Pathogenicity of exopolysaccharide-producing 
*Actinomyces oris* isolated from an apical abscess lesion

K. Yamane¹, T. Nambu¹, T. Yamanaka¹, K. Ishihara¹, T. Tatami¹, C. Mashimo¹, C. B. Walker², K.-P. Leung³ & H. Fukushima¹

¹Department of Bacteriology, Osaka Dental University, Osaka, Japan; ²Department of Oral Biology, College of Dentistry, University of Florida, Gainesville, FL; and ³US Army Dental and Trauma Research Detachment, Institute of Surgical Research, Fort Sam Houston, TX, USA

Abstract


**Aim** To demonstrate a capacity for producing exopolysaccharides (EPSs) and an ability to form biofilm on abiotic materials of *Actinomyces oris* strain K20.

**Methodology** The productivity of EPSs and the ability to form biofilm on microtitre plates, respectively. High-performance liquid chromatography was used to determine the chemical composition of the viscous materials. To examine the role of the viscous materials attributable to the pathogenicity of this organism, the ability of strain K20 to induce abscess formation was compared in mice to that of ATCC 27044.

**Results** The viscosity of the spent culture media of K20 was significantly higher than that of ATCC 27044. Strain K20 showed dense meshwork structures around the cells and formed biofilms on microtitre plates, whereas ATCC 27044 did not. Chemical analysis of the viscous materials revealed that they were mainly composed of neutral sugars with mannose constituting 77.5% of the polysaccharides. Strain K20 induced persistent abscesses in mice lasting at least 5 days at a concentration of $10^8$ cells mL⁻¹, whereas abscesses induced by ATCC 27044 healed and disappeared or decreased in size at day 5.

**Conclusions** Strain K20 produced EPSs, mainly consisting of mannose, and formed biofilms. This phenotype might play an important role for *A. oris* to express virulence through the progression of apical periodontitis.

**Keywords:** *Actinomyces oris*, apical abscess, biofilm, exopolysaccharide.

Received 3 February 2012; accepted 25 June 2012

Introduction

Bacteria remaining in the apical part of the root canal are known to be involved in the development of apical abscesses (Nair 2006, Siqueira & Roças 2008); however, it is still unclear how these bacteria survive repeat root canal treatments. The ability to produce exopolysaccharides (EPSs) could contribute to their survival and the development of persistent infections in the human body (Costerton *et al.* 1999). For example, alginate is one of several biofilm matrices produced by *Pseudomonas aeruginosa*, a prototype biofilm-forming bacterium, and is responsible for the persistence (Ryder *et al.* 2007) and antibiotic tolerance of this organism in the airway of patients with cystic fibrosis (Shankar *et al.* 1995). Accumulated evidence emphasizes that EPSs...
### Pathogenicity of exopolysaccharide-producing Actinomyces oris isolated from an apical abscess lesion

**Authors:** Yamane K., Nambu T., Yamanaka T., Ishihara K., Tatami T., Mashimo C., Walker C. B., Leung K. P., Fukushima H.,

**Performing Organization:** United States Army Institute of Surgical Research, JBSA Fort Sam Houston, TX

**DISTRIBUTION/AVAILABILITY STATEMENT**
Approved for public release, distribution unlimited

**Security Classification:**
- **REPORT:** unclassified
- **ABSTRACT:** unclassified
- **THIS PAGE:** unclassified

**LIMITATION OF ABSTRACT:** UU

**NUMBER OF PAGES:** 10
production of oral bacteria is associated with persistent infections in the oral cavity (Yamane et al. 2009, 2010, Yamanaka et al. 2010). Capnocytophaga ochracea found in periodontal disease has been shown to produce mannose-rich EPS that can suppress murine lymphocyte mitogen responses and activate human complement response (Bolton & Dyer 1983, 1986, Dyer & Bolton 1985). It was reported that some clinical isolates of Prevotella intermedia and P. nigrescens isolated from chronic periodontitis lesions produce EPSs in a sucrose-independent manner and form biofilms that contribute to their virulence (Fukushima et al. 1992, Yamane et al. 2005, Yamanaka et al. 2009). Clinical strains of Bacillus subtilis, Rothia mucilaginosa and Escherichia hermannii isolated from persistent periapical lesions have an ability to produce self-synthesized EPSs, suggesting that these organisms can cause a persistent biofilm infection in the apical region of root canal (Yamane et al. 2009, 2010, Yamanaka et al. 2010). Previous studies using scanning electron microscopy (SEM) revealed that a significant number of microorganisms were embedded in the extracellular polymeric substances in apical periodontitis lesions (Fukushima et al. 1990, Nair 2006, Rocha et al. 2008). These studies also suggested that EPSs-producing bacteria in the periapical lesion play an important role in the recurrence of acute apical periodontitis or in the development of abscess lesions from asymptomatic apical pathosis.

