The influence of fixed orthodontic retainer on oral microbiota

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Abstract---Background: The oral cavity is a wide versatile microbial community in which many microbial species coexist in harmony. However, with certain circumstances, i.e. orthodontic appliances, pathogenic biofilm outgrowth occurs. The aim: The aim of this study was to compare opportunistic pathogens, associated with dental caries and periodontal disease in addition to Candida albicans, between healthy individuals and patient wearing fixed orthodontic retainers. Method: Fourty eight subjects were recruited andwere divided into retainer wearers and control cohort groups. Biofilm samples from the lingual surface of the mandibular anterior teeth of the control group and the fixed retainer surface were evaluated to detect Streptococcus mutans, Lactobacillus acidophilus, Aggregatibacter actinomycetemcomitans, Fusobacterium nucleatum and C. albicans. Additionally, plaque (PI) and gingival (GI) indices were measured at the mandibular anterior teeth in both groups. The results: The results showed a significant increase in total micobial count in retainer group (p<0.001). The prevalence of microbial cultivity was higher in retainer group with a significant increase in PI and GI. Moreover, there was a high correlation between these genera and the GI. The conclusion: fixed retainer may increase cariogenic and periodontal pathogens and compromise oral health. Further studies on improving the fixed retainer material and plaque control measures in retainer wearers are needed.

Keywords---Fixed retainer, opportunistic pathogens, periodontal pathogens, plaque and gingival index.
**Introduction**

Retention is the step of orthodontic treatment that aims to preserve teeth in their corrected positions after active orthodontic treatment \(^1,^2\). This is by preventing the teeth from rebound to their pretreatment positions by the action of periodontal, occlusal, soft tissue forces and continued dento-facial growth \(^3\). Retainers permit reorganization of the gingival and periodontal tissues, lessen the effect of changes due to growth, allow the adaptation of neuromuscular apparatus to the treated tooth position and maintain teeth with relatively high risk of relapse i.e. compromised plan for esthetic reasons \(^4\). There are basically two types of retainers, the removable and fixed orthodontic retainers. Occasionally, the removable orthodontic appliances may be beneficial in retention as opposed to intra-arch instability and may serve as retainers in patients with growth issues i.e. modified functional appliances or part-time headgear \(^5\). However, if the case requires a permanent retention, a fixed retainer could be used which may be a probable approach for intra-arch retention when irregularity in a specified area is likely to be an issue \(^6\). In spite of the priority of the fixed retainer for long term retention, patient satisfaction and appliance invisibility with less patient compliance, it has some drawbacks such as, bonding technique sensitive, time-consuming procedure i.e. passive positioning on the lingual surface, possibility of wire fracture or bond failure and the prospect of causing periodontal issues by compromising the oral hygiene \(^7\). Furthermore, the fixed retainer wire component can act as a shelter site for bacteria adhesion, biofilm and dental calculus formation that eventually leads to caries and periodontal diseases \(^8\). The oral ecosystem comprises of oral microflora, which harbors more than 700 different bacterial species in addition to fungi and viruses. Despite the fact that the bulk of these microbiota are approved as beneficial and commensally, part of them are in charge of oral infections which varying from tooth cavities to periodontal diseases and gum related lesions \(^9\). The major notorious bacteria for dental caries are *Streptococcus mutans*, and *Lactobacillus* \(^10,^11,^12\). The *S. mutans*; that is a normal oral cavity inhabitant; is the most cariogenic type of all oral streptococci that can cause initiation and progression of dental caries\(^13\). Another important participant of the oral microflora is the *Aggregatibacter actinomyces-evertum* which is considered as one of the initial colonisers for periodontal disease in juveniles (localized aggressive periodontitis LAP) and adolescent individuals \(^14\). Additionally, *Fusobacterium nucleatum*, is recognised as a key player in biofilm figuration by its role in "bridging microorganism" by which enhances the adhesion and incorporation of pathogens responsible for periodontal lesions into the oral biofilm \(^15,^16\). To the authors' knowledge, there is limited studies regarding the effect of fixed orthodontic retainer on the oral microbial ecosystem; therefore, the aim of this study was to compare the dental caries and periodontal pathogens i.e. *S. mutans*, *L. acidophilus*, *A. actino-actinomyces* and *F. nucleatum*, in addition to *C. albicans* in fixed orthodontic retainer wearers and the oral cavity of a control group matching the same inclusion criteria.
Materials and Method

Subjects

This cohort study involves 48 individuals that were divided into two groups, with an age range of 18-37 years, who were recruited from private orthodontic clinic in Baghdad city/Iraq. The first group consisted of 24 patients with fixed orthodontic retainer fitted on the lingual surfaces of the lower anterior teeth; whereas the other group consisted of 24 normal subjects matching the retainer group with regard to age, gender distribution and the inclusion criteria but without orthodontic appliances or retainers (17,18).

