

Changes in the subgingival biofilm composition after coronally positioned flap

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ABSTRACT

Objectives: This study evaluated the effects of coronally positioned flap (CPF) on the subgingival biofilm composition. **Material and Methods:** Twenty-two subjects with gingival recessions were treated with CPF. Clinical parameters were assessed before and at 6 months after surgery. Subgingival biofilms were analyzed by checkerboard DNA-DNA hybridization technique for 40 bacterial species. **Results:** Recession height, clinical attachment level and bleeding on probing improved significantly ($p < 0.05$) at 6 months post-CPF. The proportions of 10 periodontal pathogens and the proportions of red and orange complexes decreased at 6 months. **Conclusion:** In conclusion, CPF can induce beneficial effects on the composition of the subgingival microbiota after 6 months.

Key words: Bacteria. Microbiology. DNA probes.

INTRODUCTION

It has been well documented that the exposure of root surfaces as a result of gingival recession may result in tactile and thermal sensitivity, esthetic complaints⁵, and root surface carious lesions¹⁹. Numerous longitudinal human studies have demonstrated the efficacy and predictability of different techniques to esthetically, as well as functionally, correct gingival recession²⁷. Among the various techniques employed to correct gingival exposure, the coronally positioned flap (CPF), alone or combined with other procedures, e.g. subepithelial connective tissue graft (SCTG), has been one of the most widely used procedures in the treatment of Miller Class I gingival recessions^{2,9,14}. The coverage percentage of a previously exposed root surface is the primary clinical outcome used to evaluate the effectiveness of a mucogingival procedure. A high level of success, ranging from 60% to 99% of root coverage, has been reported after CPF in Miller Class I and II recessions with appropriate case selection^{18,20,27}. Miller¹⁶ (1988) suggested that a successful plastic procedure

for root coverage should also provide shallow sulcus and no bleeding on probing¹⁶. In general, reestablishment of esthetic and reduction of dentine sensitivity are the greatest advantages for patients that receive a plastic surgery for recession coverage.

To date, reports have only focused on the clinical outcomes of the mucogingival procedures to correct gingival recessions. To our knowledge the effects of these procedures on the composition of subgingival microbiota have not been investigated. Thus, the aim of this study was to evaluate whether the treatment of gingival recession with a coronally positioned flap (CPF) could interfere with the subgingival biofilm composition at 6 months after surgery.

MATERIAL AND METHODS

Twenty-two non-smoking, non-pregnant or lactant, periodontally and systemically healthy subjects from the Periodontics Department of Guarulhos University were enrolled in this study (January 2006 to June 2006). The following inclusion criteria were used: 1-Subjects with one Miller Class I gingival recession defect (≥ 2 mm and ≤ 4 mm) in

upper canines or premolars; 2- Keratinized tissue height of at least 2 mm; 3- Probing depth \leq 2 mm; 4- Absence of caries or restorations in the area to be treated; 5- Absence of pulpal pathology and severe occlusal interferences in the teeth to be treated; 6- Radiographic evidence of sufficient interdental bone; 7- Full-mouth plaque index¹ and full-mouth bleeding on probing index scores of \leq 20%; 8- Absence of previous mucogingival surgery at the defect; 9- Dental hypersensitivity and/or impaired esthetics associated with the recession.

Patients were informed of the characteristics of the study and gave their written consent to the described procedures. The study protocol was previously approved by the Institutional Committee of Ethics in Dental Research, in accordance with the Helsinki Declaration of 1975.

