

SUPRAGINGIVAL CALCULUS: FORMATION AND CONTROL

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ABSTRACT: Dental calculus is composed of inorganic components and organic matrix. Brushite, dicalcium phosphate dihydrate, octacalcium phosphate, hydroxyapatite, and whitlockite form the mineral part of dental calculus. Salivary proteins selectively adsorb on the tooth surface to form an acquired pellicle. It is followed by the adherence of various oral micro-organisms. Fimbriae, flagella, and some other surface proteins are essential for microbial adherence. Microbial co-aggregation and co-adhesion enable some micro-organisms, which are incapable of adhering, to adhere to the pellicle-coated tooth surface. Once organisms attach to the tooth surface, new genes could be expressed so that mature dental plaque can form and biofilm bacteria assume increased resistance to antimicrobial agents. Supersaturation of saliva and plaque fluid with respect to calcium phosphates is the driving force for plaque mineralization. Both salivary flow rate and plaque pH appear to influence the saturation degree of calcium phosphates. Acidic phospholipids and specific proteolipids present in cell membranes play a key role in microbial mineralization. The roles of crystal growth inhibitors, promoters, and organic acids in calculus formation are discussed. Application of biofilm culture systems in plaque mineralization is concisely reviewed. Anti-calculus agents used—centering on triclosan plus polyvinyl methyl ether/maleic acid copolymer, pyrophosphate plus polyvinyl methyl ether/maleic acid copolymer, and zinc ion—in commercial dentifrices are also discussed in this paper.

Key words. Dental calculus, plaque mineralization, microbial mineralization, organic acid, anti-calculus agent.

(I) Composition and Structure of Dental Calculus

Dental calculus is primarily composed of mineral as well as inorganic and organic components. Supragingival and subgingival calculus contain 37% and 58% mineral content by volume, respectively (Friskopp and Isacson, 1984). The matrix of supragingival calculus constitutes 15.7% of the calculus dry weight and contains 54.9% protein and 10.2% lipid. Of the total lipids, 61.8% are neutral lipids, including a high content of free fatty acids and a smaller amount of triglycerides. Glycolipids account for 28% of the total lipids and are composed of 17.2% simple glycosphingolipids, mainly lactosyl- and glucosylceramine, and of 82.8% neutral and sulfated glyceroglucolipids. Phospholipids, representing 10.2% of the total lipid, contain 34.2% phosphatidylethanolamine, 25.5% diphosphatidylglycerol, 2.3% phosphatidylinositol, and 1.7% phosphatidylserine. Phosphatidylinositol and phosphatidylserine are two important classes of acidic phospholipids but are only minor phospholipid components of bacterial cell membrane (Goldfine, 1972), which are enriched in phosphatidylethanolamine and diphosphatidylglycerol (neutral phospholipids) (Bishop, 1971). Notably, dental calculus contains both total phospholipids and acidic phospholipids in much higher concentrations than parotid saliva (Mandel and Eisenstein, 1969; Slomiany *et al.*, 1981). In addition, the concentration of phospholipids in the saliva of heavy calculus formers is significantly higher than that of light calculus formers (Slomiany *et al.*, 1981). These findings suggest that phospholipids play an important role in calculus formation (discussed in a later section).

Dental calculus is always covered by a soft and loose layer of micro-organisms. On supragingival calculus, this layer is dominated by filamentous micro-organisms. The filaments are approximately perpendicular to and in direct contact with the

underlying dense calculus. By contrast, the mixture of cocci, rods, and filaments covering subgingival calculus has no distinct pattern of orientation (Friskopp and Hammarström, 1980). After treatment with sodium hypochlorite, however, the filaments on the surface of supragingival calculus disappear and, in scanning electron microscopy (SEM), the supragingival calculus has a honeycomb appearance (Friskopp and Hammarström, 1980). Sodium-hypochlorite-washed subgingival calculus shows holes in the surfaces which are less regular than those in the surfaces of the supragingival calculus (Friskopp and Hammarström, 1980). Beneath the layer of micro-organisms are calcifying areas, which appear to be laminated due to alternating dark- and light-staining bands (Mandel *et al.*, 1957). The properties of dental calculus were further investigated by Friskopp (1983), who reported that the calcifying areas of supragingival calculus are heterogeneous due to the presence of non-calcified layers. These non-calcified layers are usually irregular, resembling bands of cavities partially separated by calcified materials. Additionally, it has been demonstrated that plaque mineralization occurs in numerous individual foci (Mandel *et al.*, 1957; Friskopp, 1983). Mineral deposits of dental calculus are present both between and within micro-organisms, depending upon deposit age (Donald, 1997); that is, mineral initially deposits in the matrix of plaque, and gradually some plaque micro-organisms become calcified with increasing age of calculus.

It has been demonstrated that octacalcium phosphate $[\text{Ca}_8(\text{PO}_4)_4(\text{HPO}_4)_2\cdot 5\text{H}_2\text{O}]$ (OCP), hydroxyapatite $[\text{Ca}_{10}(\text{PO}_4)_6(\text{OH})_2]$ (HAP), and β -tricalcium phosphate or whitlockite $[\text{Ca}_{10}(\text{HPO}_4)(\text{PO}_4)_6]$ (WHT) form the inorganic part of both supragingival and subgingival calculus. Brushite $[\text{CaHPO}_4\cdot 2\text{H}_2\text{O}]$ (DCPD) is present only in the early-stage supragingival calculus (Rowles, 1964).

Of the calcium phosphate crystals mentioned above, only WHT contains magnesium, which may substitute for part of the calcium in WHT (Friskopp and Isacsson, 1984). Supragingival and subgingival calculus differ from each other with respect to the principal crystal constituents and inorganic elemental components; that is, the WHT-to-HAP (Jensen and Danø, 1954) and calcium-to-phosphorus ratios (Little and Hazen, 1964) in supragingival calculus are lower than those in subgingival calculus.

(II) The Distribution of Dental Calculus in the Dentition

It is well-known that the greatest amount of supragingival calculus is present on the lingual surfaces of mandibular anterior teeth and decreases toward the third molars (Parfitt, 1959). In the maxilla, supragingival calculus frequently forms on the buccal surfaces of the first molars (Parfitt, 1959). In both the mandible and maxilla, these are sites that are close to the orifices of salivary ducts.

Subgingival calculus also exhibits site-specificity, although it is much less apparent than that of its supragingival counterpart (Corbett and Dawes, 1998). It has been reported that the levels of subgingival calculus are significantly higher on the lingual than on the buccal surfaces. For the lingual surfaces of the teeth, the lower first molars have the most subgingival calculus. For the buccal surfaces of the teeth, the mandibular anterior teeth and maxillary molar teeth have the greatest amount of subgingival calculus.

For the mouth as a whole, significant associations ($P < 0.001$) have been demonstrated between the levels of supra- and subgingival calculus, between the levels of supragingival calculus on buccal and lingual surfaces, and between the levels of subgingival calculus on buccal and lingual surfaces. For individual teeth, however, only a few teeth exhibit the above associations (Corbett and Dawes, 1998).

Saliva is closely associated with the distribution pattern of supragingival calculus. As Collins and Dawes (1987) have suggested, saliva is present in the mouth not as a bulk volume but as a mobile film of 0.1 mm average thickness. For unstimulated salivary flow, the velocity of the film over tooth surfaces is estimated to vary between 0.8 and 8 mm/min, depending on different oral regions and status of saliva; for stimulated salivary flow, the velocity is from 1.3 to about 350 mm/min (Dawes *et al.*, 1989). The lowest film velocity of 0.8 to 1.3 mm/min occurs on the facial surfaces of the upper incisors, while the highest salivary film velocities are observed on the lingual surfaces of teeth (Dawes *et al.*, 1989). Based upon these results, it can be concluded that saliva is neither evenly mixed nor well-distributed in the mouth. The site-specificity distribution of calculus can be therefore explained as follows. Parotid and submandibular saliva is supersaturated with respect to calcium phosphates (Lagerlöf, 1983) and is in contact with plaque on the lingual surfaces of the lower incisors and the facial surfaces of the upper molars. As discussed in a later section on urea metabolism, the abundant supply of urea from the major salivary gland secretions tends to increase the pH in plaque (Sissons *et al.*, 1994a) and promote calcium phosphate precipitation. Furthermore, the high salivary film velocity promotes clearance of salivary sugar and acid from plaque and leads to a higher resting plaque pH. More supragingival calculus thus occurs in the regions with the most exposure to saliva. On the facial sur-

faces of upper anterior incisors, the salivary film velocity is low so that the supply of calcium phosphates and urea is limited and sugar and acid easily accumulate in plaque. In summary, calculus deposition tends to occur where plaque is exposed to a salivary film of high velocity.