The criteria for participation were as followed: subjects fitted with multistrand 0.010*0.028 stainless steel fixed orthodontic retainers, (Braided Retainer Wire, Orthotechnology, Illinois, USA), for at least 3 months. Subject must be medically fit and healthy; had no signs of oral mucosal disease; had not received oral antimicrobials or antibiotic therapy for at least 3 months before data collection; not pregnant or lactating; and do not smoke (19,20).

The study was approved by the board ethical committee at the College of Dentistry/University of Baghdad. (issue no.182, date 28/1/2020). An information sheet and written informed consents was obtained from the patients/participants involved before patient recruiting process.

Methods

Sterile swabs sticks (Bhubaneswar, Odisha, India) were used to sample the lingual surface of the mandibular anterior teeth in the control group and the fixed retainer surface that attached to the mandibular anterior teeth in retainer group. In the retainer group, a braided stainless steel wire (Orthotechnology) was bonded to the lingual surface of the mandibular six anterior teeth using flowable composite adhesive (3M Unitek, Monrovia, California). Each swab was immediately inserted in a 10ml sterile tube (sterile plain tube, Biozek, Netherland) contained 5 ml of brain heart infusion broth (BHI, Oxoid, Leicestershire, UK), and were taken immediately to the laboratory for processing (21). The samples were vortex-mixed for 1 min to make a homogenous solution. Ten-fold serial dilution was carried out in phosphate buffered saline (PBS, Sigma Aldrich, Missouri,USA) before the samples were cultured onto blood agar (Oxoid, Basingstoke, UK) to obtain the total anaerobic bacterial counts; mitis salivarius agar (MS, Himedia, Mumbai, India), a selective medium for streptococci (22); tryptic soy serum Bacitracin- Vancomycin agar (PDH, England), a selective medium for A. actinomycescomitans (23,24); De Man-Rogosa-Sharpe broth (Oxoid), a selective medium for L. acidophilus (25); crystal violet-Erythromycin agar (CVE, avongem, UK), a selective media for F. nucleatum (26) and Soubraud Dextrous agar (Neogen, Heywood, UK) for C. albicans detection (27).

The plates and tubes were stored anaerobically in CO₂ incubator for 24 hrs, except tryptic soy serum Bacitracin-Vancomycin agar plates were kept for 72 hrs (28). To identify the microorganism, microbial growth by cultural characteristics (22), Gram stain and biochemical tests using catalase, oxidase and coagulase tests
was carried out\(^{(29)}\). In addition to that, Plaque index and Gingival index were recorded according to Løe and Silness Plaque Index (PI) and Gingival Index (GI)\(^{(30)}\).

**Statistical analysis**

All data analysis was performed with the Statistical Package for Social Sciences (SPSS Inc., version 26.0) and the level of statistical significance in all analyses was set to 0.05. Shappiro-Wilk test was used to test the normality of distribution of data which revealed a non parametric distribution. A Mann Whitney-U test was used to find the difference in CFU, PI and GI between the studied groups; and Chi-square test was used to detect any difference related to gender and age. The Spearman’s rank correlation analysis was used to establish the correlation among the prevalence of studied bacteria and GI and PI.

**Results**

Data from 48 subjects were collected and analyzed. The overall gender and age comparison showed a non significant difference between the two groups (Data not included). Figure 1 shows the investigated genera isolated from the retainer and control cohort. There was no significant difference between the groups; however, the total microbial count was significantly higher in fixed retainer group.

![Figure 1: The proportions of S. mutans, L. acidophilus, C. albicans, A. actinomycescomitans and F. nucleatum in relation to total microbial count.](image)

The cultivity of these pathogens showed a higher proportions of all investigated genera in the fixed retainer group compared to controls. The prevalence of the isolated periodontal pathogens doubled in retainer group; additionally, the cultivation of C. albicans raised by approximately one third in retainer group (Figure 2).
Regarding plaque (PI) and gingival (GI) indices, there were statistically significant differences between the studied groups (p<0.001). Both indices were positively correlated to each other as seen in table 1.

Table 1: Correlation between PI and GI indices between retainer and control groups (n=24).

<table>
<thead>
<tr>
<th>group</th>
<th>Min</th>
<th>Max</th>
<th>Mean ±SD</th>
<th>Mann-Whitney U</th>
<th>P-value</th>
<th>Spearman’s correlation rho value</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>PI</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C</td>
<td>0.11</td>
<td>1.11</td>
<td>0.25 ±0.25</td>
<td>0.440</td>
<td>22.5*</td>
<td>0.877*</td>
<td>0.000</td>
</tr>
<tr>
<td></td>
<td>1</td>
<td>1</td>
<td>7</td>
<td>0.014</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>R</td>
<td>0.66</td>
<td>1.44</td>
<td>0.18 ±0.18</td>
<td>0.338</td>
<td>7.5*</td>
<td></td>
<td>0.000</td>
</tr>
<tr>
<td></td>
<td>7</td>
<td>4</td>
<td>5</td>
<td>0.19</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GI</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C</td>
<td>0.01</td>
<td>1.33</td>
<td>0.19 ±0.019</td>
<td>0.965</td>
<td>7.5*</td>
<td></td>
<td>0.000</td>
</tr>
<tr>
<td></td>
<td>0.55</td>
<td>3.22</td>
<td>2</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>R</td>
<td>6</td>
<td>3</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

* Statistically significant difference (p<0.001)
C=control group, R=retainer group

The GI was positively associated with all investigated genera. Additionally, there was a positive correlation between the PI and the periodontal pathogens i.e. *F. nucleatum*, *A. actinomyces* and *L. acidophilus* (p values were 0.002, 0.026 and 0.003 respectively), in addition to *C. albicans* (p=0.009) (Table 2).
Table 2: Correlation between PI and GI with the studied microorganisms (n=48).