Coronally-Positioned Flap

The patients were submitted to CPF procedures, performed by one of the researchers. Right before the surgery, the exposure root surfaces were mechanically treated with manual curettes. After local anesthesia (2% lidocaine with 1:100000 epinephrine), a sulcular incision was carried out at the buccal aspect and two horizontal incisions were made at right angles to the adjacent papillae. Subsequently, two divergent oblique incisions at the mesial and distal aspects of the recession, apically extending beyond mucogingival junction (MGJ), completed a trapezoidal flap design. A periosteal elevator was used to carefully reflect an initial full-thickness flap until the mucogingival junction. After the mucogingival junction, a split thickness flap was dissected mesially, distally and apically, as necessary to release any tissue tension. The papillae adjacent to the involved tooth were deepithelialized. The flap was coronally displaced, completely covering the recession and fixed with a non-resorbable suture and a mattress sling suturing technique. Finally, interrupted sutures were placed at the vertical incisions to facilitate tissue stabilization. Subjects were instructed not to brush the teeth in the treated area but to rinse with chlorhexidine gluconate (0.12%) mouthwash twice a day for 2 weeks. Analgesics were prescribed to control postoperative discomfort. The sutures were removed after 14 days and, at teeth with recession-type defects, a coronally roll technique was prescribed.

Clinical Parameters

The following parameters were assessed on the buccal aspect of all studied teeth at baseline and at 6 months after the surgeries using a manual probe (UNC15, Hu-Friedy, Chicago, IL, USA): 1- Local plaque score (PL)¹: presence (1) or absence (0) assessed using a manual periodontal probe; 2- Local bleeding on probing (BOP): presence (1) or absence

(0) of bleeding of up to 15 seconds after gentle probing; 3- Probing depth (PD): distance between the gingival margin (GM) and the bottom of the gingival sulcus; 4- Recession height (RH): distance between cemento-enamel junction to the most apical point of the GM; 5- Clinical attachment level (CAL): distance between cemento-enamel junction to the bottom of the gingival sulcus; 6- Keratinized tissue height (KH): distance between the most apical extension of GM to the MGJ, chemically disclosed with Schiller's iodine solution; 7- Keratinized tissue thickness (KT): measured at a mid-point location between GM and MGJ by penetrating the probe into the tissue and recorded to the nearest 0.5 mm.

The assessed clinical parameters were arranged in order to obtain: A- Recession reduction (RR): calculated as [preoperative RH – postoperative RH]; B- CAL gain (CALG): calculated as preoperative CAL – postoperative CAL; C- Percentage of root coverage (RC): calculated as [preoperative RH – postoperative RH]/ preoperative RH X 100. The clinical parameters were assessed by the same periodontist who was trained and calibrated (s.e.m.=0.014).

Microbiologic Assessment

Sample collection

After removing the supragingival biofilm with sterile cotton pellets, subgingival biofilm samples were collected from the mid-buccal aspect of each experimental tooth using sterile curettes (5-6 mini-Gracey curette; UNC15, Hu-Friedy) and immediately placed in separate Microtubes containing 0.15 mL TE (10 mM Tris-HCl, 1 mM EDTA, pH 7.6). Freshly prepared 0.5 M NaOH (0.1 mL) was then added to each tube and the samples were dispersed using a vortex mixer. Biofilm samples were collected from the same sites at baseline and 6 months after CPF procedures.

Checkerboard DNA-DNA hybridization

Counts of 40 bacterial species were determined in each sample, using the checkerboard DNA-DNA hybridization technique²⁴. The analyses were performed at the Laboratory of Microbiology of Guarulhos University as previously described by Matarazzo, et al.¹⁵ (2008). The 40 reference strains used to develop the DNA probes are presented in Figure 1 according to bacterial complexes²¹⁻²².

Statistical Analysis

Data were analyzed using a software (SAS for Windows V8, SAS Institute Cary, Cary, NC, USA). First, the Kolmogorov-Smirnov test was used to evaluate the normality of the data. The frequency of detection of BOP (1) and PL (1) was determined for each period. The statistical significance of the differences for BOP (1) and PL (1) over time was evaluated by Chi-square and Fisher tests. In

