Journal of Periodontology

Modulation of clinical expression of plaque-induced gingivitis I. Background review and rationale

Tatakis DN, Trombelli L: Modulation of clinical expression of plaque-induced gingivitis. I. Background review and rationale. J Clin Periodontol 2004; 31: 229–238. doi: 10.1111/j.1600-051X.2004.00477.x © Blackwell Munksgaard, 2004.

Abstract

Objectives: The purpose of this article is to provide the necessary background and rationale for the accompanying studies, which are ultimately aimed at identifying genetic and environmental factors determining gingivitis susceptibility.

Materials and methods: The literature on factors reported to modify the clinical expression of gingivitis, i.e., factors that determine individual variability in gingival inflammatory response to plaque, is presented.

Results: Clinical evidence suggests that the gingival inflammatory response to plaque accumulation may differ substantially among individuals. However, most of the available studies are of small scale and not purposely designed to address the issue. Systemic factors implicated in modulation of the clinical expression of gingivitis include metabolic, genetic, environmental and other factors. The significance of such factors in designing and conducting a large-scale experimental gingivitis trial and means to account for them are discussed.

Conclusion: Although several factors have been implicated, genetic or environmental factors underlying differences in gingivitis expression are not fully elucidated. The accompanying studies aim to identify and characterize, among participants in a specifically designed large-scale experimental gingivitis trial, subjects that differ significantly in their gingival inflammatory response to plaque. This is the first step in an effort to determine genetic or environmental factors underlying such differences.

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Key words: experimental design; gingivitis; gingivitis/etiology; gingivitis/physiopathology

Accepted for publication 25 May 2003

The two most prevalent and most investigated diseases of the periodontium, i.e. plaque-induced gingivitis, a reversible condition, and chronic periodontitis, an irreversible condition that can lead to tooth loss, are chronic inflammatory conditions an of an infectious nature (Williams 1990). The unequivocal role of dental bacterial plaque in the development of these diseases has been established since almost 40 years ago (Löe et al. 1965, Theilade et al. 1966, Lindhe et al. 1973, Slots & Hausmann 1979, Holt et al. 1988, Breuer & Cosgrove 1989, Persson et al. 1994). Although ample evidence suggests that susceptibility to periodontitis varies considerably among individuals (Löe et al. 1986), in large part because of genetic factors (Hart & Kornman 1997, Michalowicz et al. 2000), there are limited studies addressing a potential host-dependent variation in susceptibility to gingivitis.

Dental plaque-induced gingivitis (Mariotti 1999) is the most prevalent disease (Oliver et al. 1998, Albandar & Tinoco 2002, Sheiham & Netuveli 2002) among the numerous conditions that affect the periodontium (Armitage 1999, Laskaris & Scully 2003). The definitive evidence provided by the pioneering studies of Löe and colleagues in the mid 1960s confirmed that gingivitis is caused by dental bacterial plaque (Löe et al. 1965, Theilade et al. 1966). The etiologic role of plaque in the development of gingivitis is further supported by numerous intervention studies. Plaque reduction or elimination, either by mechanical means (Löe et al. 1965, Bosman & Powell 1977, Brecx et al. 1988) or chemical anti-microbial agents (Löe et al. 1967, Bosman & Powell 1977, Brecx et al. 1990), leads to disease resolution. Earlier, experimental gingivitis studies have alluded to possible subject differences in susceptibility to gingivitis. Such differences have been usually ascribed to differences in plaque accumulation rates (quantitative plaque differences) and/or differences in plaque species present (qualitative plaque differences) (Löe et al. 1965, Theilade et al. 1966). However, quantitative and/or qualitative plaque differences cannot account for all the reported differences in gingivitis susceptibility (Abbas et al. 1986a, Lie et al. 1995, 1998). The scientific evidence considered above is based on clinical studies.

