

Degree of bacterial microleakage at the implant-abutment junction in Cone Morse tapered implants under loaded and unloaded conditions

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ABSTRACT

Purpose: Different results have been reported on the internal colonization of Cone Morse connections under in vitro dynamic loading. The aim of the present in vitro study was to evaluate the bacterial leakage in Cone Morse implant-abutment connections, both under loaded and unloaded conditions.

Methods: A total of 20 implants with a Cone Morse taper internal connection were used in this study. Ten were loaded under a special testing equipment (Test Group), while 10 were left unloaded (Control Group). The inner part of all implants was inoculated with 0.1 µl of a viable *Enterococcus faecalis* suspension. A force of 120 N was applied to the loaded implants, for a total of 500,000 cycles at 1 Hz. All the samples were checked daily, for a total of 14 days, and presence or absence of turbidity recorded.

Results: In the unloaded assemblies, bacterial contamination was found in 2 out of 10 implant-abutment junctions, on the 12th and 13th days. In the loaded implant-abutment connections, bacterial contamination was found in 2 out of 10 implant-abutment assemblies, on the 13th and on the 14th days.

Conclusions: The resistance of the Cone Morse implant-abutment junction reported in the literature and confirmed in the present study, where no differences in the percentages of microbial leakage were found in assemblies unloaded and in those subjected to a dynamic loading procedure, could help to explain the histological results in man of a lack of peri-crestal bone resorption in Cone Morse implants, placed below the level of the alveolar crest.

Keywords: Bacterial leakage, Cone Morse connections, Dental implants, Dynamic loading, Implant-abutment junction

Introduction

The presence of an implant-abutment junction (IAJ) has been correlated with the presence of microorganisms (1). These bacteria may harbor inside the microgaps that can be found in the assemblies and they have been reported to be the cause of the inflammation of the soft tissues (abutment Infiltrated Connective Tissue or ICT) facing the IAJ, potentially producing a bone resorption of the surrounding peri-crestal bone (2-5). Bacterial migration has been correlated with tightening torque forces, micromovements of the different components during the masticatory cycles and the accuracy of the fit between implant and abutment (1). Even if the Cone Morse implant-abutment connection is considered

to be the most stable and the most capable to resist penetration of microorganisms when compared with flat-to-flat or tube-in-tube connections, their possible bacterial colonization is not yet well understood (6, 7). In all implant-abutment different connections the size of the IAJ microgaps can be increased during the loading of the assemblies (2, 8, 9), introducing a so-called pumping effect (10). Higher quantities of microorganisms have been reported for loaded than unloaded external hexagon and internal hexagon implants (9). On the other hand, a minimal contamination has been reported for Cone Morse connections (2, 11). Different results have been reported on the internal colonization of Cone Morse connections under in vitro dynamic loading. do Nascimento et al (9) found no significant differences in the amount of bacteria, comparing unloaded and loaded groups, while, on the contrary, Koutouzis et al (10) found that the bacterial penetration was influenced by the loading conditions. One of 20 assemblies was found to be colonized by *Escherichia coli* in the unloaded group, while the number of colonized microgaps increased to four in the loaded group. The aim of the present in vitro study was to evaluate the bacterial leakage in Cone Morse implant-abutment connections, both under loaded and unloaded conditions.