(III) Mechanism of Plaque and Microbial Mineralization

Following tooth eruption or a dental prophylaxis, salivary proteins rapidly and selectively adsorb onto the enamel surface to form an acquired enamel pellicle. It is followed by the adherence of various oral micro-organisms. Gram-positive coccoidal organisms are the first settlers to adhere to the formed enamel pellicle, and subsequently, filamentous bacteria gradually dominate the maturing plaque biofilm (Scheie, 1994).

Plaque absorbs calcium and phosphate from saliva for the formation of supragingival calculus and from crevicular fluid for the formation of subgingival calculus. Calcium phosphate supersaturation, certain membrane-associated components, and the degradation of nucleation inhibitors are required for initial mineralization of plaque and bacteria. Calculus formation begins with the deposition of kinetically favored precursor phases of calcium phosphate, OCP and DCPD, which are gradually hydrolyzed and transformed into less soluble HAP and WHT mineral phases (Rowles, 1964).

(A) ADSORPTION OF SALIVARY PROTEINS

Three types of time-dependent adsorption of salivary proteins have been observed (Lamkin *et al.*, 1996). Some salivary proteins, such as proline-rich protein-3 (PRP-3), PRP-4, and statherin, adsorb very fast onto HAP, whereas the binding rate of α -amylase, glycosylated proline-rich protein (PRG), and cystatins is slow. The third type of protein adsorption can be seen in the case of PRP-1, PRP-2, and histatins. It is characterized by a two-step process—a rapid adsorption (the direct binding of proteins to HAP), followed by a slow adsorption (protein-protein interactions). Studies on the adsorption of acidic PRPs and statherins appear frequently in the literature. Acidic PRPs and statherins possess polarity because of their highly charged amino-terminal domains, which are responsible for the adsorption of these proteins onto a tooth surface. The phosphoserine residues in the charged regions are believed to be essential for the electrostatic interactions (Raj *et al.*, 1992).

As on the tooth surface, acquired pellicle forms on an implant surface when the metal surface initially comes into contact with tissues (Baier, 1982). The acquired pellicle on an implant surface attracts and harbors various oral bacteria, which may account for the occurrence of implantitis. Notably, the adsorption of proteins does not occur on the surface of pure titanium (Ti). It is demonstrated that a 5- to 6-nm oxide layer composed of TiO_2 forms on the surface of Ti when the implant surface is exposed to air. The physicochemical characteristics of TiO_2 are quite different from those of pure Ti; at physiological pH, the TiO_2 layer carries net negative charges, which enable the TiO_2 layer to bind cations like Ca^{2+} (Abe, 1982). The bound Ca^{2+} makes the surface of an implant positively charged and, consequently, attracts the high-weight molecules carrying negative charges, notably proteins. Ti pellicle differs from enamel pellicle. It has been found that although Ti pellicle contains high-molecular-weight mucin and α -amylase that are also present in enamel pellicle, it lacks cystatins and low-molecular-weight mucin (Fisher *et al.*, 1987).

Three types of protein adsorption onto implant surfaces have been proposed (Williams and Williams, 1988). Protein adsorption onto the surfaces of Ti, Al, Mo, Ni, Ta, and Al₂O₃ implants is slow and will stop a few hours later when a single layer of protein molecules forms on the implant surfaces. The adsorption of proteins onto V and TiO₂ is also slow but continuous. The continuous adsorption can be explained by the possibility that after a single layer of protein molecules forms on the implant surfaces, additional proteins adhere to the attached ones. If the protein adsorption is not only continuous but also fast, it is likely that multi-layers of proteins will form on the implant surfaces.

(B) MICROBIAL ADHERENCE

As Van Loosdrecht *et al.* (1990) and Bollen *et al.* (1996) have described, the microbial adhesion to solid surfaces (such as tooth surfaces and various implant surfaces) may proceed as a four-stage sequence. The first stage is the initial approach of bacteria to a surface where random contact, such as Brownian motions and liquid flow, or active movement of micro-organisms may occur. The attractive van der Waals' forces and repulsive electrostatic forces are responsible for the second stage of microbial adhesion, which is a reversible process. The firm attachment of micro-organisms, "the third stage", is irreversible and is followed by the fourth stage of adhesion, *i.e.*, bacterial colonization.

(1) Surface proteins and bacterial adherence

Appendages, such as fimbriae or pili, emerging from the cell surfaces of many micro-organisms, are involved in the bacterial adherence to pellicle-coated tooth surfaces. The extension and retraction of *Pseudomonas aeruginosa* type IV fimbriae (pili) have been reported to produce forces that propel bacteria across a surface. This phenomenon is referred to as twitching motility (Darzins, 1994). *P. aeruginosa* strains with mutations in *pilB*, *pilC*, and *pilY1*, the genes required for the synthesis of type IV fimbriae (pili) (Nunn *et al.*, 1990), form a monolayer of cells only on an abiotic surface and are incapable of forming microcolonies. Fimbriae may also serve as adhesins for various receptors in the bacterial adhesion process. For example, *P. gingivalis* fimbriae can interact with human salivary components, such as PRPs and statherins (Amano *et al.*, 1996). There exist three or more binding sites in fimbriin proteins of bacteria, and the combination of all these binding sites would be essential for stable binding of salivary proteins (Amano *et al.*, 1996).

Flagellum, another microbial surface protein, is also involved in microbial adherence. It was found that few if any strains carrying a mutation in *flaK*, the gene required for the synthesis of flagellum, attach to the abiotic surface, indicating a role of flagella and/or motility in the initial cell-to-surface interactions (O'Toole and Kolter, 1998).

Flagella may have four different functions. First, flagella may be involved in chemotaxis and enable planktonic bacteria to swim toward nutrients associated with a surface or signals generated by cells attached to a surface. Second, flagella-mediated motility may overcome repulsive forces at the medium-surface interface, to allow the bacteria to reach the surface. Third, flagella-mediated motility may be required for the bacteria within a developing biofilm to move along the surface, thus facilitating growth and spread of the biofilm. Finally, flagella *per se* may be required for biofilm formation

by adhering to an abiotic surface (Pratt and Kolter, 1998). In an *Escherichia coli* model system, mutations that inhibit each of these aspects of flagella function were used to determine which aspect was critical for biofilm formation. The results showed that motile cells that were non-chemotactic formed biofilm indistinguishable from their wild-type counterpart. In contrast, cells either lacking flagella or possessing paralyzed flagella were severely defective in biofilm formation (Pratt and Kolter, 1998). It is therefore evident that chemotaxis is dispensable for biofilm formation, whereas flagella-mediated motility is critical for this process.

Collectively, flagella-mediated motility enable the bacteria to reach a distance short enough from a surface by overcoming the electrostatic interface repulsion so that fimbriae (pili) can push them onto the surface *via* adhesin-receptor interactions and/or twitching motility. The attached bacteria can move along the surface propelled again by the flagella-generated force.

In addition to fimbriae and flagella, many other surface proteins are also involved in bacterial adherence. *Streptococcus mutans* surface proteins have been most widely investigated. P1, a 117-kd and a 127-kd protein, and two glucosyltransferases (GTFase) appear to inhibit the adherence of *S. mutans* WD9463A (c) competitively to an acquired-pellicle-coated surface (Zhan *et al.*, 1998). These results indicate that the above surface proteins serve as adhesins for specific receptors on acquired pellicle. In contrast, Zhan *et al.* (1998) found that another GTFase improved the bacterial adherence, indicating that this GTFase may adsorb onto the HAP surface first and then become the receptor for *S. mutans*. An alternative possibility is that it promotes bacterial adherence by altering the properties of acquired pellicle (Zhan *et al.*, 1998). Based upon these findings, it can be concluded that a surface protein contributes to the bacterial adherence by its specific mechanism.

(2) Microbial co-aggregation and coadhesion

Oral bacteria tend to associate with one another. In suspension, the association between different oral bacteria is termed "co-aggregation", while it is referred to as co-adhesion if one partner of the pair is attached to a surface. Microbial co-adhesion would be of great importance for the colonization of oral micro-organisms on tooth surfaces. For instance, *Streptococcus sanguis* strains facilitate the adherence of *Actinomyces naeslundii*, which *per se* possess poor adhesive ability (Ganeshkumar *et al.*, 1991).

Lectin-sugar interaction is involved in co-aggregation of bacteria, which can be inhibited by sugars (McIntire *et al.*, 1978). Lectin-sugar interaction also contributes to co-aggregation between yeast and bacteria (Ganeshkumar *et al.*, 1991); the sugars in the mannoprotein-containing capsule surrounding the yeast cell may associate with bacteria and human epithelial cells by a lectin-like interaction.