<table>
<thead>
<tr>
<th>Genum</th>
<th>Spearman's correlation rho value</th>
<th>p-value</th>
<th>Genum</th>
<th>Spearman's correlation rho value</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>PI</td>
<td>0.271</td>
<td>0.062</td>
<td>GI</td>
<td>0.419*</td>
<td>0.003</td>
</tr>
<tr>
<td>Strept.</td>
<td>0.419*</td>
<td>0.003</td>
<td>AA</td>
<td>0.293*</td>
<td>0.043</td>
</tr>
<tr>
<td>Candida</td>
<td>0.374*</td>
<td>0.009</td>
<td>Candida</td>
<td>0.387*</td>
<td>0.007</td>
</tr>
<tr>
<td>Lacto.</td>
<td>0.32*</td>
<td>0.026</td>
<td>Lacto.</td>
<td>0.405*</td>
<td>0.004</td>
</tr>
<tr>
<td>Fuso.</td>
<td>0.439*</td>
<td>0.002</td>
<td>Fuso.</td>
<td>0.378*</td>
<td>0.008</td>
</tr>
</tbody>
</table>

*Statistically significant (p<0.05)

Discussion

This cohort study designed to evaluate the effect of fixed orthodontic retainers on selected organisms of the oral biofilm that considered as the main contributors in oral diseases such as dental caries, periodontitis, and oral candidiasis. The sample number was 24 which yields the alpha to 0.05 and the power of the study to 0.8 assuming the effective threshold is 10000. Streptococci, lactobacilli, actinomycetes, and fusobacteria were evaluated as these genera are often associated with oral diseases like dental cavities and periodontal issues. Additionally, candida was evaluated as it is the common opportunistic pathogen of the oral cavity and have been isolated from orthodontic retainers. The results of the current study came in accordance with that reported by other studies.

The data revealed that there was an increase in all retrieved bacteria and C. albicans. It was found that the prevalence of this fungi increased in oral microbiota of orthodontic patients and may cause candida biofilm on orthodontic retainers. The results of the current study came in accordance with that reported by other studies.

The current results showed that the majority of microbial population was S. mutans and L. acidophilus in both groups. This came in agreement with previous authors who found that these pioneer bacteria compromised the majority of dental plaque and biofilm formed on the dental prosthesis. This could be related to the surface irregularities that 'built-in' the fixed orthodontic retainers which furnish a good shelter for these aciduric and acidogenic bacteria and Mummmolo et al. who found that the levels of S. mutans and Lactobacilli in saliva increased in patients with fixed orthodontic appliance. In addition to that, Türköz et al. found that the adherence of these bacteria increased after the application of thermoplastic orthodontic appliances.

However, the proportion of retrieved microorganisms in both groups was not significant. The data of the current study agreed with that of Jing et al. who reposted that the S. mutans population was not significantly changed during the first few months of orthodontic treatment with fixed appliance.

Moreover, the cultivity of the retrieved genera in the retainer group was higher. Several factors may attribute to that. It was proposed that the fixed or removable orthodontic appliances may impede the maintenance of oral hygiene by reducing
the efficacy of normal innate plaque-removal mechanisms, such as salivary flow and the physiological cleansing movement of the tongue, resulting in plaque accumulation and a potential damage to the periodontium. This is specially true for *A. actinomycetemcomitans* and *F. nucleatum*. It was claimed that these bacteria act as bridging species among the oral microbiome and heterotypic community formation during biofilm formation. In addition to their abilities to aggregate with other bacteria, resulting in evolution of periodontitis. This was clinically reflected in periodontal parameter in retainer group. Both plaque and gingival index were significantly higher and correlated with the studied genera. These outcomes came in agreement with Klingler And Jason Robert who found an increase of these bacteria in the saliva of patients with fixed orthodontic appliance. Additionally, it was proposed that patients underwent orthodontic treatment were highly motivated regarding oral hygiene maintenance especially during the relatively short treatment recall visits, however, during retention, the follow-up visits are longer and the opportunity for microbial aggregation could be higher. This agreed with several authors who reported a positive correlation between plaque index and fixed retainers.

**Conclusion**

The cultivity of the cariogenic and periodontal pathogens was higher in retainer group with a significant increase in the total microbial counts. That was reflected through the significant correlation of the studied genera with PI and GI, especially in retainer wearer group.

Conflict of Interest: Non
Funding: Self-funding

**References**