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Methods and materials

A total of 20 implants with a Cone Morse taper internal connection (Universal II CM, Implacil, De Bortoli, Sao Paulo, Brasil) were used in this in vitro study. Ten were loaded under a special testing equipment (Test Group), while 10 were left unloaded (Control Group). To simulate cyclic loading and to mimic intraoral conditions, the Test implants were embedded in autopolymerizing acrylic resin with a custom-made stainless steel ring form. The implants were positioned using a parallelometer to ensure uniform alignment of all specimens. The implants were positioned so that the cyclic fatigue load was applied to the long axis of the implants. All embedded implants were sterilized by autoclave for 15 minutes at 121°C. After several trials, 0.1 µl was determined to be the ideal quantity of bacterial suspension for inoculation in the implant systems. *Enterococcus faecalis* (EF) is a Gram-positive coccus, nonmotile, facultatively anaerobic microbe, a human commensal and an important opportunistic pathogen inhabiting the gastrointestinal tracts, oral cavity and urinary tract, whose size ranges approximately 1.0-1.5 µm. This species is involved in the pathogenesis of secondary endodontic apical lesions and can also be found in root-filled teeth with no apical lesions and also in primary endodontic lesions. EF can survive extreme environmental conditions (acidic or basic pH, high salt concentration, the presence of heavy metals, aerobic and anaerobic). In fact, it fails to grow at 10°C and 45°C, at pH 9.6, 6.5% NaCl in broth, and it survives at 60°C for 30 min. This bacteria may invade the dentinal tubules for more than 200 microns and is able to survive without the support of other bacteria and can be resistant to a wide range of antibiotics. The inner part of ten implants was inoculated with 0.1 µl of a viable EF suspension with a 0.1 µl calibrated micropipette (Gilson, Middleton, WI, USA), with sterile gloves, under sterile conditions. A pure culture of EF (reference strain ATCC 29212) was used. For preparation of the bacterial suspension, the test organism EF was first plated onto MacConkey Agar without Crystal Violet (Oxoid LTD, Basingstoke, Hampshire, England) and then incubated for 24 hours at 37°C. Suspension was made from the culture by diluting a few colonies in nutrient solution (Nutrient Broth, Oxoid LTD, Basingstoke, Hampshire, England) for EF to a density of 0.5 McFarland Standard (1×10^8 colony forming units per mL - CFU/mL), confirmed by spectrophotometer analysis (Agilent Technologies 8453 UV, Santa Clara, USA) and subjected to a series (two series) of 10-fold dilutions in broth. As a positive control, a test tube was used with only nutrient solution and inoculated with 0.1 µl of EF. It showed bacterial growth with solution cloudiness, and this confirmed the viability of the microorganism throughout the experiment. As a negative control, an identified test tube was used, with only sterile nutrient solution. This was confirmed by the transparency of the solution and conventional microbial culturing techniques. Before the implants and abutments were connected, the inner part of implants was inoculated with bacterial suspension, under sterile conditions. After the implant inoculation, the abutment was carefully connected to the implant, according to the manufacturer's protocol, without touching the outer surface of the implant and using sterile gloves. An implant torque controller pre-calibrated at 30 N cm as manufacturer-recommended was

used to ensure proper seating torque for all implants (Fig. 1). Subsequently, implant systems were placed in sterile cylinders. The specimens were partially immersed in Nutrient Broth (Nutrient Broth, Oxoid LTD, Basingstoke, Hampshire, England), so that only half of each element, comprising the connection implant/abutment area, was immersed in the nutrient solution and subjected to the load (Fig. 2). The whole system was maintained at a temperature of 37°C. Then, the specimens were mounted, each at a time, in a chewing simulator (Figs. 3A and B). A cyclic fatigue load (Universal Testing Machine, LLOYD LR 30K, UK) was individually applied to each specimen with a stainless steel stem through the rubber ring. A force of 120 N was applied for a total of 500,000 cycles at 1 Hz, which is within the physiologic clinical range. The samples were checked daily, for a total of 14 days, and presence or absence of turbidity recorded. Such leakage caused bacterial colonization and resulted in a cloudy solution; 1 µl of the solution was drawn with sterile 1-time-use syringes and analyzed with a Gram stain and by colony morphology in MacConkey Agar without Crystal Violet (Oxoid LTD, Basingstoke, Hampshire, England), incubated at 37°C for 24 h to confirm the purity of the microorganism which had been inoculated in the inner part of the implant and determining the presence of EF. A resulting growth of EF, confirmed that bacteria had indeed escaped from the inner part of the implant along the tested interface into the surrounding solution. The same procedure was performed with the Control implants, but without the loading.

Results

The frequencies of contamination in both tested groups were shown in Table I. In the unloaded assemblies, bacterial contamination was found in 2 out of 10 IAJ, on the 12th and 13th days. In the loaded implant-abutment connections, bacterial contamination was found in 2 out of 10 implant-abutment assemblies, on the 13th and on the 14th days. No assemblies showed contamination of the outer portions. The positive controls remained positive. The negative controls remained negative.