Adhesin-receptor interaction also has a role in microbial co-aggregation. The adhesin molecule on the surface of one partner cell and the receptor molecule on the surface of the other partner cell have to be complementary for a specific binding interaction to occur (Simmonds *et al.*, 2000). Numerous genes encoding the adhesin molecules have been cloned and sequenced, and the encoded proteins have been investigated. For instance, Ssp, a major streptococcal surface protein, mediate adhesion to *Porphyromonas gingivalis* *via* a C-

terminal functional domain (Brooks *et al.*, 1997).

Co-aggregation between some pairs appears to be cation-dependent. For example, calcium ion is required for co-aggregation reactions between actinomycete (such as *Actinomyces viscosus* and *A. naeslundii*) and streptococcal cells (such as *S. sanguis* and *Streptococcus mitis*) (Cisar *et al.*, 1979). Since the surfaces of most bacteria are negatively charged, it is not surprising that electrostatic repulsion exists between bacteria. Cations may therefore make the bacterial zeta potentials less negative, and consequently the weakened electrostatic repulsion allows co-aggregation to occur. Co-aggregation between bacteria can also be non-cation-dependent, such as the co-aggregation of *Prevotella intermedia* and *Prevotella nigrescens* with *A. naeslundii* (Cookson *et al.*, 1995), which can be explained by surface thermodynamic analysis based on interfacial free energies.

In addition to cations, extracellular vesicles also have a role in bacterial co-aggregation (Grenier and Mayrand, 1987). *P. gingivalis*-released vesicles, which have been reported to agglutinate sheep erythrocytes strongly, can promote the co-aggregation of *Eubacterium saburreum* and *Capnocytophaga ochracea*, which otherwise cannot co-aggregate. Electron microscopy reveals that these vesicles act as a physical link or bridge between the two bacterial cells. The vesicle-mediated co-aggregation is unaffected by variation in pH from 4.5 to 8.5, the presence of sugars, uronic acids at a concentration of 100 mM, or EDTA at a concentration of 10 mM. L-arginine and L-lysine, however, are able to inhibit these co-aggregations mediated by vesicles (Grenier and Mayrand, 1987).

Non-specific interaction between micro-organisms is another mechanism of co-aggregation and co-adhesion. For instance, cell-surface lipoproteins are responsible for the non-specific adhesive interaction of *Streptococcus gordonii* with other bacterial species (Jenkinson, 1992).

(3) New behavior pattern of attached bacteria

Once attached to a surface, bacteria stop dividing for hours, during which time genetic changes occur in the planktonic cells, and a biofilm phenotype comes into existence (Rice *et al.*, 2000). With the passage of time, a process termed cell-density-dependent gene expression is required for the formation of more mature biofilm. In Gram-positive bacteria, cell-density-dependent gene expression proceeds as follows. With increasing numbers of bacteria, the pheromone (signal molecules) accumulates up to a critical threshold. The signal is received by the sensing component of a two-component signal transduction system (Kleerebezem *et al.*, 1997). The second component of the transduction system is a cognate response regulator, which is activated by the sensing component and initiates the expression of target genes (Morrison, 1997). Take *S. gordonii* as an example. ComD, an autophosphorylating histidine kinase, is the sensing component that acts as a receptor for Com C, the pheromone that induces competence in a bacterial population. ComE serves as the response regulator, which receives the phosphoryl group from Com D and then binds to specific promoter regions of appropriate target genes (Morrison, 1997). When the level of the molecule approaches a critical point, a transcriptional activator is triggered and expression of the target gene occurs, which is followed by a new behavior pattern of oral bacteria. In the process of cell-density-dependent gene expression, the required concentration of signal molecules depends on the critical level of bacterial population density. The sensing process is therefore

referred to as a quorum sensing system (Pulcini, 2001).

While ComD acts as a signal molecule for Gram-positive bacteria, N-acyl homoserine lactone is involved in the cell-to-cell signaling system as a signal molecule for most Gram-negative bacteria (Pulcini, 2001). Many studies have been conducted to explore the cell-density-dependent gene expression of Gram-negative *P. aeruginosa*. There are at least two signals in *P. aeruginosa* serving the cell-to-cell signaling system, *lasR-lasI* and *rhlR-rhlI* (*vsmR-vsmI*) systems (Davies *et al.*, 1998). The *las I* gene product directs the synthesis of a diffusible extracellular signal, N-(3-oxododecanoyl)-L-homoserine lactone (3OC₁₂-HSL). When 3OC₁₂-HSL reaches a critical level, the *lasR* product activates the *LasI* and *rhlR-rhlI* system. The *rhlI* product directs the synthesis of another signal molecule, N-butyl-L-homoserine lactone. A sufficient level of N-butyl-L-homoserine lactone is required by the *rhlR* gene product for the expression of target gene. Research has shown that *lasI* is essential for biofilm differentiation, resistance to antimicrobial agents, and secretion of an extracellular polysaccharide (EPS) matrix that differs from the planktonic cell matrix (Davies *et al.*, 1998).

Biofilm bacteria exhibit stronger resistance to biocides and antimicrobial agents than does their planktonic counterpart. Four hypotheses may help to explain the increased resistance. First, the exopolysaccharide of the biofilm matrix may inhibit the diffusion of biocides and antimicrobial agents. Second, the physiological differences among biofilm bacteria leave only part of biofilm bacteria susceptible to growth-dependent antibiotics. Third, the genetic changes occurring in transition from planktonic to biofilm bacteria make the biofilm bacteria insensitive to various biocides and antimicrobial agents (Pulcini, 2001). Finally, the low level of metabolisms due to nutrient limitation may also contribute to the resistance to antimicrobial agents by biofilm bacteria (Chen, 2001).

(4) Microbial adherence to implant surfaces

Microbial adherence to Ti implant surfaces has been investigated in various studies (Nakazato *et al.*, 1989; Edgerton *et al.*, 1996). Since Ti pellicle differs from enamel pellicle with regard to protein composition, it is understandable that the microbial adherence and the subsequent microbial colonization on a Ti surface may be different from that on an enamel surface (Edgerton *et al.*, 1996). Indeed, there is evidence that Ti implants harbor less mature plaque, which is composed of more coccoid cells and fewer motile rods, as compared with natural teeth (Quirynen and Listgarten, 1990). In addition, Nakou *et al.* (1987) found six species of oral bacteria unique to dental implants. Microbial adherence to an implant surface can be promoted by an increase in the roughness of the implant surface. Higher levels of microbial adherence have been found on rough Ti surfaces than on smooth and polished surfaces in *in vivo* studies (Nakazato *et al.*, 1989).

(C) THE DRIVING FORCE FOR PLAQUE MINERALIZATION

Calcium and phosphate are two salivary ions which are "raw materials" for dental calculus formation. Theoretically, supersaturation of saliva, especially plaque fluid, with respect to calcium phosphate salts is the driving force for dental plaque mineralization. Ion product (Ip) and solubility product (Ksp) are important for estimation of saturation degree (SD) with

respect to a given salt. If $I_p > K_{sp}$, the fluid is supersaturated and precipitation of the salt can occur; if $I_p = K_{sp}$, the fluid is just saturated with respect to the salt, and if $I_p < K_{sp}$, the salt tends to dissolve, since the fluid is not saturated with respect to the salt (Dawes, 1998). As a matter of fact, parotid and submandibular saliva is generally supersaturated with respect to HAP, OCP, and WHT (Lagerlöf, 1983). Submandibular secretions are even more highly supersaturated with respect to HAP than are parotid secretions. Like saliva, rested and starved plaque fluids are also supersaturated with respect to HAP and other calcium phosphate phases (Carey *et al.*, 1986). HAP is the least soluble phase at pH values above about 4.0, and its solubility varies more quickly with pH than DCPD (Barone and Nancollas, 1978). Therefore, saliva and plaque fluid are more supersaturated with respect to HAP, and consequently, the precipitation of HAP is more likely to occur compared with other calcium phosphate phases. In a recent study (Poff *et al.*, 1997), however, it was reported that no significant correlation existed between calcium phosphate supersaturation in saliva and the rate of calculus formation for both unstimulated and stimulated saliva. However, calculus formation is influenced by a variety of factors, such as salivary flow rate, and inhibitors and promoters of calculus formation, other than salivary supersaturation with calcium phosphate salts. Unfortunately, all these factors were not controlled in this study. Therefore, there is an inherent limitation in the study by Poff *et al. in vivo* (1997). The lack of correlation between the degree of supersaturation and calculus formation may result from the effects of these factors. For example, if the subjects in one group have a higher degree of supersaturation with respect to calcium phosphates but lower salivary flow rate compared with those in the other group, the two groups may have similar calculus levels; that is, the correlation between the degree of supersaturation and calculus formation is overshadowed by the inter-group difference in salivary flow rate.