In this study, the unique characteristics of Actinomyces oris strain K20 that was isolated as a dominant bacterium in an apical abscess lesion and has been maintained in a culture collection since its initial isolation have been described. This strain spontaneously formed mucoid-type colonies despite subculturing under laboratory condition and produces viscous materials in spent culture medium. Here, the physical characteristic (viscosity) and chemical nature of the EPS produced by this organism, its ability to form biofilm and its pathogenicity as determined by abscess formation in mice, were characterized. Additionally, re-identification was performed on this clinical strain because it has been reported that strains of A. naeslundii, A. viscosus, A. johnsonii and A. oris cannot be identified by conventional phenotype testing and 16S ribosomal ribonucleic acid (rRNA) sequencing (Henssge et al. 2009).

**Materials and methods**

**Bacteria**

A clinically isolated A. oris strain K20 from an apical abscess lesion was used. Strain K20 was found to be one of the dominant bacteria in an oral abscess lesion and identified as A. oris by phylogenetic analysis. The isolation of a bacterium from an apical abscess was performed according to routine procedures as follows: The patient was asked to rinse the mouth with 0.2% chlorhexidine solution for 30 s. The apical abscess (submucosa abscess) was then isolated with sterile cotton wool rolls, and the overlying mucosa was wiped with a cotton wool bud soaked in 0.2% chlorhexidine solution. The suppuration sample was collected from the apical abscess by aspiration with a sterile 21-gauge needle. The sample was diluted with 0.1 mol L⁻¹ phosphate-buffered saline (PBS) and plated on anaerobe 5% sheep blood agar plates (BAPs) formulated by the Center for Disease Control (BBL Microbiology Systems, Cockeysville, MD, USA). The inoculated plates were incubated anaerobically with the atmosphere composed of 10% carbon dioxide (CO₂), 10% hydrogen (H₂) and 80% nitrogen (N₂) in an anaerobic chamber (ANX-3, Hirasawa, Tokyo, Japan) or aerobically at 37 °C for 24–72 h. Grown colonies were used to recover biofilm-forming bacteria. The samplings of bacteria were from patients who were referred to the Department of Endodontics at Osaka Dental University Hospital and approved by the ethics committee of Osaka Dental University (Ethics Committee Approval No. 060641). Informed consent was obtained from the patients after the nature of the procedure, possible discomfort and risks had been fully explained.

**Re-identification of strain K20 by phylogenetic analysis using atpA sequence**

Strain K20 was presumptively identified as A. viscosus by 16S rRNA gene sequencing and catalase productivity (Ellen 1976). However, a recent study suggested that partial sequences of two housekeeping genes (atpA and metG) can separate A. naeslundii genospecies 1, genospecies 2 (including A. viscosus serotype 2) and genospecies 3 (Henssge et al. 2009). Therefore, pyrosequencing of strain K20 genome was performed to obtain data necessary for re-identification of this organism. A draft genome sequence of strain K20 was determined with a combined strategy of 454-pyrosequencing and paired-end sequencing using the Roche Genome Sequencer FLX System (Roche Diagnostics, Tokyo, Japan) and Applied Biosystem 3730 DNA (deoxyribonucleic acid) Analyzer (Applied Biosystems, Tokyo, Japan). The sequence data were assembled with GS De Novo Assembler Software (version 1.1.03.24) (Roche Diagnostics) and deposited in the databases of the DNA Data Bank.
Phylogenetic analysis was performed by using the sequence of the housekeeping gene actpA (ATP synthase F1 α subunit: accession number: AB573870, DDBJ) and the data set of actpA for Actinomyces as described elsewhere (Henssge et al. 2009). The nucleotide sequences of actpA were aligned by MAFFT (version 6) with FFT-NS-I option (Katoh & Toh 2010). Then, a neighbour-joining tree was constructed using Jukes–Cantor distance matrices. Bootstrap values were determined with 500 replicates. A phylogenetic tree constructed in this study indicated that strain K20 belonged to the A. oris clade with high bootstrap values (Fig. 1). Therefore, A. oris ATCC 27044 (an ATCC type strain for A. oris) was chosen as a reference strain for comparison in this study.