The clinical assessment of the gingival inflammatory response to plaque accumulation relies on several parameters (Mariotti 1999). Visually assessed parameters include the extent and severity of changes in the physical status of the gingiva, such as changes in color, in surface anatomy (contour), and in bleeding tendency. Several indices have been proposed for the clinical evaluation of gingival inflammation, including the papilla, marginal, attached (PMA) index (Schour & Massler 1947), the papillary bleeding index (PBI) (Mühlemann 1977), and the gingival index (GI) of Löe & Silness (1963), among others (Ciancio 1986). The visually assessed signs of inflammation correspond to histopathological tissue changes (Hancock et al. 1979, Greenstein et al. 1981, Engelberger et al. 1983). The various indices have been shown to each have its own limitations, a major common one being the subjective nature of some of the required determinations. To add objective means to the clinical assessment of the status of the gingiva, quantitation of the gingival crevicular fluid has been developed and employed widely (Golub & Kleinberg 1976, Hinrichs et al. 1984a, b. Ciantar & Caruana 1998). Abundant evidence indicates that crevicular fluid flow correlates with both the clinical and the histological status of the gingiva (Oliver et al. 1969, Rüdin et al. 1970, Daneshmand & Wade 1976, Engelberger et al. 1983). Methods and indices to quantify the accumulated dental bacterial plaque, such as the Debris index (Greene & Vermillion 1960), the Quigley and Hein plaque index (Quigley & Hein 1962), and the plaque index by Silness & Löe (1964), among others (Fischman 1986), have been necessary in the assessment of the gingival response to the plaque. Use of these indices has helped demonstrate the relationship between the extent of plaque deposits and the severity of gingivitis (Breuer & Cosgrove 1989, Müller et al. 2000). Beyond the quantitative aspects of this relationship, there are important qualitative aspects, as well. Analysis of the microbial composition of plaque has led to the implication of certain bacterial species in the development of gingivitis (Moore et al. 1984). Furthermore, more sophisticated means to evaluate the inter-relationship between accumulated plaque and gingival response have been pursued from the qualitative aspect of the host response, such as characterization of relevant molecular determinants present in crevicular fluid (Lamster et al. 1985, Patters et al. 1989, Heasman et al. 1993a, Herrmann et al. 2001, Lundy et al. 2001).

Regardless of the methods used for the clinical evaluation of the gingiva (or plaque) in either natural or experimental gingivitis, it is clear that there are significant differences among individuals with respect to the severity and extent of gingivitis they experience. Characterization of the nature of these differences and identification of the underlying etiology have been the impetus for the design and implementation by the authors of a series of studies (Trombelli et al. 2003a, b, unpublished data) whose immediate goal is to define and characterize subpopulations with significant differences in gingivitis susceptibility. The ultimate goal of these studies is the identification of hostdependent variables, possibly genetic ones, which characterize the two subpopulations. The purpose of this review paper is to provide the necessary background and the rationale for these studies.

Factors modulating gingivitis expression

To undertake a series of large-scale experimental gingivitis studies aimed at identifying individual characteristics associated with increased susceptibility to plaque-induced gingival inflammation requires, at the very least, acknowledgement of the factors already established or implicated in the literature as modifiers of gingival inflammation. It also requires, whenever possible, measures to control for such factors, either during study design or during data analysis. In the following paragraphs, these factors are reviewed and discussed in relation to the design and execution of such an experimental gingivitis trial.

Local factors modulating gingivitis expression

In many circumstances, particularly in relation to natural gingivitis, the reasons for exacerbated manifestation of gingivitis have been attributed to local factors, implicated in increased plaque accumulation or hindrance of oral hygiene procedures. Such local factors reported to lead to deteriorating gingival/periodontal health include developmental or anatomical tooth variations [palato-gingival groove (Hou & Tsai 1993), enamel pearls (Goldstein 1979), crowding (Chung et al. 2000)], pathological tooth conditions [fractures (Polson 1977), caries (Albandar et al. 1995)], gingival anatomical conditions [recession defects (Smukler & Machtei 1987, Goutoudi et al. 1997), frenum position (Addy et al. 1987)], and iatrogenic factors [subgingival restoration margins (Waerhaug 1975, Bader et al. 1991), overhangs (Rodriguez-Ferrer et al. 1980, Lang et al. 1983), partial dentures (Bissada et al. 1974, Yeung et al. 2000), orthodontic appliances (Boyd & Baumrind 1992), etc.]. Maxillofacial anatomic variants (e.g. inadequate upper lip coverage) and/or upper respiratory obstructions (e.g. epipharyngeal adenoids, deviated nasal septum) may lead to mouthbreathing. Mouthbreathing and inadequate upper lip coverage have been shown to lead to changes in plaque accumulation and gingivitis expression, particularly in the maxillary anterior segment of the dentition (Jacobson 1973, Addy et al. 1987, Wagaiyu & Ashley 1991, Gulati et al. 1998).