Salivary flow rate affects calcium phosphate saturation in glandular saliva. According to a study by Lagerlöf (1983), although parotid saliva is supersaturated with respect to HAP and WHT at all flow rates tested, ranging from 0.1 to 2.0 mL/min, parotid saliva is undersaturated with OCP at flow rates lower than 0.2 mL/min. With respect to DCPD, parotid saliva is undersaturated or just saturated at all flow rates, and the degree of saturation is only weakly influenced by flow rate. The higher salivary pH or possibly the increased secretions of calcium phosphate salts in glandular saliva may contribute to the effect of salivary flow rate on the degree of calcium phosphate saturation in saliva.

The effect of pH on calcium phosphate saturation degree in parotid saliva has also been investigated (Lagerlöf, 1983). As Lagerlöf reported, the degree of saturation with respect to HAP, OCP, and DCPD increased with increasing salivary pH at various flow rates. The parotid saliva was supersaturated with respect to HAP, WHT, and OCP at pH values above 5.5, 6.4, and 6.9, respectively. A close correlation ($r = 0.91$) between salivary pH and degree of supersaturation was observed by Poff *et al.* (1997). Take HAP, for example. The next equation shows why the degree of calcium phosphate saturation is influenced by pH: $Ca_{10}(PO_4)_6 \leftrightarrow (OH)_2 \cdot 10Ca^{2+} + 6PO_4^{3-} + 2OH^-$. When the calcium phosphate crystals in solution are in kinetic equilibrium, the rate of precipitation is equal to that of dissolution. If pH in solution drops (the concentration of hydrogen ions increases), OH^- and PO_4^{3-} tend to be removed by H^+ by

forming water and more acidic forms of phosphate, respectively. As a result, the equilibrium is broken and is pulled to the right; that is, the rate of dissolution exceeds that of precipitation, and the net result is the dissolution of HAP crystals and a decrease in the HAP saturation degree. If pH in solution rises, the opposite event will occur: OH^- forces the equilibrium in the equation to the left, thus resulting in an increased degree of HAP saturation in solution (Cotton and Wilkinson, 1966).

(D) THE INVOLVEMENT OF BACTERIA IN CALCULUS FORMATION

Although calculus can be induced in germ-free animals (Theilade *et al.*, 1964), human calculus development invariably involves plaque bacterial calcification. *In vitro* mineralization in specific plaque bacteria has been observed by various studies. Microbial mineralization occurs even within "acidogenic" and cariogenic bacteria (Moorer *et al.*, 1993). Studies have shown that filamentous micro-organisms are predominant in supragingival calculus and calculus-associated plaque, whereas micro-organisms of various morphologies are found in the plaque adjacent to subgingival calculus (Friskopp and Hammarström, 1980). In addition to dead micro-organisms, live and degenerated ones can also calcify in synthetic calcifying media (Sidaway, 1980). Additionally, deoxyribose and ribose have never been found in calculus, suggesting the absence of nucleic acids in calculus (Little *et al.*, 1966). The lack of nucleic acids in calculus indicates that the oral micro-organisms undergo extensive degradation, leaving only the cell walls for calculus formation.

Importantly, some oral bacteria seem to be associated with the absence of calculus. Pockets harboring undetectable subgingival calculus deposits appeared to have a significantly greater percentage of coccoid forms and fewer motile rods and total motile morphotypes (CM Brown *et al.*, 1991). In addition, non-calculus sites are associated with a significantly higher level of *Actinobacillus actinomycetemcomitans* (Aa) and a lower level of black-pigmented anaerobic rods than sites presenting with calculus. Aa has, therefore, been proposed to exert an inhibitory effect on the colonization of plaque-producing and calcifiable bacteria (Listgarten, 1987).

(E) MICROBIAL MINERALIZATION

It has been reported that initial deposition of apatite in calcifying bacteria is associated with membrane or acidic membrane-associated components (Boyan and Boskey, 1984). Although the levels of acidic phospholipids, principally phosphatidylserine and phosphatidylinositol, are generally low in biologic membranes compared with some other lipids, acidic phospholipids are the key components of the membranes involved in microbial calcification. Acidic phospholipid has a net negative charge at physiological pH and is amphipathic, with a hydrophobic tail and a hydrophilic head, so that either a spherical micelle or a bilayer can be formed, depending upon the type of molecule involved (Vogel and Boyan-Salyers, 1976). The functions of acidic phospholipids in calcification depend upon their ability to bind calcium by their negatively charged groups. Calcium is bound to adjacent phospholipid molecules in the membrane through a two-point electrostatic attachment to form a stern layer which facilitates the interaction of calcium with inorganic phosphate ions in solution (Hauster *et al.*, 1969). The bound calcium ions can cause

loss in water of hydration due to the neutralization of electrostatic charge of membrane (Hauster *et al.*, 1975). Desolvation helps to concentrate calcium and phosphate ions, thus facilitating their interactions. Inorganic phosphate associates with the bound calcium to form a Calcium-phospholipid-phosphate complex (CPLX). Once CPLX has formed, apatite deposition follows when sufficient calcium and phosphate ions are present and the concentration of inhibitors is low. The availability of CPLX exhibits bacterium-specificity; CPLX is always present in *Corynebacterium matruchotii* whether or not they are cultured in calcification-permissive media, while CPLX is absent from bacteria that do not calcify in culture (Boyan and Boskey, 1984). Moreover, not all of the acidic phospholipids are involved in the formation of CPLX, and phosphatidylserine is the predominant acidic phospholipid in CPLX isolated from *C. matruchotii*.

However, acidic phospholipids not associated with specific proteolipids in the membrane neither form CPLX nor support HAP deposition. It has been shown that when various lipid extracts from *C. matruchotii* are incubated in metastable calcium phosphate solutions, CPLX formation occurs predominantly on proteolipid-associated phospholipids (Boyan and Boskey, 1984). Calcifiability of bacteria is positively correlated with the increasing concentration of proteolipids. Further, the membrane-associated proteolipid purified from *C. matruchotii* can induce calcium precipitation *in vitro* (Van Dijk *et al.*, 1998). These observations suggest that acidic phospholipids must associate with specific non-polar protein to form a proteolipid complex, through hydrophobic interactions, to initiate calcification. Moreover, although all micro-organisms have proteolipids, not all micro-organisms calcify, suggesting a diversity in bacterial proteolipids. For example, proteolipids isolated from *A. naeslundii*, which are different in concentration and composition from those isolated from *C. matruchotii*, do not calcify in culture (Howell and Boyan-Salyers, 1980). This is consistent with the observation discussed above that not all the bacteria can form CPLX. Additionally, the amount of proteolipids has been observed to increase with culture time, and this finding partially explains the observation that there is a maturation of plaque before calcification can take place (Mandel, 1957).

The transport of ions through cell membrane is much faster than that predicted by the solubility constants of ions in lipids, suggesting that ions may be transported across biological membranes through ion channels within the phospholipid bilayer (Benga and Holmes, 1984). Proteolipid proteins exhibit conformational flexibility, which may contribute to their capacity to form ion channels in phospholipid bilayers. It has been demonstrated that protein ion channels are composed of several membranes spanning α -helices and that these α -helices are amphiphilic, with a hydrophilic face and a hydrophobic face. The α -helices are parallel, and their hydrophilic faces aggregate to form an ion channel (Lear *et al.*, 1988). A 10,000 M_r proteolipid and an 8500 M_r dicyclohexylcarbodiimide-binding proteolipid have been proposed to be two components of the ion channel (Swain and Boyan, 1989). Collectively, proteolipids may function not only as sites for CPLX formation but also as ion channels for transport of calcium and phosphate to the calcification sites or of protons away from the sites.

Lipids are also responsible for the calcification of dental calculus matrix. It has been known that although membranes

are mainly cellular constituents, either as the plasma membranes or as intracellular structures, membranes are also present in extracellular matrix of calcifying tissues in the form of matrix vesicles produced by a budding off the plasma membranes or by extrusion (Wuthier and Cummins, 1974). Therefore, the calcifiable proteolipids in the membranes of matrix vesicles can initiate the mineralization process in the calculus matrix as they do in calcifying bacteria.

(F) NUCLEATION INHIBITORS

It has been demonstrated that magnesium (Mg) prevents apatite nucleation by *C. matruchotii* proteolipid during incubation in a metastable calcium phosphate solution. At a level that does not alter calcium binding by *C. matruchotii* proteolipid, Mg blocks apatite crystallization and stabilizes calcium phosphate as amorphous mineral (Ennever and Vogel, 1981). Diphosphonates, such as ethane-1-hydroxy-1, 1-Diphosphonate (EHDP), inhibit both apatite nucleation (Fleisch *et al.*, 1970) and crystal growth (Francis, 1969). Importantly, nucleation inhibitors should not be used clinically, because they have been found to interfere with normal mineralization of hard tissues (Schenk *et al.*, 1973).

