Oral Microbiology and Immunology

# Role of gene *E2f1* in susceptibility to bacterial adherence of oral streptococci to tooth surfaces in mice

Matsumoto N, Salam MA, Watanabe H, Amagasa T, Senpuku H. Role of gene E2f1 in susceptibility to bacterial adherence of oral streptococci to tooth surfaces in mice. Oral Microbiol Immunol 2004: 19: 270–276. © Blackwell Munksgaard, 2004.

Dental plaque is composed of a biofilm community of microorganisms on teeth that coats the oral cavity, including attaching to the teeth, and provides a protective reservoir for oral microbial pathogens, which are the primary cause of persistent and chronic infectious diseases. Oral streptococci are pioneering organisms that play an important role in biofilm formation on tooth surfaces as well as being primary causative agents of dental caries. The purpose of this study was to clarify the role of the E2f1 gene in susceptibility to dry mouth and bacterial adherence of oral streptococci to tooth surfaces in animal model experiments. A mutation of the *E2f1* gene in mice is known to cause enhanced T-lymphocyte proliferation, leading to testicular atrophy, splenomegaly, salivary gland dysplasia, and other systemic and organ-specific autoimmunity. We found a decreased volume of saliva production and protein production rate, along with increased amylase activity, IgA concentration, and mucin 1 concentration in E2F-1<sup>-/-</sup> mice as compared with the control C57BL/6 mice. Further, we quantified the recolonization of oral streptococci in E2F-1<sup>-/-</sup> mice and found that a higher number of some oral streptococci were colonized on the teeth of these mice. In particular, following oral ingestion of 1% sucrose in water, the colonization of Streptococcus mutans increased in comparison with other streptococci. Our results suggest that the *E2f1* gene may affect susceptibility for oral biofilm formation by streptococci in humans with dry mouth.

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Key words: biofilm; dry mouth; E2F-1 knockout mouse; saliva; *Streptococcus mutans* 

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Dental plaque is a complex biofilm community of microorganisms that attaches to the surfaces of teeth, consisting of more than 500 bacterial species in a mature state and providing a reservoir for oral microbial pathogens (8, 19, 26). The initial colonizers of freshly cleaned enamel surfaces are predominantly streptococci (24), which account for approximately 20% of the total number of salivary bacteria (18). Oral streptococci are pioneering organisms that play an important role in biofilm formation on tooth surfaces as well as being primary causative agents of dental caries (8, 13, 17). The ability of oral streptococci to bind to the salivary pellicle on the tooth surface protein is of considerable etiologic significance (6, 23, 30), and *Streptococcus mutans* and *Streptococcus sanguis* are known to be primarily involved when bacterial flora forms on tooth surfaces. In addition, *S. sanguis* and *Streptococcus mitis* are early colonizers of the salivary pellicle, while *S. mutans* colonizes later on. The ability of each to bind to salivary proteins and glycoproteins is strong and important in biofilm development (15, 39).

The mechanical forces of salivary flow and tongue movement tend to dislodge and expel bacteria from tooth surfaces and the oral cavity (3, 4). It has been proposed that streptococci bind the salivary component receptors via adhesins present on the cell wall. In addition, there is direct evidence that streptococcal cell wall components mediate adherence to various salivary receptors (5, 23, 29, 31, 33). Their importance in controlling microbial colonization in the oral cavity has also been well demonstrated in individuals with diabetes mellitus, Sjögren's syndrome, and dry mouth, who suffer from a rapid overgrowth of biofilm and rampant caries, making them highly susceptible to oral infections (1, 2).

Members of the E2F transcription factor family (E2f1-E2f5) are important regulators of cell proliferation, differentiation, and apoptosis. The best characterized member of the E2F family is *E2f1* (9, 11, 36), which controls the initiation of DNA synthesis and subsequent transition of cells from the G0/G1 to S phase of the cell cycle (14, 22). Several recent studies have demonstrated that a mutation of the E2f1 gene in mice causes enhanced T-lymphocyte proliferation, leading to testicular atrophy, splenomegaly, salivary gland dysplasia, and other types of systemic and organ-specific autoimmunity (7, 16, 20, 28, 42, 44). We propose E2F-1 to be a host defense regulator acting against oral pathogens, and attempted to clarify the role of the *E2f1* gene in susceptibility to dry mouth and the initial adhesion of oral streptococci on tooth surfaces in animal model experiments with E2F-1<sup>-/-</sup> and wild type mice.