The frequent intake of sucrose, acting on a local level, is well established as responsible for increased plaque accumulation and concomitantly increased gingival inflammation (Jalil et al. 1983, Sidi & Ashley 1984). Although addressed here, under local factors, because of its action, increased sucrose intake can also be considered an environmental factor.

Under the above conditions, the cause of the increased (or decreased) gingival inflammation manifested in the subjects is identifiable and amenable to corrective action. In the context of experimental gingivitis trial design, such factors are controllable by establishing appropriate inclusion and exclusion criteria, as well as detailed instructions for study participants. More importantly, such local factors are not expected to alter the inherent manner in which the host tissues respond to plaque accumulation, which is the main thrust of the present studies. Therefore, these local factors, although critical for individual patient management in daily practice, do not merit further discussion here.

Systemic factors modulating gingivitis expression

The most recent classification of periodontal diseases (Armitage 1999)

acknowledges that the clinical expression of plaque-induced gingival inflammation can be substantially modified by systemic factors, either inherent to the host or related to environmental influences (Mariotti 1999). For the purposes of this review, the systemic factors will be divided in the following categories: metabolic, genetic, environmental, and others. These systemic factors are summarized in Table 1.

Metabolic factors

Physiologic and pathologic endocrine changes have long been established as significant modifying factors in the expression of gingivitis (Sooriyamoorthy & Gower 1989). Among physiologic changes, the variation in sex hormone levels occurring during puberty (Mombelli et al. 1989, Bimstein & Matsson 1999) and pregnancy (Hugoson 1971) have been shown to alter the plaquegingivitis relationship. With the onset of puberty, both female and male children exhibit significantly increased gingival bleeding (mean whole mouth score and percent of bleeding sites) that reaches a peak after 1-5 years, without any noticeable changes in plaque levels (Mombelli et al. 1989). Pregnancy was one of the

Table 1.	Systemic	factors	shown	to	modulate
gingiviti	s expressi	on			

Factor/condition	Gingival inflammation modulation
Metabolic	
puberty	+
pregnancy	+
diabetes	+
Genetic	
Down's syndrome	+
Environmental	
smoking	_
vitamin C deficiency	+
antibiotics	_
calcium channel blockers	+
corticosteroids	_
cyclosporin	+
NSAIDs	_
phenytoin	+
Other	
immune (PMN) deficiency	+
leukemia	+
HIV/AIDS	+
psychological stress	+

⁺Heightened gingival response to plaque.
– Reduced gingival response to plaque.
NSAIDs: non-steroidal anti-inflammatory drugs.
PMN: polymorphonuclear granulocytes.
HIV: human immunodeficiency virus.
AIDS: acquired immune deficiency syndrome.

first conditions identified as impacting gingivitis (Löe & Silness 1963, Silness & Löe 1964). Experimental gingivitis studies that compared the same women during pregnancy and 6 months postpartum indicate that, without any significant changes in the amount of plaque accumulated during the two phases, there is more swelling, redness, and bleeding during pregnancy (Raber-Durlacher et al. 1994). Besides puberty and gingivitis, the physiologic changes in hormone levels observed during the menstrual cycle have been associated with infrequent and subtle changes in gingivitis expression (Mariotti 1994). The evidence suggests that hormonal variations during the cycle do not affect clinically normal gingiva but do exacerbate existing chronic gingivitis (Holm-Pedersen & Löe 1967, Kovar et al. 1985, Niemi et al. 1986). The effect of the menstrual cycle has not been investigated in the experimental gingivitis model. Despite the significant impact of menopause on both the systemic and oral health of women (Samsioe 1995, Friedlander 2002), there are limited studies addressing the possible effects of the postmenopausal lack of estrogen or of hormone replacement therapy (HRT) on gingival inflammation. In a study on postmenopausal women receiving HRT, gingival bleeding was significantly decreased compared with the control group of postmenopausal women not receiving HRT; however, oral hygiene was also better in the group receiving HRT (Norderyd et al. 1993). The effect of menopause or HRT has not been investigated in the experimental gingivitis model. The significance of menopause as a metabolic factor modulating the clinical expression of plaque-induced gingival inflammation remains unknown. The effect of contraceptive agents on gingivitis expression, although related to changes in sex hormone levels, will be reviewed under environmental factors.