Species	Strain	Species	Strain
Blue complex		Orange complex	
<i>Actinomyces gerencseriae</i>	23860a	<i>Eubacterium nodatum</i>	33099a
<i>Actinomyces israelii</i>	12102a	<i>Fusobacterium nucleatum ss. nucleatum</i>	25586a
<i>Actinomyces naeslundii sp. 1</i>	12104a	<i>Fusobacterium nucleatum ss. polymorphum</i>	10953a
<i>Actinomyces naeslundii sp. 2</i>	43146 a	<i>Fusobacterium nucleatum ss. vincentii</i>	49256a
Purple complex		<i>Fusobacterium periodonticum</i>	
<i>Actinomyces odontolyticus</i>	17929a	<i>Parvimonas micra</i>	33270a
<i>Veillonella parvula</i>	10790a	<i>Prevotella intermedia</i>	25611a
Yellow complex		<i>Prevotella nigrescens</i>	
<i>Streptococcus gordonii</i>	10558a	<i>Streptococcus constellatus</i>	27823a
<i>Streptococcus intermedius</i>	27335a	Red complex	
<i>Streptococcus mitis</i>	49456a	<i>Tannerella forsythia</i>	43037a
<i>Streptococcus oralis</i>	35037a	<i>Porphyromonas gingivalis</i>	33277a
<i>Streptococcus sanguinis</i>	10556a	<i>Treponema denticola</i>	B1b
Green complex		Other species	
<i>Aggregatibacter actinomycetemcomitans a+b</i>	43718/29523 a	<i>Eubacterium saburreum</i>	33271a
<i>Capnocytophaga gingivalis</i>	33624a	<i>Gemella morbillorum</i>	27824a
<i>Capnocytophaga ochracea</i>	33596a	<i>Leptotrichia buccalis</i>	14201a
<i>Capnocytophaga sputigena</i>	33612a	<i>Prevotella melaninogenica</i>	25845a
<i>Eikenella corrodens</i>	23834a	<i>Propionibacterium acnes</i>	11827/11828a
Orange complex		<i>Selenomonas noxia</i>	
<i>Campylobacter gracilis</i>	33236a	<i>Streptococcus anginosus</i>	33397a
<i>Campylobacter rectus</i>	33238a	<i>Treponema socranskii</i>	S1b
<i>Campylobacter showae</i>	51146a	<i>Neisseria mucosa</i>	19696 a

Figure 1- Bacterial strains employed for the development of the DNA probes
a- ATCC (American Type Collection, Rockville, MD); b- Forsyth Institute, Boston, MA

addition, Wilcoxon test was carried out to evaluate changes in PD, KT, KH, RH and CAL.

The proportions of each species and of each microbial complex were determined for each experimental site and averaged within baseline and six-month periods. The significance of differences between baseline and 6 months in mean proportion of each species and of the different microbial complexes was determined using the Wilcoxon signed-rank test.

Adjustments were made for multiple comparisons as described by Socransky, et al.²³ (1991) when the mean proportions of individual species were evaluated. The significance level established for all analyses was 5% ($p < 0.05$).

RESULTS

Twenty-two patients, 18 females and 4 males, aged between 25 and 60 years (mean age 42.66 ± 12.01) were included in the present study. Twenty-two maxillary Miller Class I gingival recession defects, one from each patient, were treated: 4 right canines, 1 left canine, 8 right first premolars, 7 left first premolars and 2 left second premolars.

The mean values of the clinical parameters at baseline and at 6 months post-surgery are summarized in Table 1. No significant differences were observed for PD, KT and KH ($p > 0.05$) between experimental periods. In addition, no significant changes in the frequencies of PL were observed

Table 1- Clinical parameters at baseline and at 6 months after coronally positioned flap

Clinical parameters	BASELINE	6 MONTHS
PD (mm)	1.4 \pm 0.5	1.2 \pm 0.5
KT (mm)	0.93 \pm 0.34	0.88 \pm 0.28
KH (mm)	3.18 \pm 0.91	3.40 \pm 0.94
RH (mm)	2.75 \pm 0.55 ^a	0.55 \pm 0.69 ^b
CAL (mm)	4.13 \pm 0.72 ^a	1.75 \pm 0.91 ^b
Number of site with:		
PL	12	7
BOP	9 ^a	0 ^b

Data are means \pm standard deviation. No statistically significant differences for PD, KT and KH were determined by Wilcoxon test ($p > 0.05$). Different letters (a,b) mean differences for RH and CAL determined by Wilcoxon test ($p = 0.0001$) and for BOP determined by the Chi-square or Fisher tests between the two time points. No statistically significant differences for PL was found by Chi-square or Fisher tests ($p > 0.05$).