(G) CRYSTAL GROWTH INHIBITORS

Some salivary proteins containing negatively charged sequences may adsorb at active sites on the crystallite surfaces and thereby inhibit the growth and dissolution of calcium phosphate crystals. Of these negatively charged salivary proteins, statherin and PRP are two representatives of salivary inhibitors of crystal growth.

It is notable that macromolecules like PRP cannot completely block crystal growth (Margolis *et al.*, 1982). In an *in vitro* study by Margolis *et al.* (1982), the kinetics of crystal growth of HAP was studied with seed crystals of HAP coated with PRP-3 (a proline-rich phosphoprotein salivary inhibitor of crystal growth). In a solution with the nominal composition (1.06 mM in Ca, 0.63 mM in Pi, pH 7.4, 50 mM NaCl) as background electrolyte, a saturation degree with respect to HAP of 16.1, and an initial PRP3 concentration of 3.87×10^{-5} mol/L, PRP should be able to cover 99.9% adsorption sites available on the surface of HAP, and the application of NaF should not be able to initiate crystal growth. However, when a solution of NaF (1 ppm) was added, precipitation occurred as evidenced by the continuous decrease of the phosphate concentration. As Margolis *et al.* (1982) discussed, there may be three explanations for this phenomenon. First, other crystal growth sites are available for the fluoride to induce the formation of fluoridated HAP. Second, PRP-3 and fluoride bind to the same sites, and PRP-3 is displaced from the binding site by fluoride. Third, PRP-3 cannot cover all the available crystal growth sites. To elucidate this issue, Margolis *et al.* (1982) studied the kinetics of crystal growth of 40 mg HAP seed crystals with 50% growth sites covered with PRP-3 and 21 mg HAP seed crystals uncovered with PRP-3. After calculation, the number of 50% growth sites on 40 mg HAP seed crystals was equal to that of 100% growth sites on 21 mg HAP seed crystals. The results showed that the two groups of HAP seed crystals, in the solutions of various saturation degrees with respect to HAP, exhibited nearly identical kinetic behavior both before and after the addition of 0.75 ppm fluoride. If other crystal growth sites were available for fluoride, the number of these sites on 40 mg HAP seed crystals should have been nearly

twice that of the sites on 21 mg HAP. When NaF was applied, the growth of 40 mg HAP seed crystals should have been promoted much better than that of 21 mg HAP seed crystals. Thus, the kinetics of crystal growth after the addition of NaF should have been different. Since the kinetics was identical, the hypothesis of other crystal growth sites for fluoride is rejected. If PRP-3 were displaced from binding sites by fluoride, the growth of 40 mg HAP seed crystals should also have been better promoted by fluoride than that of 20 mg HAP seed crystals. The identical kinetics of crystal growth is best explained by the possibility that at maximum PRP-3 coverage, a small number of crystal growth sites remain uncovered. The two groups of seed crystals have equal numbers of growth sites and therefore exhibit a similar kinetics of crystal growth (Margolis *et al.*, 1982). The incomplete coverage of growth sites by PRP-3 may be due to steric reasons: PRPs are macromolecules, and one PRP molecule will impede the adsorption of other PRP molecules in proximity, thus leaving sites available for crystal growth.

The adsorption sites of PRPs have been investigated by Bennick *et al.* (1979). They have found that the adsorption sites are located in the proline-poor N-terminal part of PRPs between residues 3 and 25. Further, phosphoserine within the proline-poor N-terminal part may be necessary for optimal adsorption of the proteins to HAP, in that digestion of proteins by phosphatase resulted in considerable decrease in the adsorption by PRP. Additionally, the adsorbed PRP resists proteolytic digestion, since some of the susceptible bonds in the N-terminal part of proteins were protected from digestion by proteolytic enzymes, leaving the proline-rich C-terminal part to be removed gradually.

In addition to PRP and statherin, other salivary molecules with negatively charged residues close to one another may also have a high affinity for apatite surfaces and inhibit crystal growth. Some cystatins in saliva, such as the acidic cystatin S and the neutral cystatin SN, have several negatively charged residues in their N-termini and thus may be able to bind strongly to HAP. In a study by Johnsson *et al.* (1991), the adsorption, at HAP surfaces, of these two cystatins was compared with that of statherin. Although the affinity of cystatins for HAP surfaces was lower than that of statherin, their inhibitory effect on crystal growth was considerably greater.

The neutral histatin 1 also has high affinity for apatite surfaces, partly due to the presence of a phosphoserine residue in the vicinity of other negatively charged side-groups in the N-terminal part. The neutral histatin 1 from human parotid saliva has been demonstrated to inhibit crystal growth of HAP (Oppenheim *et al.*, 1986).

Immunoglobulins present in dental plaque and calculus may also have an inhibitory effect on plaque mineralization. The IgG and IgA detected in dental calculus are mainly distributed along the incremental lines, which contain fewer minerals. The IgG and IgA can also retard the growth rate of crystals through adsorption. So, the incremental lines are likely to be caused, at least in part, by the accumulation of IgG and IgA on the surface of dental calculus (Lindskog and Friskopp, 1983). Despite these findings, the exact role of these molecules in plaque mineralization remains obscure.

It was found that albumin could bind to HAP surfaces and inhibit crystal growth *in vitro* (Robinson *et al.*, 1998), indicating the possibility that albumin may be involved in the inhibition of crystal growth in dental plaque.

In addition to these salivary proteins, pyrophosphate

and zinc ions act as crystal growth inhibitors (discussed in a later section).

(H) ORGANIC ACID AND CALCULUS FORMATION

Numerous plaque bacteria produce organic acids through degradation of carbohydrates. The roles of organic acids in caries formation have been investigated by many investigators. However, a correlation between organic acids and dental calculus has seldom been reported. Despite that fact, our knowledge of dental caries can lead one to assume the potential role of organic acids in calculus formation, since both the enamel and supragingival calculus contain apatites and have similar oral environments.

The importance of lactic acid in demineralization of enamel is well-known. Other weak acids, however, also play a role in caries formation. As Featherstone and Rodgers (1981) observed, acetic, propionic, isobutyric, and succinic acids could produce subsurface caries-like lesions similar to those produced with lactic acid. It has been proposed that the un-ionized form of high pKa organic acid, such as acetic, butyric, and propionic acids, diffuse most rapidly into enamel compared with the ionized form of low pKa organic acids. As the un-ionized organic acids diffuse, they continually dissociate partially into hydrogen ions and acid anions. The released hydrogen ions attack the enamel crystals. Collectively, the process of demineralization is composed of the following three steps: diffusion of the un-ionized acids into enamel, partial dissolution of acids, and diffusion of calcium and phosphate out of enamel (Featherstone and Rodgers, 1981).

The preferential diffusion of high pKa organic acid into enamel was confirmed by Geddes *et al.* (1984). Enamel slices were immersed in a solution containing acetic acid and lactic acid. Dental plaque was not used in this system, to eliminate the possibility of acid loss other than into the enamel (such as bacterial utilization). The results showed that the acetate concentration decreased linearly with time, whereas lactate concentration did not change beyond experimental error. Since there was no dental plaque in this system, the only place where the acetate could have gone is into the enamel.

Importantly, the preferential diffusion of high pKa organic acid may also apply to our understanding of the effects of organic acids on the dental calculus formation. One could assume that the un-ionized, high pKa organic acids diffuse into dental calculus, dissolve the calcium phosphate crystals, and counteract the calculus calcification.

(I) ENZYMES DEGRADING CALCIFICATION INHIBITORS

According to a study by Watanabe *et al.* (1982), calculus level was positively correlated with protease activity in human saliva. In addition, supragingival plaque from calculus formers has also been found to show significantly higher protease activity than that from non-calculus formers (Morita and Watanabe, 1986). These findings are understandable, since protease in saliva and plaque can degrade calcification inhibitors such as statherin and PRP. Moreover, proteases may increase dental plaque pH through the production of ammonia, one of the proteolytic end-products of proteins (Frostell and Söder, 1970).

Acid and alkaline phosphatases are present in oral microorganisms, dental plaque, dental calculus, and saliva (Bercy

and Vreven, 1979). Acid and alkaline phosphatases may promote crystal growth by degrading pyrophosphate, which is an inhibitor of crystal growth and will be discussed in a later section (Francis, 1969). They hydrolyze phosphoproteins to produce inorganic phosphate ions (Poirier and Holt, 1983) and transport inorganic phosphate ions (Melani *et al.*, 1967) for mineralization. The levels of alkaline phosphatases in bones increase where mineralization occurs (Martland and Robinson, 1924), indicating a close relationship of alkaline phosphatases with biomineralization. In the field of dentistry, a positive correlation has been found between the amounts of dental calculus and salivary phosphatases (Bercy and Vreven, 1979).

