Osseointegrated implants placed at supracrestal level may harbour higher counts of *A. gerencseriae* and *S. constellatus* – a randomized, controlled pilot study

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**Purpose**: This study aimed at evaluating the bacterial colonization in dental implants inserted in the crestal or supracrestal position and correlated it to radiographic bone measurements.

**Methods**: Thirty-five implants with regular platform in nine patients (mean age 62.4 ± 11.2 years) were inserted either at the bone crest level (control group) or at a suprecrestal level (test group). Radiographic examination was performed at baseline (implant installation) and after 6 months. Clinical and microbiological data were collected after 6 months. Digital radiography was used to assess bone remodeling (marginal bone loss and optical alveolar density). Bacterial profile was analyzed by checkerboard DNA–DNA hybridization, including a panel of 40 bacterial species.

**Results**: After 6 months, there were significantly higher counts of *Actinomyces gerencseriae* (*p* = 0.009) and *Streptococcus constellatus* (*p* = 0.05) in the test group. No significant differences between test and control groups were observed for marginal bone loss (*p* = 0.725) and optical alveolar density (*p* = 0.975). Probing depth was similar in both groups.

**Conclusion**: Significantly higher counts of *A. gerencseriae* and *S. constellatus* were found in implants placed at the supracrestal level compared to the ones placed at the bone level. No relation was found between the installation level of dental implants and peri-implant bone remodeling.

**Keywords**: dental implants; bacteria; Actinomyces; Streptococcus; bone remodeling

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Replacing missing teeth by dental implants represents a breakthrough in dentistry, being a well-accepted treatment modality (1) and one of the first treatment options for rehabilitation of edentulous patients (2). Despite high successful rates achieved by implant-supported restorations and osseointegration of different implant designs (3–5), some factors are still not elucidated, especially concerning the bacterial colonization that might influence bone remodeling around dental implants.

At the moment of installation, pristine implant surfaces are devoid of an indigenous microbiota. The colonization of the implant-supported restoration leads to an increase in peri-implant inflammation, altering the local habitat. In turn, the composition of the microbiota adjacent to implants is influenced by the local environment established at the interface between peri-implant mucosa and implant surface (6). Peri-implant microbiota composition has been evaluated along with clinical parameters. Some studies have shown that an increase in proportions of spirochetes and motile organisms was associated with an increase in probing depth around implants (7, 8). Other studies have found greater levels of periodontal pathogens around implants with marginal bone loss (9, 10).

A supracrestal position of the implant platform may favor the establishment of a biological width, and also
push aside the microgap and its bacterial contamination from peri-implant bone crest, besides reducing the inflammatory infiltrate. Studies have shown that the absence of a microgap at the bone crest level, obtained with a supra-crestal installation, was associated with reduced inflammatory peri-implant cells and with minimal bone loss (11, 12). Authors have shown that minor bone loss is observed when the microgap is coronally moved away from the bone crest. In turn, the contrary may be observed as far as the microgap is moved apically (13).

Peri-implant bone remodeling is one of the healing phases, which involves hemostasis, an inflammatory phase, a proliferative phase, and finally, a remodeling phase (14), and occurs since the implant is exposed to the oral environment, in a second surgical procedure or when the prosthetic component is immediately connected after implant installation (immediate loading protocol) (15). The remodeling process implicates marginal bone resorption both horizontally and vertically that may be affected by several biological and mechanical factors, for instance, a traumatic surgical technique, excessive loading conditions, peri-implant and microgap bacterial colonization, and implant-neck geometry (6, 15, 16). Thus, the aim of this study was to evaluate the bacterial colonization in dental implants inserted in crestal or supracrestal position and correlated it to radiographic measurements of bone remodeling.

Methods
A total of 10 patients were included in this study. They were selected from the Prosthodontics Clinic at the Rio de Janeiro State University. All patients signed a written informed consent. The study protocol was approved by the Ethics Committee in Research of the Pedro Ernesto University Hospital – Rio de Janeiro State University. Patients were totally edentulous; had absence of medical conditions that compromise the immunological status; wore complete superior and inferior dentures, which were aesthetically and functionally adequate; and had enough bone to support four dental implants with 4.1 mm in diameter in the anterior portion of the mandible between the mental foramina. Exclusion criteria were use of anti-inflammatory peri-implant cells and with minimal bone loss (11, 12). Authors have shown that minor bone loss is observed when the microgap is coronally moved away from the bone crest. In turn, the contrary may be observed as far as the microgap is moved apically (13).