PD- probing depth; KT-keratinized tissue thickness; KH-keratinized tissue height; RH-recession height; CAL-clinical attachment level; plaque accumulation (PL); bleeding on probing (BOP).

between baseline and six months ($p > 0.05$). RH, CAL and BOP levels improved significantly at 6 months

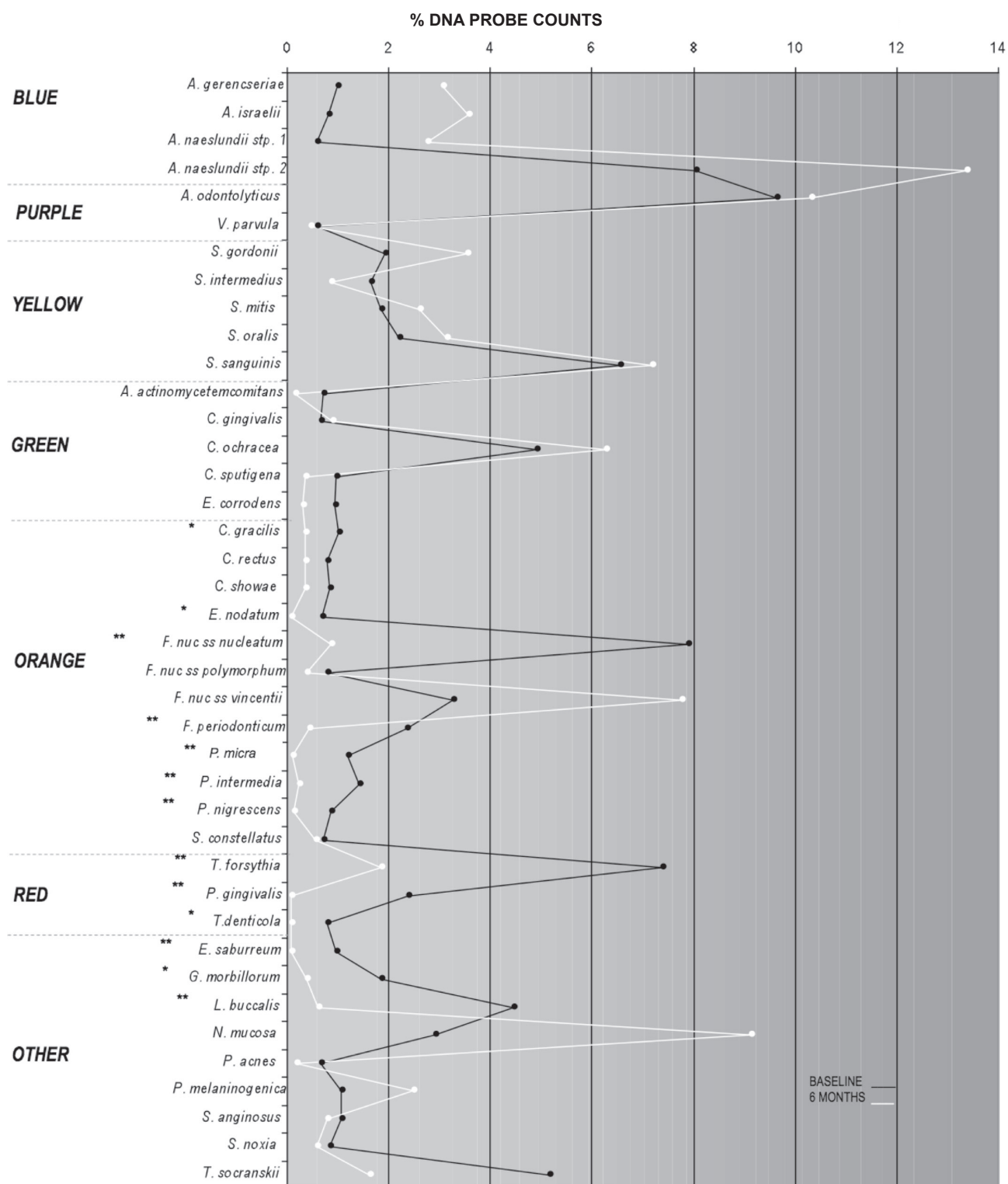


Figure 2- Mean proportions of 40 subgingival bacterial species at baseline and at 6 months after coronally positioned flap. The significance of differences in mean proportions between baseline and 6 months for each species was tested using the Wilcoxon signed-rank test after adjusting for multiple comparisons (*= $p < 0.05$, **= $p < 0.01$). The black line represents the mean data at baseline and the white line represents the mean data at 6 months post-surgery

($p < 0.05$).

At six months, the mean RR, CALG and RC were respectively, 2.2 ± 0.6 , 2.4 ± 1.0 and $81.64\% \pm 21\%$. Twelve sites (55%) achieved complete RC, 7 sites (32%) reached from 60% to 100% and only 3 sites

(13%) showed less than 60% of coverage.

The mean proportions of the 40 species evaluated in subgingival biofilm at baseline and at 6 months after CPF are presented in Figure 2. The results revealed that the proportions of 13 species

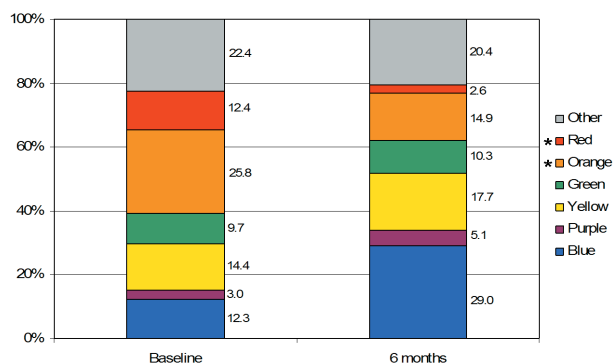


Figure 3- Mean proportions of microbial complexes (12) at baseline and at 6 months after coronally positioned flap. The significance of differences was tested using the Wilcoxon signed-rank test (* $p < 0.05$)