Viscosity of spent culture media

The viscous material- or EPS-producing ability of strains K20 and ATCC 27044 was first evaluated by measuring the viscosity of spent culture media. Strains K20 and ATCC 27044 were grown in tryptic soy broth (TSB) for 20 h. For the viscosity measurement, an automated microviscometer (AMVN) (Anton Paar, Graz, Austria) was used in accordance with the manual provided by the manufacturer. The culture supernatant was filled into a glass capillary and maintained at 20 °C. The measurement was performed five times on each sample, and the results of four independent cultures were averaged to give the final value.

SEM observation

SEM was used to observe the cell surface structure of strains K20 and ATCC 27044. Each strain grown on a BAP aerobically for 24 h was collected on a piece of filter paper (Glass fiber GA55, Toyo Roshi, Tokyo, Japan) and fixed with 2% glutaraldehyde in 0.1 mol/L PBS. Samples were dehydrated and fixed with 1% osmium tetroxide (OsO4) in 0.1 mol/L PBS for 1 h at 4 °C. Samples were dehydrated and followed by platinum ion coating (E-1030, Hitachi, Tokyo, Japan). Specimens were examined with an SEM (S-4800, Hitachi) at an accelerating voltage of 1 kV.

Biofilm assay

The ability to form biofilm was determined as described previously (O’Toole et al. 2000). Briefly, strains K20 and ATCC 27044 were inoculated into 1 mL of Actinomyces broth (BBL Microbiology Systems) in 24-well flat-bottomed microplates (AGC Techno Glass, Chiba, Japan) at a concentration of 10^7 CFU per well. The plates were incubated statically for 2 days at 37 °C without shaking to establish biofilms. Unbound cells were removed by inversion of the microplate and tapping on an absorbent paper towel. Adherent cells were then stained with 0.1% crystal violet for 15 min. Excess stain was removed by washing with PBS. The stained wells were recorded photographically using a camera set (PC1089, Canon, Tokyo, Japan). The crystal violet was then solubilized by the addition of 95% ethanol, and absorbance at 570 nm was measured by spectrophotometer U-2000 (Hitachi). Quadruplicate wells for each strain were measured, and the experiments were performed in triplicate.

Preparation of EPSs

The EPS (viscous materials) was prepared from culture supernatants by the method described by Campbell & Pappenheimer (1966). Briefly, strains K20 and ATCC 27044 were grown at 37 °C in TSB for 24 h, respectively. Supernatants were separated by centrifuging the liquid culture at 8,000 rpm for 30 min, and sodium acetate was added to supernatants at a final concentration of 5%. The mixture was stirred for 30 min at 22 °C, and the EPS was isolated by ethanol precipitation from the reaction mixture. The ethanol-precipitated material was collected by centrifugation (12,000 rpm for 15 min, at 22 °C), resolved in 5% sodium acetate and then treated with chloroform–1-butanol (1:5, by volume). Water-soluble and chloroform–butanol layers were separated by centrifugation, equal amounts of ethanol were added to the water-soluble layer, and the ethanol-precipitated material was freeze-dried until use.

High-performance liquid chromatography (HPLC) analysis of EPSs

Neutral monosaccharides were released from purified EPS (5 mg) by hydrolysis in a sealed tube with 2 N trifluoroacetic acid (200 mL) at 100 °C for 6 h. The hydrolysate was concentrated in vacuo and dissolved in 500 mL of distilled water. The sugars were identified by HPLC (LC-9A, Shimadzu, Kyoto, Japan) with a TSKgel® sugar AXG column (15 cm × 4.6 mm) (Tosoh, Tokyo, Japan) using 0.5 mol L^-1 potassium tetraborate buffer (pH 8.7) as a carrier at a flow rate of 0.4 mL min^-1 and a column temperature of 70 °C. The amount of extracellular DNA in purified viscous
materials or EPSs (1 mg) was determined by absorbance measurement at 190–360 nm with a U-2001 spectrophotometer (Hitachi).

