pulps are immune cells and endothelial cells not associated with fibroblasts (102–104). The suppression of cytokine expression in vivo may arise from immunomodulation by other cytokines (105).

*S. mutans* induces peripheral blood mononuclear cells to produce high titers of proinflammatory cytokines including IFN- $\gamma$ , IL-12 (77), TNF- $\alpha$  (106), and chemokines including IL-8 and MCP-1 (107). Cell wall molecules of *S. mutans*, such as proteins of the *I*/II family and the serotype f antigen (a polysaccharide rhamnose-glucose polysaccharide covalently bound to bacterial cell wall peptidoglycans that facilitates the adherence of *S. mutans* to hard surfaces) are responsible for the induction of these inflammatory cytokines (108). Cell-free supernatant fluids of viridans Streptococci induce IL-1, IL-6, IL-8, and TNF- $\alpha$ (109, 110). Extracellular products of *Streptococcus mitis* and *S. sanguinis* also induce proinflammatory cytokines (109, 111). The inflammatory mediators induced by these proinflammatory cytokines sensitize C fibers in the dental pulp (112–116). These findings may partially explain why some deep caries loaded with oral streptococci exhibit prolonged pain to heat testing (42).

Other bacterial components such as cell surface polysaccharides, heat shock proteins, and extracellular products (such as proteases and exotoxins) also induce pro-inflammatory cytokines (117). Superantigens, which are potent cytokine inducers that do not require antigen presentation, have been associated with certain strains of Streptococcus mitis, P. intermedia, and periodontal pathogens (109, 118–120). Proinflammatory cytokines play a crucial role in tissue destruction and the pathogenesis of many diseases (121-124). Anti-inflammatory cytokine treatment has proved beneficial in arthritis, inflammatory bowel disease, and periodontal disease (125-128). Similarly, proinflammatory cytokine induction by caries bacteria is an important virulence factor in pulpitis. Patients with compromised proinflammatory cytokine responses may react to caries invasion with little inflammation, whereas severe inflammation and pulpal tissue destruction may result in patients with a hyper-response of proinflammatory cytokines to bacterial challenge. Local application of anti-inflammatory cytokine reagents can therefore be therapeutic in early pulpitis by reducing tissue destruction as well as pain.

## Anti-inflammatory Cytokine (IL-10) Induction by Caries Bacteria

Lactobacilli are the dominant flora in certain deep carious lesions and are negatively associated with thermal sensitivity (26). Clinically, we sometimes encounter teeth with vital pulps and deep caries but no obvious thermal sensitivity. This clinical finding may be explained partly by the direct suppressive effect of the organic acids produced by these

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bacteria and partly by their induction of anti-inflammatory cytokine IL-10 and regulatory T cells (Treg) (129-131). L. casei, one of the Lactobacillus species most frequently recovered from deep caries, induced a significantly higher IL-10 titer than that of other caries bacteria at a low concentration  $(10^5 \text{ per mL})$  (132). Certain Lactobacilli such as L. casei and L. paracasei bind to DC-specific intercellular adhesion molecule 3-grabbing non-integrin (DC-SIGN) on immature DCs and induce Treg and IL-10 production (129-131). Furthermore, Lactobacilli appear to generate tolerogenic DCs, a phenotype characterized by increased co-stimulatory marker expression but low production of proinflammatory cytokines (133). Such tolerogenic DCs may contribute to the production of Treg in vivo (134, 135). This immune suppressive effect of *L. casei* helps explain the significantly reduced IFN- $\gamma$  titer that we observed when peripheral blood mononuclear cells were challenged with S. mutans and L. casei together (unpublished data). P. alactolyticus, another representative isolate from caries, exhibits a strong type-2 cytokine profile, inducing more IL-10 than IFN- $\gamma$ . The IL-10 induced by these bacteria and Treg can contribute to the significant elevation of IL-10 mRNA in pulps beneath deep caries (77).

# **Other Virulence Factors of Caries Pathogens**

Gram-positive caries bacteria are capable of activating complement pathways. Based on experimental pulpitis studies, C3a and C5a generated from complement activation were thought to be important virulence factors of cariogenic bacteria that elicit neutrophil accumulation in the Arthus reaction (2, 3, 5, 30). Recent studies, however, showed that the presence of an intact complement cascade is neither necessary nor sufficient to trigger or propagate the Arthus reaction (136, 137). Thus, tissue damage from direct complement activation by Gram-positive bacteria in shallow caries may be minimal.