Among pathologic changes, diabetes is an endocrine condition with a wellcharacterized effect on gingivitis. Diabetic subjects, whether affected by the insulin-dependent (de Pommereau et al. 1992) or the non-insulin dependent (Cutler et al. 1999) form of the disease, have significantly higher gingival inflammation compared with non-diabetics with similar plaque levels.

In the course of implementing an experimental gingivitis trial, appropriate subject selection criteria (age, gender, health status) can eliminate the above factors. From a practical point of view, elimination of the menstrual cycle factor may be difficult, if not impossible, if both male and female subjects are expected to participate in the trial.

Genetic factors

As mentioned earlier, the gingival response to plaque may vary significantly between individuals with neither quantitative nor qualitative differences in plaque accumulation (Abbas et al. 1986a, Lie et al. 1995, 1998). Evidence from experimental gingivitis studies indicates genetic conditions can predispose subjects to increased susceptibility to plaque-induced inflammation. One such genetic condition is Down's syndrome (trisomy 21). In the course of experimental gingivitis trials, Down's syndrome subjects manifest more extensive and severe gingival inflammation and much earlier than age- and sexmatched genetically healthy controls, despite no differences in plaque accumulation rates (Reuland-Bosma et al. 1986, 1988). Although the specific gene(s) responsible for the various phenotypic changes observed in Down's syndrome subjects are not known, the genetic basis is unquestionable.

Recent preliminary evidence suggests that specific genetic characteristics may contribute to exacerbated gingival inflammation in response to plaque accumulation. Subjects positive for a specific interleukin-1 (IL-1) genotype exhibit increased susceptibility to experimental gingivitis (Goodson et al. 2000). The report was based on the comparison of seven periodontally healthy IL-1 genotype-positive subjects to 13 periodontally healthy IL-1 genotype-negative subjects. Significant differences in bleeding on probing between the two groups were found after 10 days of comparable plaque accumulation (Goodson et al. 2000). In contrast to these results, another experimental gingivitis report on ten periodontally healthy IL-1 genotype-positive subjects and ten periodontally healthy IL-1 genotype-negative subjects found no differences in gingival inflammation (including bleeding on probing) between the two groups after 21 days of plaque accumulation (Jepsen et al. 2003). Therefore, the significance of IL-1 genotype as a genetic factor modulating the clinical expression of plaque-induced gingival inflammation remains to be established.

The concept of host-dependent variation in gingivitis susceptibility is supported by the results of a published large-scale experimental gingivitis study. In 1979, Wiedemann et al. reported that in a group of 62 subjects, eight did not develop gingivitis within the time of the study (21 days), despite plaque accumulation. Another group of 25 subjects exhibited substantial gingival inflammation within 2 weeks. The remaining 29 subjects formed an intermediate group, which developed gingival inflammation by day 21 (Wiedemann et al. 1979). In a study involving 45 subjects, van der Weijden and colleagues found that 10 subjects consistently exhibited greater than average gingival inflammation, representing a "susceptible" group, while six subjects were consistently below average, representing a "resistant" group (van der Weijden et al. 1994). The difference in gingivitis susceptibility between the two groups could not be ascribed to qualitative plaque differences (Lie et al. 1995). In accordance with the above, Watts reported that only 18 of 29 subjects were consistent in bleeding from certain sites in response to plaque accumulation (Watts 1978). From the data reported in the above experimental gingivitis studies (Wiedemann et al. 1979, van der Weijden et al. 1994), one can estimate that approximately 13% of subjects represent a "resistant" group.

Collectively, the studies reviewed in this section are consistent with the hypothesis of a host-dependent and genetically based modulation of gingivitis expression and provided the motivation for initiating the present series of investigations.