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Clinical examination
The same examiner assessed probing depth using a calibrated periodontal probe (Colorvue®, Hu-Friedy, Chicago, IL) at the mesial and distal sites of each implant.

Microbiological analysis
Two sites for each implant were selected. After removing the supragingival plaque, the most apical subgingival biofilm was collected using sterile Teflon curettes (Implacare™ IC4R/4L – Hu-Friedy Co., Chicago, IL). Samples were placed in separate microtubes containing 0.15 mM TE (10 mM Tris-HCl and 1 mM EDTA, pH 7.6). Freshly prepared 0.5 M NaOH was added to each tube so that the bacterial DNA remained viable. Samples were kept under −20°C until analysis.

Counts of 40 bacterial species were determined in each sample using checkerboard DNA–DNA hybridization. The analyses were performed at the Laboratory of Microbiology of Guarulhos University as previously described (17). A single blinded examiner performed radiography films readings twice in two different days. Each signal produced by a probe in the sample was compared, in its intensity, to the signal produced by the same probe in two
control lines that contained $10^5$ and $10^6$ bacteria. Thus, score 0 was attributed to sample when no signal was detected; score 1 corresponded to a signal with intensity lower than $10^5$ control; score 2 corresponded to $10^5$ cells; score 3 corresponded between $10^5$ and $10^6$ cells; score 4 corresponded to approximately $10^6$ cells; and score 5 was attributed to more than $10^6$ cells.

Radiographic bone assessment

Radiographic assessment was performed in the mesial and distal sites of each implant. Periapical radiographies were obtained at baseline and after 6 months, by paralleling distal sites of each implant. Periapical radiographies were used to assess linear correlations between variables. Statistical significance was set at 0.05.

Statistical analysis

Since many implants were analyzed in each patient, imposing an internal dependence of observed data, generalized linear models with generalized estimation equation were used to analyze marginal bone loss and optical alveolar density variations. In these models, the statistical significance was obtained by Wald statistics. Pearson and Spearman’s rank coefficients, whenever appropriate, were used to assess linear correlations between variables. Statistical significance was set at 0.05.

Based on the difference observed for the counts of Actinomyces gerencseriae, this study presented power >80% with 16 implants per group and $\alpha = 0.05$.

Results

From the 10 patients who were included, 9 remained in the T2 analysis. One patient from test group dropped out of the study. One implant was lost in a patient from the control group. Mean age ($\pm$ standard deviation) of the nine patients (three males and six females) was 62.4 $\pm$ 11.2 years.

Checkerboard DNA–DNA hybridization

Regarding the microbial profile, the test group showed higher counts of A. gerencseriae ($p = 0.009$) and Streptococcus constellatus ($p = 0.05$). There was also a trend to higher counts of S. oralis ($p = 0.06$) and Gemella morbillorum ($p = 0.08$) in the test group. No bacterial count was significantly higher than in the control group. Microbiological data are depicted in Fig. 1.

The median total bacterial load was $59.5 \times 10^5$ ($32.5–93.0 \times 10^5$) in the test group and $57.0 \times 10^5$ ($36.0–65.0 \times 10^5$) in the control group, without significant difference between the groups.

Radiographic bone analysis

Bone loss and optical alveolar density in T1 and T2 are presented in Table 1. Both groups showed a statistically significant increase in radiographic bone loss from T1 to T2 ($p < 0.001$). There was no significant difference in optical alveolar density from T1 to T2 in the test and control groups ($p > 0.05$). There were no significant differences between groups in both parameters at T2 ($p > 0.05$).

In T2, the probing depth was 2.32 mm ($\pm 0.37$) and 2.38 mm ($\pm 0.71$) for the test and the control groups, respectively. There was no significant difference in probing depth between groups ($p > 0.05$).

Correlational analysis

No significant correlation was established between bone loss ($\Delta T2 – T1$) and variation in optical alveolar density ($\Delta T2 – T1$) in both groups. Also, there was no significant correlation between radiographic measures and probing depth. There was no significant correlation between bacterial profile and radiographic measures and between bacterial profile and probing depth.