Like phosphatase, acid and alkaline pyrophosphatases can also promote crystal growth by hydrolyzing pyrophosphate. The presence and activity of pyrophosphatases in plaque and calculus have been demonstrated (Pellat and Grand, 1986). Alkaline pyrophosphatase activity occurring in dental plaque is positively correlated with calculus formation *in vivo* (Bercy and Vreven, 1979), despite the fact that the pH of dental plaque does not favor alkaline pyrophosphatase activity (pH optimum, 8.5). A positive correlation between calculus formation and acid pyrophosphatase activity of saliva has also been observed (Bercy and Vreven, 1979). Importantly, a magnesium pyrophosphate complex appears to be the real substrate for pyrophosphatase, based upon the findings that pyrophosphatase activity is substantially decreased in the absence of magnesium and completely inhibited by calcium (Pellat and Grand, 1986).

(J) CALCIFICATION PROMOTERS

(1) Urea

Urea is a product from the metabolism of nitrogen-containing substances. Urea can be secreted in normal saliva at concentrations of between 5 and 10 mmol/L (Macpherson and Dawes, 1991), but can be as high as 30 mmol/L in patients with renal disease (Peterson *et al.*, 1985). Gingival crevicular fluid contains up to 60 mmol/L urea (Golub *et al.*, 1971). It has been demonstrated that urease is responsible for bacterial urea hydrolysis (Sissons *et al.*, 1995). At a neutral pH, urea is hydrolyzed by urease to NH_4^+ and bicarbonate (Sissons *et al.*, 1985). At an acidic pH, aqueous and gaseous carbon dioxide equilibrate with bicarbonate, while, at an alkaline pH, aqueous and gaseous NH_3 co-exists with NH_4^+ (Sissons *et al.*, 1994b).

(a) The effect of urea metabolism on plaque pH

Ammonia produced from ureolysis of urea contributes to an increased plaque pH that is an essential factor in natural calculus formation. A urea-induced pH response, which was the inverse of the Stephan pH curve induced by sucrose, has been observed in an *in vitro* biofilm culture system called "artificial mouth" (discussed in a later section) (Sissons *et al.*, 1991). The curve of a urea-induced pH response contains two parts—a rising part and a return part; that is, following the application of urea to plaque, plaque pH rises quickly and then returns slowly to the level before the urea application.

In the "artificial mouth", when supplemented with urea, sucrose-dependent pH responses in the microcosm plaque shift to a more alkaline range and result in a higher resting pH than in that without urea supplementation (Sissons *et al.*, 1998). Urea metabolism can also contribute to the pH gradient in plaque, because the diffusion and removal of ammonia are

very slow (Sissons *et al.*, 1994a). When urea is initially applied to a plaque, the plaque bacteria produce ammonia through urease, and ammonia accumulates on the plaque surface due to the slow removal and diffusion of ammonia. At this point, the pH gradient ranges from low at the "artificial mouth" plaque base to high at the surface. During the following minutes, some ammonia diffuses and arrives at the middle-layer plaque, and this leads to an increase in the pH at the middle layer. Meanwhile, some ammonia is removed from the plaque surface. Both the diffusion into plaque and the removal of ammonia decrease the pH at the surface. Therefore, the mid-plaque-to-surface pH gradient gradually reverses. As ammonia continuously diffuses inside the plaque, pH at the base gradually increases and becomes the highest. The pH at the base remains high for hours, and then pH in all layers in the plaque slowly returns to the resting pH level. The ureolytic pH response is roughly proportional to the urea concentration (Sissons *et al.*, 1994a). The ureolytic pH response (an increase in plaque pH by the production of ammonia from urea) promotes calculus formation by increasing the saturation degree of calcium phosphate in plaque fluid. (The mechanism is discussed in a previous section.)

(b) The effect of salivary flow rate on urea-dependent pH response in plaque

In the oral cavity, urea available to supragingival plaque bacteria comes from both saliva and extra-oral factors such as urea-containing chewing gum and toothpastes. The effects of salivary flow rate on the pH responses induced by urea from these two sources are different. In the absence of exogenous urea, increased salivary flow rate promotes the urea-dependent pH response in plaque. Although a negative correlation has been demonstrated between salivary flow rate and urea concentration in whole saliva (Macpherson and Dawes, 1991) and parotid saliva (Shannon and Prigmore, 1960), higher flow rate means that more urea is available to dental plaque bacteria, because the effects of low urea concentrations are outweighed by an increased availability of urea. Thus, the sites with higher salivary film velocity have an increased urea-dependent pH response in plaque (Shannon and Prigmore, 1960). That can partially explain the site-specificity of calculus distribution in the dentition.

Sissons *et al.* (1994a), using an "artificial mouth", investigated the effect of flow rate of a basal medium containing 0.25% mucin (BMM) on urea-induced pH responses in plaques. In the "artificial mouth", BMM works as artificial saliva. It was found that increasing the BMM flow rate significantly decreased the urea-induced pH response with respect to the maximum pH reached and pH curve area (representing the total amount of ammonia produced). The reduced pH response can be explained by urea dilution caused by an increasing BMM flow. Additionally, the urea-induced pH response varies more quickly with BMM flow rate when the flow rate is lower, indicating that ammonia clearance may be limited at higher flow rates.

(c) Bacteria responsible for the ureolysis in dental plaque

According to Frostell (1960), none of the known species of oral ureolytic bacteria contributes significantly to plaque ureolysis, and most plaque ureolysis is due to an unknown but highly active sub-population of the total ureolytic flora. However, in a study by Sissons *et al.* (1988a), using "artificial mouth" plaque, the ureolytic-

ic fraction of the flora detected on urea-agar plates explained 86% of the plaque urea metabolism rate. The plaque bacteria giving strong ureolytic reactions on agar plates were all Gram-positive cocci, and in 6 of the 9 plaques were streptococci only. Since Gram-positive cocci, mainly streptococci, comprise a high proportion of the total ureolytic flora, the hypothesis that plaque ureolysis is derived mainly from an unidentified active segment of the total ureolytic flora is rejected. Bacteria suspected of having a role in ureolysis include *S. salivarius*, coagulase-negative staphylococci (CNS) (Sissons *et al.*, 1988a,b), *Actinomyces viscosus/naeslundii* (Gallagher *et al.*, 1984), transient *Enterobacteriaceae*, unknown anaerobes (Frostell, 1960), and *Haemophilus sp.* (Salako and Kleinberg, 1989). Among these ureolytic bacteria, *S. salivarius* has attracted the most attention. It has been accepted that *S. salivarius* is a major contributor to ureolysis in natural saliva (Sissons *et al.*, 1989) and in artificial plaque cultured from saliva (Sissons *et al.*, 1988a,b). Although *S. salivarius* is regarded as a minor component of mature plaques, its number is sufficient to enable it to play a major potential role in ureolysis in plaque (Denepitiya and Kleinberg, 1982).

(2) Fluoride and calculus formation

The caries-inhibiting ability of fluoride is well-known. Fluoride not only counteracts demineralization of hard tissue through the formation of lower-solubility fluorapatite by fluoride substitution for hydroxyl ions and adsorption onto apatite surfaces (Wong *et al.*, 1987), but also contributes to remineralization by precipitation of a fluoride-enriched apatite or calcium fluoride (Nelsons *et al.*, 1984). Moreover, fluoride has been found to curtail greatly or even eliminate the appearance of OCP-like precursor phases during spontaneous calcium phosphate precipitation. In other words, fluoride may promote the maturation of spontaneous precipitated calcium phosphate at physiological pH by reducing the stability of OCP (Eanes and Meyer, 1978). Fluoride has also been demonstrated to affect the morphology of the apatite crystals as it converts them from thin plates to short and slender needles (Eanes and Meyer, 1978).

Fluoride has an antimicrobial function: A mixed culture with 19 ppm fluoride can inhibit the growth of *S. mutans* (Marsh and Bradshaw, 1990). A decreased level of *S. mutans* in dental plaque adjacent to fluoride-releasing glass ionomers has also been reported (Berg *et al.*, 1990). In addition, acid production by plaque bacteria can be inhibited by the presence of fluoride (Briner and Francis, 1962). Therefore, fluoride may have the potential to increase the plaque pH, which may be another mechanism by which fluoride inhibits demineralization and promotes remineralization of hard tissue.

Taken together, one can assume that fluoride present in saliva from the diet, toothpastes, and mouthrinses should be able to facilitate calculus formation. However, few studies or reports have shown that fluoride increases the prevalence or severity of dental calculus in humans. Instead, sodium fluoride may be able to inhibit the bacterial phosphatases and pyrophosphatases, the enzymes that are well-known to promote calculus formation (Lobene and Volpe, 1987).