**Animal study**

Strains K20 and ATCC 27044 were grown in TSB for 48 h. The optical density of each bacterial culture was measured at 600 nm, and their cell concentrations were adjusted to $5.0 \times 10^8$ CFU mL$^{-1}$ according to a standard curve. A total of 500 µL of bacteria was injected subcutaneously into the inguinal region of each BALB/c mouse (male, 4 weeks, five mice per strain). The development of abscess lesions was recorded photographically using a camera (Nikon FIII, Nikon, Tokyo, Japan) set at a fixed magnification for

---

**Figure 1** Phylogenetic tree based on *atpA* region for *Actinomyces* species. A data set provided by Henssge et al. (2009) was used (accession numbers are indicated at the tree). Four clusters, each cluster containing a representative type strain (asterisk), were marked. The scale bar represents a difference of 0.002 substitution per site.
five consecutive days. Sizes of abscess areas were
determined with Photoshop (version 7.0, Adobe, San
Jose, USA) and expressed in number of pixels. Statisti-
cal analysis was performed with Student’s t-test. This
examination was repeated three times. The animal
protocol used in this study was approved by the Ani-
mal Research Committee of Osaka Dental University
(Confirmation numbers 05-21001 and 06-05003).

Results

Re-identification of strain K20

The phylogenetic tree constructed with the data set of
atpA for Actinomyces spp. (Henssge et al. 2009) and
the sequence of atpA of strain K20 obtained by the
genome pyrosequencing of strain K20 indicated that
it belonged to the A. oris clade as described previously
(Fig. 1).

Viscosity of spent culture medium

The viscosity of the spent culture medium obtained
from strain K20 cultures was significantly higher
than that of the control TSB medium or ATCC 27044
cultures (Fig. 2).

Chemical compositions of viscous materials (EPSs)

Chemical analyses showed that the purified viscous
materials obtained from strain K20 were composed of
neutral sugars with mannose constituting 77.5% of
sugars recovered from the culture supernatant
(Table 1), suggesting that these materials are EPSs.
This fraction did not contain extracellular DNA at a
substantial level (data not shown). Viscous materials
were not isolated from the spent culture of ATCC
27044 at a substantial level.

SEM observation of cell surface structures

SEM revealed that strain K20, but not ATCC 27044,
had dense meshwork-like structures around the cells
(Fig. 3).

Biofilm formation in microtitre plates

The ability to form biofilm was investigated for
strains K20 and ATCC 27044 using a crystal violet
microtitre plate assay. Figures 4 (a,b) show that
strain K20 was able to form biofilm on polystyrene
plates, whereas ATCC 27044 showed poor biofilm
formation. Quantitative analysis measuring the optical
density of destained biofilms at 570 nm revealed
that the ability of strain K20 to form biofilm was
significantly greater than that of ATCC 27044
(Fig. 4c).

Animal study

Two days after the injection of suspended bacterial
cells at a concentration of $10^8$ CFU mL$^{-1}$, strains K20
and ATCC 27044 induced distinct abscesses in mice
(Fig. 5a,b). The abscesses induced by strain K20 lasted
at least 5 days in all mice, whereas the abscess lesions

![Figure 2](viscosities.png)

**Figure 2** Viscosities of spent culture media obtained from
strains K20 (a clinical isolate from an apical abscess lesion)
and ATCC 27044 (a reference strain for Actinomyces oris).
Medium: control tryptic soy broth without bacterial inocula-
tion. Statistical analysis was performed with Student’s t-test.
*P < 0.05.

![Table 1](table1.png)

**Table 1** Sugar composition of exopolysaccharides isolated
from strain K20

<table>
<thead>
<tr>
<th>Neutral sugars</th>
<th>Contents (µg)</th>
<th>Content ratio (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mannose</td>
<td>165.31</td>
<td>77.5</td>
</tr>
<tr>
<td>Glucose</td>
<td>21.71</td>
<td>10.2</td>
</tr>
<tr>
<td>Ribose</td>
<td>13.28</td>
<td>6.2</td>
</tr>
<tr>
<td>Galactose</td>
<td>6.15</td>
<td>2.9</td>
</tr>
<tr>
<td>Fructose</td>
<td>3.32</td>
<td>1.6</td>
</tr>
<tr>
<td>Xylose</td>
<td>2.10</td>
<td>1.0</td>
</tr>
<tr>
<td>Arabinose</td>
<td>1.14</td>
<td>0.5</td>
</tr>
<tr>
<td>Rhamnose</td>
<td>&lt;1.00</td>
<td>&lt;0.5</td>
</tr>
<tr>
<td>Total</td>
<td>213.28</td>
<td>100</td>
</tr>
</tbody>
</table>
induced by ATCC 27044 healed and disappeared in three of five mice or decreased in size in the other two mice at day 5 (Fig. 5c,d). The average abscess size induced by strain K20 at day 5 was significantly larger than that of ATCC 27044 (Fig. 5e).