Discussion

There is a discordance regarding insertion level of dental implants related to bone crest (and consequently the localization of the microgap) and its influence on peri-implant bone remodeling. Thus, this study was performed to evaluate the effect of the implant insertion level on bacterial profile and peri-implant bone remodeling. We found higher counts of A. gerencseriae and S. constellatus in the supracrestal group. There were also trends for higher counts of S. oralis and G. morbillorum in the supracrestal group. The total bacterial count was not significantly different between the groups. Canullo et al. (6) have found no significant differences in bacterial colonization between
implants restored with a traditional approach and those restored with a platform-switching approach. Our patients were fully edentulous, while Canullo’s patients were partially edentulous. This may account for the conflicting results. A systematic review has shown a difference in submucosal peri-implant microbiota between fully edentulous and partially edentulous patients (18).

Fig. 1. Bacterial profiles in test and control groups (*p < 0.05).

Table 1. Bone remodeling in baseline (T1) and after 6 months (T2) in test and control groups

|                         | Test group (n = 16) | Control group (n = 19) |
|-------------------------|---------------------|------------------------|
| Linear measurement      |                     |                        |
| T1                      | 4.00 (± 0.33)       | 4.31 (± 0.31)**        |
| T2                      | 4.68 (± 0.48)       | 4.93 (± 0.33)          |
| Bone loss (A T2 – T1)   | 0.68 (± 0.36)*      | 0.62 (± 0.54)*         |
| Optical alveolar density|                     |                        |
| T1                      | 53.4 (± 18.5)       | 61.3 (± 16.0)          |
| T2                      | 57.0 (± 28.1)       | 64.6 (± 30.4)          |

Data are presented as mean (standard deviation). p-Value calculated using Wald test applying generalized estimating equations. *p < 0.001 for intragroup difference. **p < 0.001 for the difference between test and control group.
implants (21). On the other side, higher proportions of *Actinomyces* were shown in healthy implants (22). However, we found no significant correlation between bacterial profiles and bone remodeling and between bacterial profiles and probing depth. In agreement with our finding, Canullo et al. (6) also found no significant correlation between mean bone loss and levels and proportions of subgingival species in the peri-implant plaque.

Some studies have shown that the absence of microgap at the bone level with non-submerged implant systems would result in minimal peri-implant bone loss (4, 11, 23). Meanwhile, other studies have found advantages with subcrestal placement of dental implants (24, 25). Our study found no statistically significant difference when bone remodeling was evaluated in two-piece implants inserted 1 mm supracrestally and at the bone level. This result is in accordance with other studies, which found no effect of the microgap location in bone remodeling (26–28).

Several factors may account for the conflicting results, as *interface implant/abutment and healing time*. In our study, the interface implant/abutment in the supracrestal group was located 1 mm above the bone crest. Boyneugri et al. (1) have located the microgap 2.8 mm above the bone crest. Piattelli et al. (13) located the interface implant/abutment 1–2 mm above the bone crest and found that this position was favorable to a minor bone loss in this group. In addition, our study used an immediate loading protocol. On the other hand, healing time before loading varied from 3 to 6 months in other studies (1, 11, 12), which may account for the different results. However, Guruprasada et al. (29) and Piattelli et al. (13) found no significant differences in peri-implant bone remodeling comparing conventional loading and immediate loading protocols.

We also found no significant difference in probing depth between the groups. This result is in agreement with that showed by Boyneugri et al. (1).

A reduced sample and analysis of multiple implants in the same patient are limitations of our study, which imposed an internal dependence in the observed data. Nevertheless, we used generalized linear models with a generalized estimation equation to take into account this dependence. Also, we assessed a limited number of bacterial species, and modern sequencing techniques have been shedding light on the peri-implant microbiome complexity (21, 30). Studies with greater sample sizes and longer follow-ups are needed to investigate if these results and their clinical implications will be maintained. In conclusion, significantly higher counts of *A. gerencseriae* and *S. constellatus* were found in implants placed at the supracrestal level compared to the ones placed at the bone level. No relation was found between the installation level of dental implants and peri-implant bone remodeling.

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**Conflict of interest and funding**

The authors declare no conflict of interest.

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