(3) Silicon

Silicon is present in saliva, dental plaque, and dental calculus. The source of silicon is mainly drinking water and food. Silicic acid and silica represent most of the silicon found in drinking water and food, respectively. The family of silicic acid contains

monosilicic acid and polysilicic acid. Monosilicic acid is not stable at concentrations above 2 mmol/L and may polymerize to form polysilicic acid or silica (Damen and ten Cate, 1989).

Silicic acid has been reported as a strong promoter of both spontaneous precipitation of calcium phosphates and the growth of seeded crystals (Damen and ten Cate, 1989). In a study by Damen and ten Cate (1989), silicic acid initiated spontaneous calcium phosphate precipitation earlier by reducing the induction time of precipitation. Additionally, the stimulating effect of silicic acid on the growth of seeded HAP crystals was concentration-dependent. Silicic acid was also found to stimulate the transformation from amorphous calcium phosphate to HAP (Hidaka *et al.*, 1993). The capacity of silicic acid to promote HAP transformation stems from the presence of polysilicic acid, rather than monosilicic acid (Hidaka *et al.*, 1993). It was found that the polysilicic-acid-containing solutions stimulated HAP transformation, whereas those containing monosilicic acid failed to do so (Hidaka *et al.*, 1993).

Silica at a concentration of 0-2 mg/mL was reported to exert stimulating effects on calcium phosphate precipitation (Damen and ten Cate, 1989). Silica was also reported to increase the rate of calculus formation 35 days after its incorporation into food. Silica can also promote calculus formation when ingested by stomach tube. When provided *via* a stomach tube, silica does not promote calculus formation until 49 days after this treatment (Rølla *et al.*, 1989). Thus, it is likely that silica may remain in plaque for a long period of time and serve as a trigger for the increase of calculus formation.

There is also epidemiological evidence for the effect of silicon on calculus formation. It has been reported that the rate of calculus formation is higher in the Indonesian than in the Norwegian population (Gaare *et al.*, 1989). The difference in the silicon content of the food may partially explain the phenomenon. Indonesian people consume larger amounts of rice, which is enriched in silicon. The rice-based diet of Indonesian people contributes to the greater calculus formation in this population.

The calcium-binding capacity of silicic acids and silica may account for their ability to stimulate calcium phosphate precipitation. The stimulating effects may derive from the OH-silanol groups on their surfaces, which bind cations at neutral and acid pH (Ruvillac, 1982). On the other hand, it is noteworthy that 3.0-30.0 mmol/L silicic acid and 10 mg/mL silica inhibit the rates of both amorphous calcium phosphate formation and HAP transformation. The inhibitory effects on calcium phosphate precipitation of silicic acid and silica at relatively high concentration may be due to their increased chelating effects.

(IV) Anti-calculus Agents Used in Commercial Dentifrices

Toothbrushing is relatively effective in dental plaque removal, but it is still inadequate for the maintenance of gingival health. Chemotherapeutic agents have been used to supplement the mechanical removal of dental plaque (Aleece and Forscher, 1954; Grossman, 1954; Zacherl *et al.*, 1985; Volpe *et al.*, 1992). Early attempts with chemotherapeutic agents focused on the removal of dental calculus from teeth. The use of mucinase is an example, since it was believed that enzymatic dissolution of the organic matter in calculus could help destroy the calculus structures (Aleece and Forscher, 1954). Chelating agents are known to sequester and dissolve calcium salts by forming stable and soluble calcium complexes. For instance, Ex347, a chelating

agent, has been confirmed to effectively prevent the formation of calculus (Grossman, 1954). However, chelating agents can cause enamel damage (Weinstein and Mandel, 1964). Antimicrobial agents have also been used for calculus reduction, since micro-organisms are important for dental calculus formation. However, their usefulness is undermined because of the potential problem of developing resistance to antibiotics (Mandel, 1995). Since the 1970s, the major anti-calculus strategy has focused on inhibiting crystal growth and preventing development of mineralized plaque. At present, anti-calculus agents in use include triclosan (antimicrobial agent) with polyvinyl methyl ether (PVM) and maleic acid (MA) copolymer, and crystal growth inhibitors, including pyrophosphate with PVM/MA copolymer, zinc citrate, and zinc chloride.

Prior to a further discussion of anti-calculus agents, several important points must be elucidated. First, crystal growth inhibitors, such as pyrophosphate salts and zinc salts, have been proven effective only for the control of supragingival calculus and not for subgingival calculus. Second, the use of crystal growth inhibitors is directed at the prevention of deposit formation. Third, crystal growth inhibitors such as pyrophosphate and zinc ion typically decrease the dissolution rate of chemisorbed surfaces (Fleisch, 1981). The decrease in dissolution rate, as in the case of fluoride, could provide cariostatic benefits to treated teeth. Therefore, crystal growth inhibitors may exhibit two simultaneous functions by binding to apatite surfaces—preventing remineralization and preventing demineralization. Fourth, the currently marketed anti-calculus dentifrices and mouthrinses do not include crystal growth inhibitors alone, but also contain some form of fluoride as an anti-caries agent. Last, it is noteworthy that the concentration of crystal growth inhibitors used in dentifrices and mouthrinses is actually 2-3 orders of magnitude higher than that required for maximum coverage of growth sites on the HAP, OCP, and DCPD phases as calculated by the Langmuir analysis of crystallization rates (White *et al.*, 1989). Overgrowth of crystals (discussed in the paragraph “Pyrophosphate and PVM/MA”, below) and degradation of inhibitors have been demonstrated to impede the function of these agents. Therefore, a residual action of inhibitors retained as a “reservoir” within the plaque fluid between treatments is necessary. The residual inhibitors from the “reservoir” can coat the newly developing crystals within the plaque matrix and thus counteract the effect of overgrowth of crystals (White *et al.*, 1989).

(A) TRICLOSAN WITH A PVM/MA COPOLYMER

Triclosan is a broad-spectrum antibacterial agent active on both Gram-positive and -negative micro-organisms. The target is the cytoplasmic membrane. At bacteriostatic concentration, triclosan prevents bacterial uptake of essential amino acids, while at bactericidal concentrations, triclosan destroys the integrity of the cytoplasmic membrane and causes leakage of cellular contents (Regos and Hitz, 1974). Besides acting as an antibacterial agent, triclosan may have an anti-inflammatory function, since it can neutralize the products of bacteria that can provoke inflammation. It is also a potent inhibitor of both cyclo-oxygenase and lipoxygenase pathways (Gaffar *et al.*, 1995).

For triclosan to be effective, a delivery system is necessary to increase its residence time in the oral cavity. PVM/MA copolymer (trade name Gantrez) has been utilized as a delivery system for triclosan. PVM/MA copolymer promotes uptake of triclosan by enamel and buccal epithelial cells (Nabi *et al.*, 1989).

Increased retention of triclosan has also been observed in both plaque and saliva when a dentifrice containing triclosan and PVM/MA copolymer was used (Afflitto *et al.*, 1990). The mechanism by which PVM/MA copolymer enhances the delivery of triclosan has been elucidated (Gaffar *et al.*, 1997). The copolymer is composed of two groups: an attachment group and a solubilizing group. The solubilizing group retains triclosan in surfactant micelles so that the attachment group can have enough time to react with tooth surfaces *via* calcium in the liquid adherent layer. Triclosan is then slowly released *via* interactions with the salivary environment. In addition to the above functions, the copolymer also has weak crystal growth inhibitory property and is effective against calculus formation (Gaffar *et al.*, 1990). The copolymer can strongly complex and sequester magnesium and therefore inhibit the hydrolysis of pyrophosphate by alkaline phosphatases (Mandel, 1992).

The effects on calculus formation of triclosan and copolymer have been substantiated by clinical studies (Volpe *et al.*, 1992; Bánóczy *et al.*, 1995). It was found that 0.3% triclosan and 2.0% PVM/MA copolymer in a 0.243% sodium fluoride/silica base dentifrice significantly reduced the severity and occurrence of supragingival calculus after complete prophylaxis compared with placebo dentifrice.