Environmental factors

An environmental factor with welldocumented influence on the clinical expression of plaque-induced gingival inflammation is smoking. Smokers, when compared with non-smokers in experimental gingivitis studies, accumulate plaque at the same rate and exhibit significantly less gingival inflammation at similar plaque levels (Bergstrom & Preber 1986, Danielsen et al. 1990, Lie et al. 1998). When examined, no qualitative plaque differences were found between smokers and non-smokers during experimental gingivitis (Lie et al. 1998). It appears that the effect of smoking is mediated, at least in

part, by modulation of the local vascular response (Bergstrom et al. 1988). Because of the smoking-induced altered relationship between plaque and gingivitis (Bergstrom & Preber 1986, Danielsen et al. 1990, Lie et al. 1998, Müller et al. 2002), smokers have to be either excluded from experimental gingivitis trials or included as a separate group.

Various medications have been found to have adverse effects involving the gingiva. Several drugs (Seymour 1993), including anti-convulsants such as phenytoin (Angelopoulos 1975), anti-hypertensive calcium channel blockers such as nifedipine (Nery et al. 1995, O'Valle et al. 1995), and the immunosuppressant cyclosporin (Seymour & Jacobs 1992, O'Valle et al. 1995) cause severe gingival enlargement, a reaction related to the plaque-induced gingival inflammation (Seymour et al. 1996). In a way, the above medications can be considered as factors that exacerbate the plaque-induced gingival inflammatory response.

An equally large number of drugs have been shown to curtail plaqueinduced gingival inflammation. These include both steroidal (Sutton & Smales 1983, Vogel et al. 1984, Markitziu et al. 1990) and non-steroidal (Vogel et al. 1984, Heasman et al. 1993b, Jones et al. 1999) anti-inflammatory medications, whether systemically administered (Sutton & Smales 1983, Vogel et al. 1984, Markitziu et al. 1990, Heasman et al. 1993b) or topically applied (Vogel et al. 1984, Jones et al. 1999). These agents act by inhibiting the host response, thus inhibiting the clinical expression of gingivitis. Finally, systemic or topical anti-microbial agents inhibit plaque formation and the ensuing gingivitis. Various systemic antibiotics, such as erythromycin (Helovuo & Paunio 1989), metronidazole (Listgarten et al. 1979, Heijl & Lindhe 1980), clindamycin (Heijl & Lindhe 1980), vancomycin (Jensen et al. 1968, Heijl & Lindhe 1980), and tetracyclines (Listgarten et al. 1979, Atilla et al. 1996) have been shown to inhibit plaque accumulation and gingivitis development. Antibiotics such as tetracycline and vancomycin have also been used as a rinse, and have been shown to reduce plaque and inflammation (Löe et al. 1967). The use of an adjunctive antimicrobial product as a rinse [e.g. chlorhexidine (Bosman & Powell 1977, Hefti & Huber 1987, Cappelli et al. 2000), phenolic compounds (Mankodi et al. 1987, Brecx et al. 1990), amine fluoride/stannous fluoride (Hefti & Huber 1987, Brecx et al. 1990, Zimmermann et al. 1993)] or a toothpaste [e.g. triclosan-containing toothpaste (Lindhe et al. 1993)], to complement mechanical oral hygiene procedures has been associated with decreases in plaque and gingival inflammation.

In addition to their direct anti-microbial (anti-plaque) activity, some of these agents appear capable of modulating the host response by altering functions of several cell types involved in inflammatory reactions (Shapira et al. 1997, Yanagihara et al. 1997, Scaglione & Rossoni 1998, Solomon et al. 2000, Kuzin et al. 2001). Therefore, certain anti-microbial agents, whether systemically or locally used, can truly modulate the clinical expression of plaque-induced gingivitis, and not simply reduce gingival inflammation by reducing plaque levels. In this context, it is noteworthy that mouth irrigation as an adjunct to toothbrushing, even if it is performed with just water (Flemmig et al. 1990, Cutler et al. 2000), appears to inhibit plaque-induced inflammation, an effect attributed to reduction in the concentration of inflammatory mediators (such as IL-1 β and prostaglandin E₂) in gingival crevicular fluid (Cutler et al. 2000). Obviously, use of any of the above products needs to be excluded in an experimental gingivitis trial aimed at identifying the inherent characteristics of the host response to plaque accumulation.