decreased at 6 months after surgery ($p < 0.05$), including 3 pathogens of the red complex (*Tannerella forsythia*, *Porphyromonas gingivalis*, *Treponema denticola*) and 7 putative periodontal pathogens of the orange complex (*Fusobacterium periodonticum*, *Parvimonas micra*, *Prevotella intermedia*, *Prevotella nigrescens*, *Eubacterium nodatum*, *Fusobacterium ss. nucleatum nucleatum* and *Campylobacter gracilis*) There was also a trend towards an increase in the proportion of some beneficial microorganisms of the blue, purple, and yellow complexes, such as *Actinomyces gerencseriae*, *Actinomyces israelii*, *Actinomyces naeslundii* 1 and 2, *Streptococcus gordonii*, *Streptococcus mitis* and *Streptococcus oralis*.

Mean proportions of microbial complexes at baseline and at 6 months after CPF are presented in Figure 3. A significant decrease in the proportion of red (from 12.4% to 2.6%) and orange (from 25.8% to 14.9%) complexes was observed at 6 months after CPF ($p < 0.05$). A tendency for an increase in the proportion of beneficial complexes (blue, purple, yellow and green) was also observed post-therapy. This was particularly noted for the blue complex, which accounted for 12.3% of the evaluated species at baseline and for 29% at 6 months after CPF.

DISCUSSION

Despite the common clinical application of periodontal plastic therapies, to our knowledge, no study to date has described the microbiological impact of these treatments. Therefore, this is the first study to evaluate whether the treatment of gingival recession with a CPF could influence the subgingival biofilm composition at 6 months following surgery. In addition, CPF clinical effectiveness was determined as a function of root coverage percentage and periodontal tissue health.