(B) PYROPHOSPHATE AND PVM/MA COPOLYMER

EHDP has been demonstrated as an effective inhibitor of HAP crystal growth when adsorbed onto the surfaces of HAP crystals (Francis, 1969). However, because of its hydrolytic stability in the oral cavity and its inhibitory effect on apatite nucleation (discussed in a previous section), it can interfere with normal mineralization of hard tissues (Schenk *et al.*, 1973). Therefore, EHDP should not be used in dentifrices. Pyrophosphate is a biologically stable analog of diphosphonates in which the stable P-C-P linkage has been replaced with a labile P-O-P linkage (Fleisch *et al.*, 1969). In contrast to EHDP, pyrophosphate is hydrolytically labile. Its hydrolytic instability is enhanced by high temperature, low pH, and certain enzymes (Francis, 1969). Pyrophosphate, which is present in serum and saliva, is a soluble inorganic small molecule (Stookey *et al.*, 1989). Unlike macromolecules such as PRP, a small molecule is able to block all the adsorption sites of apatitic surface available for crystal growth, and therefore can block crystal growth even when the precipitation driving force is enhanced by the addition of fluoride (Moreno *et al.*, 1989). Pyrophosphate is a small molecule and has been reported to inhibit crystal growth by binding to the surface of crystal. Pyrophosphate binds to two sites on the HAP surface, and one of the two sites needs to be bound by phosphate ion to permit crystal growth to occur. If this site is bound by pyrophosphate, phosphate ion cannot adsorb onto crystal, and thus crystal growth is inhibited. To inhibit crystal growth effectively, the concentration of pyrophosphate has to reach a critical level. Below this level, the addition of NaF can induce a short period of slow precipitation, which is followed by rapid crystal growth (Moreno *et al.*, 1989). One explanation for the induction of crystal growth by NaF is overgrowth of crystals. During crystal growth, the inhibitors are gradually buried by continuously growing fronts, and consequently a new surface without inhibitors is generated, and crystal growth proceeds at rates comparable with those of the control systems (in the absence of inhibitors). This phenomenon does not take place if inhibitors are macromolecules such as PRP,

since it is difficult to bury these large molecules. In addition to the inhibitory effect on crystal growth, pyrophosphate can also delay the initiation of conversion of DCPD to HAP by over three-fold (White *et al.*, 1989) and reduce acquired pellicle formation. It has been reported that pyrophosphate can desorb the acquired enamel pellicle, due to its ability to displace anions and negatively charged macromolecules from tooth surfaces (Rølla and Melsen, 1975).

Pyrophosphate at various concentrations has been widely used as an anti-calculus agent in dentifrices and mouthrinses. Dentifrices containing pyrophosphate have been confirmed to produce significant calculus reduction (Zacherl *et al.*, 1985). However, pyrophosphate is susceptible to rapid breakdown in the oral cavity by bacterial and host phosphatases and pyrophosphatases. The addition of copolymer seems to be a good idea, since PVM/MA copolymer is believed to prevent hydrolysis of the pyrophosphate (Gaffar and Esposito, 1986). In clinical studies, a dentifrice containing 3.3% pyrophosphate and 1% PVM/MA copolymer significantly inhibited the formation of supragingival calculus as compared with a placebo dentifrice (Triratana *et al.*, 1991), and it was more effective than a dentifrice containing 3.3% pyrophosphate alone (Schiff, 1987).

(C) ZINC ION

Zinc salts are thought to reduce plaque acidogenicity (Oppermann *et al.*, 1980) and plaque growth *in vivo* (Saxton *et al.*, 1986) and to increase resistance of HAP to acid dissolution (Brudevold *et al.*, 1963). Zinc can also inhibit crystal growth by binding to the surfaces of solid calcium phosphates (Gilbert and Ingram, 1988). This binding is reversible and may be caused by ion exchange between zinc ions and surface calcium ions (Brudevold *et al.*, 1963). Zinc ions also have an effect on the types and amounts of calcium phosphate crystals (Le Geros *et al.*, 1999). Zinc at a concentration of 0.1 mmol/L inhibits the formation of DCPD, OCP, and amorphous calcium phosphate, but at higher concentrations varying from 0.5 mmol/L to 2 mmol/L, it promotes the formation of amorphous calcium phosphate or zinc-substituted tricalcium phosphate (beta-TCP), depending on the pH values and temperature. Additionally, the degree of crystallinity may be affected by the presence of zinc (Leskovar and Hartung, 1977).

The effect of zinc ion on calculus formation has been examined by Kazmierczak *et al.* (1990) in a six-month clinical study. A 2.0% zinc citrate dentifrice significantly reduced supragingival calculus deposits, compared with a placebo dentifrice. However, commercialization of dentifrice containing 2.0% zinc chloride is difficult, due to its disagreeable taste. Therefore, 0.5% zinc citrate was tested. In a three-month study, the mean calculus score (Volpe-Manhold Index) in a group using the 0.5% zinc citrate dentifrice was significantly (13.7%) lower than that in a group using a control dentifrice (Segreto *et al.*, 1991). However, Scruggs *et al.* (1991) found that a 0.5% zinc citrate dentifrice did not significantly reduce supragingival calculus. Because of these conflicting results, it would seem preferable to use a higher concentration of zinc citrate for dentifrice formulation.

(D) PREVENTION OF CALCULUS FORMATION IN IMPLANT PATIENTS

Although bacterial adhesion and plaque formation on implant surfaces, especially on Ti surfaces, have been investigated in

many studies (Nakazato *et al.*, 1989; Edgerton *et al.*, 1996), our knowledge of the plaque mineralization on implant surfaces is very limited. Theoretically, various anti-calculus agents, which were discussed above, can also be active against calculus formation on implant surfaces in addition to tooth surfaces. This needs to be confirmed by future studies.

(V) Application of Biofilm Culture Systems in the Study of Plaque Mineralization

In vivo studies of dental plaque present many difficulties. Natural dental plaque has a biodiverse, heterogeneous structure which changes over time. The oral environment is almost uncontrollable and site-specific. Therefore, it is necessary to develop *in vitro* culture systems which simulate the natural oral environment and are more controllable. Various biofilm culture systems have been developed in recent decades, and each system has strengths and limitations. The major oral biofilm culture system technologies in current use include chemostat-base systems (Keevil *et al.*, 1987), a growth-rate-controlled biofilm fermenter (GRBF) (Gilbert *et al.*, 1997), an artificial mouth (Dibdin *et al.*, 1976), the constant-depth biofilm fermenter (CDFF) (Sarah *et al.*, 1996), and the multiplaque artificial mouth (Sissons *et al.*, 1991). Due to the design of the multiplaque artificial mouth, which permits the long-term independent growth of multiple replicate plaques from the same inoculum, at the same temperature, and in the same gas phases (Sissons *et al.*, 1991), it appears to be suitable for studies of plaque growth, metabolism, plaque pH, and mineralization (Sissons *et al.*, 1991). The use of LabVIEW[®] software on a Macintosh computer makes it easier to control separate delivery of nutrients, reagents, and gas required for the culture of multiple dental plaque as well as continuous acquisition of pH data (Wong *et al.*, 1994). Urea metabolism, which is closely associated with plaque mineralization, has been investigated in studies where the multiplaque artificial mouth system was used to detect the ureolytic bacteria and to examine the properties of urea metabolism in microcosm plaques. According to a study by Sissons *et al.* (1988b), ureolytic bacteria could be detected on Christensen's urea-agar plates or urea segregation agar. On the two plates, colonies growing rapidly to 0.5 mm or larger are assumed to be involved in ureolytic activity. Moreover, the average rate of ureolysis of microcosm plaques is similar to that in natural plaques (Sissons *et al.*, 1988b). Therefore, the validity of the multiplaque artificial mouth system in the study of urea metabolism is further reinforced. In other studies where the multiplaque artificial mouth was used (Sissons *et al.*, 1988a, 1990, 1994a,b), the effects of pH, bicarbonate, carbon dioxide, ammonia, and glucose on bacterial urease levels were examined. Urea-induced pH responses and intra-microcosm plaque pH gradient were observed and discussed. Furthermore, regulation of urease level in microcosm plaque became possible by controlled applications of urea, ammonia, and arginine (Sissons *et al.*, 1995). Direct studies of microcosm plaque mineralization were also performed (Pearce and Sissons, 1987; Pearce *et al.*, 1991; Sissons *et al.*, 1991; Wong, 1998), and a urea-based calcium-phosphate-monofluorophosphate-urea (CPMU) mineralization solution was used for plaque mineralization research (Sissons *et al.*, 1991; Wong, 1998).

(VI) Future Research on Supragingival Calculus

There have been many advances in our understanding of how plaque and microbial mineralization is initiated and regulated by various inhibitors and promoters, and how it can be con-

trolled and prevented. However, there are many questions left to be answered. The roles of macromolecules, such as PRPs, statherins, immunoglobulins, and other glycoproteins, in the initiation of biomineralization remain incompletely understood. The potential effects of fluoride on calculus formation need to be clarified. More studies on ureolytic bacteria, which contribute to the increase in plaque pH, are also recommended. Development of biofilm culture systems, notably the "artificial mouth", could lead to breakthroughs in our understanding of mineralization mechanisms. In the "artificial mouth", various antibacterial agents, which may be used in future dentifrices and mouthrinses, can be evaluated with regard to their possible inhibitory effects on plaque and calculus formation. The effects of the various anti-calculus agents on plaque mineralization on implant surfaces also need to be confirmed. In addition, safer and more effective calculus control formulations should be a long-term goal in this field.

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