Discussion

A gap of 1-5 microns has been reported to be present, in retrieved human implants, between the implant and the healing screw, and this space was found to be filled by bacteria and calculus (12). Bacteria were also present inside the most apical portion of the hollow part of the implant (12). In another histological study on retrieved human implants, it has been reported that bacteria were often present between the implant and the abutment and in the internal portion of the implants (13). These results, with the finding of hollow spaces, in screw-retained abutments, between the implant and abutment components with the presence of fluids and bacteria inside the internal hollow cavities of the implants, serve to validate, in man, the results obtained from in vitro studies or animal experiments (12). Moreover, in retrieved human implants, an inflammatory infiltrate was present below the implant-abutment microgap (12), and, in an experimental study in monkeys, many inflammatory cells were found below the



Fig. 1 - Abutment connected to the implant at 30 N cm.



Fig. 2 - Implant/abutment immersed in nutrient solution, drawn by sterile syringe and analyzed by culture method.

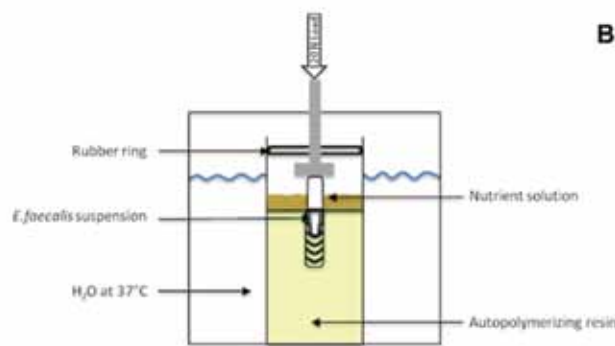


Fig. 3 - (A) Picture of the chewing simulator, (B) drawing of the chewing simulator: an implant was embedded in resin, immersed in a nutrient solution, inoculated with *E. Faecalis* suspension and subjected to cyclic fatigue load (120 N) with a stainless steel stem through the rubber ring.

TABLE I - Bacterial leakage in loaded and unloaded implants

Implants	Bacterial species	% of contamination	Days	Total
Loaded	<i>E. Faecalis</i>	2 out of 10	13 th and 14 th days	2 contaminated samples out of 10
Unloaded	<i>E. Faecalis</i>	2 out of 10	12 th and 13 th days	2 contaminated samples out of 10

implant-abutment microgap (5). There is evidence that with cyclic loading the microgap can increase in size, especially in external hexagon implants, facilitating the flow of fluids and bacteria (6, 9, 13-15). The results of the present study did not agree with the statement that overall loaded implants showed higher bacterial counts than unloaded implants (9). The present data may reflect the possible differences in the precision fit, in the degrees of taper and in the internal connection features of the implant components in the different implant systems (10, 14, 16). The reported average misfits between implant components vary greatly, from 1 to 100 microns (9). The occlusal loading of the implants produces bending forces and/or micromovements within the implant systems (9) increasing the microgap and promoting a higher fluid and bacteria infiltration (15), especially if imprecise machining of implant components or improper male-female adaptation exist (17). Cone Morse connections have been proposed to get a better stability, with reduced micromovements, of the components, providing a better bacterial seal (18). The bacterial microleakage can, probably, be reduced by obtaining an optimal component fit, a reducing or absence of micromovements, and a minimal prosthetic misfit (19-28). The length of the IAJ in Cone Morse implant-abutment assemblies could be a reason for the reported differences in the bacterial leakage (14). Implants that use mechanical frictional abutments seem to present precise adaptation in the deeper inner portions of the system, resulting in reduced micromovements (9).

Conclusion

The resistance of the Cone Morse IAJ reported in the literature and confirmed in the present study, where no differences in the percentages of microbial leakage were found in assemblies unloaded and in those subjected to a dynamic loading procedure, could help to explain the histological results in man of a lack of peri-crestal bone resorption in Cone Morse implants, placed below the level of the alveolar crest.

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