It has been recognized that the success of a periodontal plastic procedure is reached when the

gingival margin is at the cemento-enamel junction with sulcus depth ≤ 2 mm, no bleeding on probing and with presence of remained clinically attached gingiva¹⁷. In the present study, the exposed root surfaces demonstrated improvements in clinical outcomes in terms of recession height reduction (RR), clinical attachment gain (CALG), shallow probing depth (1.2 ± 0.5), reduced frequency of bleeding on probing (BOP) and unchanged KT and KH at 6 months post-surgery. In addition, as previously described^{2-3,10}, the mean percentage of root coverage ($81.64\% \pm 21\%$) observed in the present study confirmed that CPF is a predictable procedure to treat Miller Class I mucogingival defects.

Reports describing the composition of subgingival biofilm associated with gingival recessions are very scarce. A previous report demonstrated that subgingival biofilm of buccal sites with progressive gingival recession is colonized by gram positive species and, therefore, the microbiota resembled that of healthy sites. None of the predominant species, including *A. naeslundii*, *S. oralis* and *A. israelii* has been described as periodontal pathogens²⁶. These microbiological data are consistent with clinical and epidemiological observations that suggest a major traumatic, not infectious, etiology for gingival recessions^{4,11}. Contrary to the above-mentioned findings²⁶, in the present study, some of the predominant species observed on subgingival biofilm of the initially selected recessions, including *T. forsythia* and *F. nucleatum ss. nucleatum*, have been described as periodontal pathogens^{12,22,24,28}. A possible explanation for these contradictory results may be the fact that, in contrast to the aforementioned study²⁶ that found no recessions with BOP, 45% of our recessions presented BOP on the beginning of the study. Various microbiological studies have reported a range of gram negative species at bleeding sites, including *F. nucleatum* species, *T. forsythia*, *P. intermedia*, *P. nigrescens* and *P. gingivalis*^{6,25-26}. Additionally, it has been recognized that the three red complex species (*T. forsythia*, *P. gingivalis*, *T. denticola*) and various species of the orange complex are related to the presence of BOP^{7,21}.

In addition to BOP reduction and CALG, our longitudinal analyses demonstrated a decrease in the proportion of some pathogens and a trend for an increase in the proportions of health-associated microorganisms after CPF. Although the presence of BOP is not a reliable predictor or indicator for additional periodontal attachment loss^{7-8,13}, it has been demonstrated that the presence of some periodontal pathogens (e.g. *T. forsythia*) may be associated with sites converting to periodontal disease⁷. Thus, the microbial changes observed in the subgingival biofilm after CPF, especially those

related to the reduction of pathogens, seem to be a previously unreported advantage of mucogingival therapy. One could argue that these microbial changes would be related to the oral hygiene instruction and not to the CPF. Because of ethical reasons, we did not include a contralateral tooth that received only hygiene instruction to observe the effect of brushing technique alone on the subgingival composition. Although it was not observed a statistically significant reduction on visible plaque accumulation at 6 months post-surgery, we speculated that our clinical and microbiological findings could be due to the improvement of hygiene in the treated areas due to the surgery. Some hypothesis can be made in an attempt to explain the improvement of biofilm control after CPF: the reduction of dentin hypersensitivity; the shift of the gingival margin from apical to coronal position that could aid the brushing of the dento-gingival interface of the treated teeth; the increased motivation for hygiene since the volunteers were enrolled in a clinical study; and/or the fact that the subjects were submitted to a surgical procedure and, thus, they were concerned in preserving the initial benefits achieved.

CONCLUSION

It may be concluded that coronally positioned flap can have a beneficial effect on the composition of the subgingival microbiota, reducing the periodontal pathogen proportions in a six-month evaluation, maybe by the improvement of hygiene control in the treated area. Thus, besides to the esthetic improvement and the possible reduction in sensitivity, favorable changes in the subgingival biofilm adjacent to the gingival tissue seem to be an additional advantage for coronally positioned flap.

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