*S. mutans* and *L. casei* are cytotoxic and are capable of causing total pulpal necrosis of rat teeth (138, 139). *S. sanguinis* and *Entero*-

*coccus faecalis* were the most cytotoxic isolates of the commonly isolated endodontic pathogens to fibroblasts and macrophages (140, 141). The molecular mechanism of the cytotoxicity of *S. sanguinis* is not yet understood. Thorough reviews of the virulence factors of *E. faecalis* are available (142, 143). Virulence factors of other Gramnegative bacteria associated with infected canals are reviewed by Baumgartner (144).

#### **Remaining Dentin Thickness and Tubular Permeability**

Remaining dentin thickness and tubular permeability are the most important determining factors of the pulpal inflammatory response (145–147). Although bacteria or their cell-wall components such as LPS are capable of passing through tubules to induce inflammatory responses in the dental pulp (2, 4, 5, 30), the thickness of dentin can greatly reduce the concentration of bacterial proteins and the amount of LPS that reaches the pulp (31, 148, 149). We infiltrated commercial LTA preparations from S. mutans through dentinal blocks using fluid filtration and observed that the dentin permeability decreased with time (Fig. 3), probably as a result of LTA binding to the dentinal walls (150). An increase of inflammatory cells in pulps of teeth with pulpitis was observed only when the caries front was <1.5 mm from the pulp (151, 152). McLachlan et al. (153) observed less expression of PMN-associated S100 family genes (which code for S-100 proteins, calcium-binding proteins, some of which have been shown to have intracellular and extracellular functions associated with inflammation) in inflamed pulps when the remaining dentin thickness was >2 mm. It appears that 2 mm of sound dentin provides a safeguard for a speedy recovery of the dental pulp to health.

# **Future Directions**

Caries are composed of complex and dynamic flora, all of which the dental pulp is exposed to at one time or another. Studies of individ-

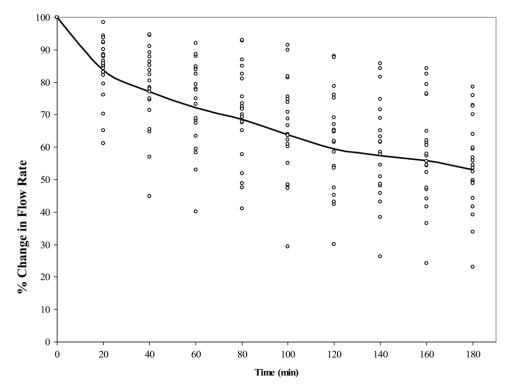


Figure 3. LTA reduces fluid flow. Human dentinal disks (2.6-1.3 mm in thickness) were infiltrated with LTA from *S. mutans* (50 µg/ml, Sigma) for 180 minutes according to the method described previously (157). Each dot represents one data point of one sample block at the specific time point. Percentage changes of flow rate of 23 blocks were plotted over time. There was a significant decrease of flow rate with time.

ual cultivable caries bacteria and their cell wall components have been undertaken to better understand the pathogenesis of pulpitis. Unfortunately, roughly 50% of caries bacteria are not cultivable (154), and we have not yet identified the molecules in shallow caries that diffuse through the dentinal tubules. Molecular approaches to characterize these chemicals are much needed. Analysis of a large sample population of carious dentin and the infecting microbes using a high-throughput gene expression technique such as microarrays would provide a more comprehensive understanding of the multitude of genes of importance in caries. Therefore, conclusions drawn from current studies may need to be modified or abandoned as more data are collected through future research.

Caries bacteria elicit various concentrations of both proinflammatory and anti-inflammatory cytokines. These bacteria have usually been grown and studied in a planktonic state. Gene and protein expressions, however, are clearly different when comparing bacteria grown in the planktonic state to those in a biofilm community (155–157). Bacteria grown in a biofilm environment, closely simulating the in vivo carious lesion, would provide valuable information about virulence factors involved in the caries invasion. Furthermore, dynamics of the polymicrobial caries infection and its evolution over time have not been addressed with these in vitro studies. Thus, a more realistic in vivo and/or in vitro caries model is needed to critically examine the virulence factors of caries bacteria.

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