Although effects similar to the ones associated with changes in sex hormone levels during pregnancy (see section on Metabolic factors above) have been reported in earlier studies on young women taking contraceptives (Lindhe et al. 1969, Kalkwarf 1978), recent studies suggest that the effect of newer contraceptive pills (which have much reduced hormone content compared with the ones used in the past) on gingivitis is practically nil (Klinger et al. 1998, Preshaw et al. 2001). Therefore, exclusion of females of childbearing age that use contraceptive medication from experimental gingivitis trials is not supported by current evidence.

Among the various nutritional factors examined during the last 35 years, only ascorbate (vitamin C) has been consistently shown to affect gingivitis expression. In both humans (Leggott et al. 1986, 1991) and non-human primates (Alvares et al. 1981) subclinical deficiency of ascorbate results in increased gingivitis, relative to non-deficient controls with similar plaque levels and the same type of microflora. Subclinical ascorbate deficiency resulted in 36% increase of mean GI score in non-human primates (Alvares et al. 1981), while in humans it reduced in half the percentage of sites that were healthy (non-bleeding) (Leggott et al. 1986). Elimination of a possible subclinical vitamin C deficiency among participants of an experimental gingivitis trial can be easily accomplished by nutritional supplementation.

Other systemic factors

Other systemic conditions identified as affecting the plaque-gingivitis relationship include neutropenia (Andrews et al. 1965, Rylander et al. 1975, Reichart & Dornow 1978), leukemia (Levin & Kennedy 1973, Bergmann et al. 1992), and human immunodeficiency virus/ acquired immune deficiency syndrome (Glick et al. 1990). Psychological stress, at least in adults (Deinzer et al. 1998, 1999, Vanderas et al. 1998), leads to heightened gingivitis expression. However, because psychological (academic) stress results in greater plaque accumulation, the relationship between stress and gingivitis may be mediated, at least in part, by this increase in plaque accumulation (Deinzer et al. 2001). Of the above conditions, stress is the one that is impractical, if not impossible, to eliminate during implementation of an experimental gingivitis trial. However, stress levels can be ascertained through appropriate questionnaires and taken into consideration during data analysis.

Rationale and significance

The susceptibility of an animal host to inflammation in response to various stimuli, including bacteria and bacterial products, varies significantly between strains of the same species (Allen et al. 1983, Yoshino et al. 1985). The physiological and/or genetic basis for the differences among strains has been partly elucidated (Sternberg et al. 1989, Listwak et al. 1999, Kandil et al. 2000), and genetic manipulations, e.g. introduction of a single immune response gene, can significantly alter such host inflammatory responses (Hammer et al. 1990, Tatakis et al. 2002). This biological phenomenon is consistent with the reported variation in plaqueinduced gingival inflammation in humans. However, in the case of humans, very limited information is available regarding the physiological/genetic reasons underlying this variation.

It was hypothesized that within a large-scale experimental gingivitis trial, conducted in a manner that minimizes confounding from parameters already known to influence gingivitis expression as described above, it would be possible to identify two subpopulations of individuals with inherently significantly different response to plaque-induced gingival inflammation (Trombelli et al. 2004b). Identification of these two groups of subjects should allow further study of other relevant clinical characteristics, such as their response to therapy (Trombelli et al. 2004a) and characterization of physiologic or genetic parameters that set them apart. Being able to identify physiologic and/ or genetic factors that account for increased susceptibility or resistance to the development of plaque-induced inflammation could have implications well beyond gingivitis prevention and management. Such potential far-reaching implications rest upon the unique relationship between gingivitis and periodontitis.

The epidemiology and natural history of gingivitis and periodontitis indicate that gingival inflammation is invariably a component of periodontitis and that gingivitis precedes the onset of periodontitis (Lindhe et al. 1973, Löe et al. 1986). However, it is also incontestable that all gingivitis cases do not proceed to develop into periodontitis (Lindhe et al. 1973, Listgarten et al. 1985, Löe et al. 1986, Prayitno et al. 1993). The reason for this is that plaque bacteria are necessary but not sufficient for the development of periodontitis, a susceptible host is necessary (Lindhe et al. 1973, Page & Schroeder 1982, Löe et al. 1986, Page 1999). The fact that gingivitis is a very poor predictor of periodontitis in persons under the age of 30 years (Prayitno et al. 1993) may also be related to the fact that gingivitis will manifest itself only days or weeks (Löe et al. 1965, Theilade et al. 1966) after the onset of plaque accumulation, while periodontitis is a condition that in the majority of cases requires far longer periods (few years to decades) (Lindhe et al. 1973, Löe et al. 1978) to evolve. At the present time, there are no reliable means to predict susceptibility to periodontitis. What the literature suggests, however, is that susceptibility to periodontitis may be linked to susceptibility to gingivitis.