Discussion

*Actinomyces* spp. are predominant members of human oral commensal microbiota and are known as initial colonizers on tooth surfaces (Nyvad & Kilian 1987, Kolenbrander et al. 2005, Diaz et al. 2006). Oral *Actinomyces* spp. are branching, non-spore-forming rods that belong to the high G + C division of Gram-positive bacteria. On the basis of 16S rRNA gene sequencing and DNA–DNA relatedness, six oral *Actinomyces* spp. (*A. georgae, A. odontolyticus, A. israelii, A. gerencseriae, A. meyeri and A. naeslundii* genospecies I, II and III) have been recognized (Tanner et al. 1994). The identification of *A. naeslundii* strains, including *A. viscosus* serotype 2 strains, had been problematic because neither 16S rRNA gene sequencing nor biochemical tests can differentiate these three genospecies. Recently, Henssge et al. (2009) demonstrated that the partial sequences of two housekeeping genes (*atpA* and *metG*) can separate three *A. naeslundii* genospecies. They proposed the name *A. oris* for *A. naeslundii* genospecies II and the name *A. johnsonii* for *A. naeslundii* genospecies III (genospecies WVA 963). In this classification, *A. naeslundii* genospecies I remains as *A. naeslundii*. According to this classification, strain K20 was re-identified as *A. oris*.

In persistent apical periodontitis lesions, actinomycosis has been known as an important cause (Nair & Schroeder 1984), and several oral *Actinomyces* spp. such as *A. israelii* (Nair 2006), *A. radicidentis* (Collins et al. 2000, Kallas et al. 2001) and *A. naeslundii* (Happonen et al. 1985) are frequently isolated from human apical periodontitis lesions. Figdor et al.
(1992) demonstrated that the pathogenicity of A. israelii isolated from a case of failed endodontic therapy (Sundqvist & Reuterving 1980) was because of its ability to establish characteristic cohesive colonies consisting of cells enmeshed in an extracellular matrix. The organisms existing in such colonies can collectively evade destruction and elimination by host phagocytic cells. A recent study demonstrated the capacity of A. israelii to produce EPSs and to form biofilm in vitro on copper surfaces of intrauterine contraceptive devices immersed in a simulated uterine fluid under anaerobic conditions (Carrillo et al. 2010). Actinomyces radicidentis is known to form biofilms, in which bacteria seem to be embedded in the extracellular matrix and exhibit antiphagocytic activity against rat neutrophils (Nair et al. 2008). Actinomyces meyeri, the cause of various actinomycosis or abscess lesions (Apotheloz & Regamey 1996, Liaudet et al. 1996), was also found in periapical lesions refractory to endodontic therapy (Sunde et al. 2002).

The present study found that A. oris strain K20 has a capacity to produce EPSs in liquid culture condition as viscous materials and meshwork structures around the cells on agar plates. These activities are similar to those of slime-producing bacteria with biofilm-forming ability such as P. aeruginosa (Kobayashi 1995, Yasuda et al. 1999), Salmonella enterica Serovar Typhimurium (Anriany et al. 2001), Staphylococcus epidermidis (Vuong et al. 2004) and Vibrio cholerae (Wai et al. 1998). Considering the fact that ATCC 27044, a type strain for A. oris, which lacks an ability to produce EPS at a substantial level and does not form meshwork on its cell surface, failed to form apparent biofilm on microtitre plates and could not induce persistent abscess lesions in mice, the self-synthesized polysaccharide seems to be important to biofilm formation on abiotic materials as well as to the pathogenicity of A. oris strain K20. It is important to note that other constituents such as extracellular nucleic acids (Wu & Xi 2009) or secreted proteins (Latasa et al. 2006) found in other organisms are also used as part of the biofilm matrix that are essential in their biofilm development. Similar tubule-like structures are formed by bacterial nanotubes (Dubey & Ben-Yehuda 2011) or amyloids (Dueholm et al. 2010). Several Actinomyces spp. are armed with type 1 and type 2 fimbriae for being able to adhere to a host surface, and fimbriated strains including A. naeslundii genospecies 1 (A. naeslundii), genospecies 2 (A. oris) and A. viscosus exert higher cell surface interactive force than those devoid of fimbriae (Tang et al. 2004). Therefore, how the EPSs produced by A. oris interacts with other cell surface structures and how many structural elements contribute to the establishment of persistent infection caused by A. oris are still to be elucidated. A study in the early 1970s clearly showed that the addition of slime from P. aeruginosa cultures to E. coli or Staphylococcus aureus dramatically inhibited phagocytosis by neutrophils (Schwarzmann & Boring 1971). As described previously, it has also reported that a clinical isolate of P. nigrescens, which has the