Studies by van der Velden and colleagues indicate that susceptibility to experimental gingivitis (in terms of development of bleeding on probing) differs between two groups of patients with apparently different susceptibility to periodontitis. The group with greater periodontitis susceptibility exhibited greater susceptibility to gingivitis (van der Velden et al. 1985), a result that could not be explained by the age difference between the two groups (Winkel et al. 1987), or by the specific microbial (plaque) or immune (host) parameters examined in these subjects (Abbas et al. 1986b). The notion of a link between susceptibility to periodontitis and susceptibility to gingivitis is also supported by studies on patients with Down's syndrome. Down's syndrome patients exhibit severe periodontal disease early on (Reuland-Bosma & van Dijk 1986, Modeer et al. 1990), with no evidence of qualitative plaque differences between Down's syndrome patients and controls (Amano et al. 2001). and also manifest greater gingival inflammation much sooner than ageand sex-matched controls despite similar plaque accumulation (Reuland-Bosma et al. 1986, 1988). Therefore, it is possible that identification of factors related to increased susceptibility to gingivitis may help identify, at an early age, subjects at risk for periodontitis. If this were the case, it would permit the targeted application of public health resources and private practice-based preventive measures in a cost-effective manner.

Given the relationship between gingivitis and periodontitis (gingivitis precedes periodontitis, gingivitis is a prerequisite for periodontitis development), efforts to develop effective antigingivitis agents remains an important objective in the prevention of periodontitis (Robinson 1995). In the context of such efforts, the development of criteria/tests to identify subjects highly susceptible/resistant to gingivitis may provide a unique and powerful tool for designing and implementing clinical trials aimed at the evaluation of systemic and topical anti-microbial agents and host response modifying agents.

Conclusions

The experimental gingivitis model, initially developed by Löe and coworkers as a tool to demonstrate the relationship between plaque and gingivitis (Löe et al. 1965, 1967, Theilade et al. 1966), has been repeatedly validated and extensively used in the last 37 years for the study of gingivitis pathogenesis and evaluation of therapeutic interventions, mostly in smaller scale studies. Over the years, several factors that modulate clinical expression of gingivitis have been identified. However, the experimental gingivitis model has never been specifically used in a large-scale trial for the identification of individuals susceptible or resistant to gingivitis development and the subsequent characterization of inherent factors associated with gingivitis susceptibility.

Our choice of this model for the study of genetic and environmental factors modulating susceptibility to gingivitis was based on the following criteria: (a) the model has been extensively validated in humans, non-human primates, and other mammals, (b) the general qualitative and quantitative characteristics of the anticipated responses are well documented and provide an invaluable historical control, and (c) it is the only periodontal disease model that is fully reversible, thus precluding any permanent damage to the participants.

The accompanying papers report: (a) the findings of a large-scale experimental gingivitis trial that was designed and conducted taking into consideration the factors known to modulate the clinical expression of plaque-induced gingivitis, as reviewed here, and the identification of two subpopulations of subjects with significantly different gingival inflammatory response to plaque accumulation (Trombelli et al. 2004b), and (b) the response of these two subpopulations to self-administered plaque control (Trombelli et al. 2004a). Identification of host characteristics associated with resistance or high susceptibility to plaque-induced gingivitis remains the long-term goal of these studies.

Acknowledgments

We wish to express our gratitude to Dr. Arthur F. Hefti, College of Dentistry, The Ohio State University, for his critical review of the manuscript. This study was supported by funds from The Ohio State University College of Dentistry and by funds ex 60% 2000 from Ministero dell'Istruzione, dell'Università e della Ricerca, Italy.

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