Figure 5 Abscess induction in mice by strains K20 and ATCC 27044. Two days after the injection of suspended bacterial cells at a concentration of $10^8$ CFU mL$^{-1}$, strains K20 and ATCC 27044 induced distinct abscesses in mice (a,b). Persistent abscesses (white arrows) were observed at day 5 (c). In contrast, the abscess lesions induced by ATCC 27044 decreased in size or cured and disappeared (d). The average abscess size induced by strain K20 was significantly larger than that of ATCC 27044 (e). Statistical analysis was performed with Student’s t-test. *P < 0.05.
ability to produce a copious amount of EPS, is far more potent in causing abscess formation in mice and far more resistant to the phagocytic activity of human polymorphonuclear leukocytes than the chemically induced mutant that lacked the EPS (Yamane et al. 2005). When the purified EPS from the biofilm-forming strain of *P. nigrescens* was added to the cultures of a biofilm-non-forming mutant, these cells exhibit the presence of these meshwork structures on their surfaces and demonstrate the ability to induce abscess formation in mice (Yamane et al. 2005). As *A. israelii* packed in alginate gel induced actinomycotic lesions in mice efficiently, but a bacterial suspension without alginate or gel particles without bacteria did not induce lesions (Sumita et al. 1998), it seems that polysaccharide coating of bacterial mass might determine the persistence of *Actinomyces* spp. in the host as described in many pathogens (Costerton et al. 1987).

Although the bacteria in biofilms are considered to be dormant in general, it is conceivable that *Actinomyces* spp. have another strategy to survive when the nutrients are scarce such as in apical lesions. Paddock et al. (2005) showed that *A. oris* (*A. naeslundii* genospecies 2) together with *Streptococcus oralis*, *Streptococcus mitis* and *Streptococcus intermedius* form the bulk of the microbiota that are present in dentine samples beneath the restoration for 5 months. *Actinomyces oris* in the 5-month-old dentine samples was positive for sialidase, β-galactosidase and N-acetyl-β-glucosaminidase activities (Paddock et al. 2005). The complete function of sialidase of this organism is not fully understood. Recent studies suggested that this enzyme could provide nutrients to the cells by liberating sialic acid from glycans (Paddock et al. 2005, Do et al. 2008) and mediate the adhesion of the bacteria to host tissues by exposing the galactose residues of glycoprotein glycans that are present on mucosal surfaces (Ruhl et al. 2000, Do et al. 2008). The importance of *A. oris* type 2 fimbriae, which possess the lectin-like activity, in biofilm formation and co-aggregation has been vigorously studied (Yeung 1999, Mishra et al. 2010).

The factors controlling the establishment of bacterial biofilm in the periapical lesion that can lead to the recurrence of an acute abscess lesion in the site appear to be complex. Further thorough investigations are needed to elucidate the nature of biofilm matrices, extracellular enzymatic activities and the interaction between the host and *A. oris* in the apical abscess lesion.

**Conclusions**

Taken together, the results in this study suggest that the EPS productivity of *A. oris* strain K20 affects its pathogenicity and that this organism could be involved in the development of apical abscess lesions.

**Acknowledgements**

We would like to thank Mr. Hideaki Hori for his excellent assistance with the electron microscopy. This study was performed at the Institute of Dental Research, Osaka Dental University, and supported in part by the Osaka Dental University Joint Research Funds (B08-01) and JSPS KAKENHI Grant Number 23592724, 23792118, 24791975 and 80368159.

**References**


FEMS Microbiology Letters 288, 156–62.


Henssge U, Do T, Radford DR, Gilbert SC, Clark D, Beighton D (2009) Emended description of Actinomyces naeslundii and descriptions of Actinomyces oris sp. nov. and Actinomyces johnsonii sp. nov., previously identified as Actinomyces naeslundii genospecies 1, 2 and WVA 963. International Journal of Systematic and Evolutionary Microbiology 59, 509–16.


Oral Surgery, Oral Medicine, Oral Pathology, Oral Radiology, and Endodontics 92, 208–14.