#### Material and methods Bacterial strains and culture conditions

The streptococci strains used in this study were *S. mutans* MT8148, *S. sanguis* ATCC 10556, *Streptococcus salivarius* ATCC 9759, *S. mitis* ATCC 6249, and *Streptococcus sobrinus* 6715. All bacteria were grown in an atmosphere of  $H_2$  and  $CO_2$  (GasPack, Becton/Dickinson, Sparks, MD) in Brain Heart Infusion broth (BHI, Difco Laboratory, Detroit, MI) at 37°C.

#### Animals

E2F-1<sup>-/-</sup> and NOD/LtJ mice were purchased from The Jackson Laboratory (Bar Harbor, ME), B10.D2 and C57BL/6 mice came from Japan SLC (Shizuoka, Japan), and 129SV mice were provided by Atsuo Ogura, Department of Veterinary Science, National Institute of Infectious Diseases. NOD mice, which naturally develop Sjögren's syndrome and dry mouth, were used to compare the susceptibility for adherence of oral streptococci with E2F-1<sup>-/-</sup> mice. C57BL/6 and 129SV mice were parental lines to produce E2F-1-/ mice (7) and were used as control mice in saliva and bacterial inoculation experiments. B10.D2 mice having different background genes from E2F-1<sup>-/-</sup> mice and which did not present salivary gland abnormalities were also used as control mice. All were maintained in accordance with the guidelines of the National Institute of Infectious Diseases. Clinical onset of diabetes in NOD mice was determined by the presence of glucose in urine and blood. Urine was tested weekly using Uristix reagent strips (Bayer Medical Ltd, Newbury, UK) and diabetes was confirmed to be positive by blood glucose measurements. All mice were provided with sterile food and water, and were used in the experiments at 16–24 weeks of age.

# Bacterial sampling and colony-forming unit (CFU) counting

Bacterial inoculation, sampling and CFU counting were performed using procedures and conditions described previously (32). All oral streptococci were cultured in BHI broth overnight and then washed twice with sterile phosphate-buffered saline (PBS). Chlorhexidine (0.2%) soaked sterile cotton swabs were used for disinfecting the oral cavities of the mice. including the mandibular incisor teeth. which were immediately washed with sterile PBS. Oral streptococci were introduced to the oral cavities of all females at 4 months of age at a final concentration of  $7.5 \times 10^9$  in 250 µl of PBS for 2.5 min. As an additional experiment to reproduce the early adherence of streptococci in conditions closer to a natural state, mice were given drinking water containing 1% sucrose, which is similar to the concentration in juice, 1 day before the streptococci inoculation (34). Following inoculation, samples were collected from the labial surfaces of the mandibular incisor teeth with a sterile cotton ball and then dipped in 2 ml of PBS. The samples in PBS were sonicated by ultrasonic dispersion (power output, 60 W) for 10 s, and then poured onto Mitis-Salivarius agar plates (MS) or MS containing 0.02 M bacitracin (MSB) using an EDDY JET spiral system (Gunze Sangyo, Inc., Tokyo). The number of CFU was determined after 48 h of anaerobic incubation at 37°C.

#### Saliva collection and flow volume

Under anesthesia, the mice were injected with a cocktail of isoproterenol (0.20 mg/ 100 g body weight) and pilocarpine (0.05 mg/100 g body weight) (Sigma Chemical, St. Louis, MO) in PBS (0.1 ml) as a secretagogue. Following the intraperitoneal injection, whole saliva was collected from each mouth by a micropipette for 15 min and stored at  $-20^{\circ}$ C in preweighed tubes. Volume was calculated

based on the weight change of the tubes using a Mettler analytical balance.

### α-Amylase activity and protein concentration

Whole saliva samples were examined for amylase activity to hydrolyze starch using an amylase kit (Sigma Chemical). Aliquots of 20 µl of saliva diluted 1000-fold in PBS were added to 1 ml of an amylase reagent, which contained 4, 6-ethylidene (g7)-pnitrophenyl (G1)-a, and D-maltoheptaside (ET-G7PNP), and mixed immediately by inversion then incubated at 37°C for 2 min. Absorbance was measured at 405 nm. Protein concentrations in saliva were also measured using a Bio-Rad Protein Assay (Bio-Rad Laboratory, Hercules, CA) based on the method of Bradford, an accurate procedure for determining the concentration of solubilized protein that involves the addition of an acidic dve to the protein solution and subsequent measurement at 595 nm.

# SDS, and native PAGE, and Western blot analysis

Prior to electrophoretic analysis, the saliva samples were diluted with an equal volume of SDS-PAGE sample buffer [0.06 M Tris-HCl (Amersham Pharmacia Biotech, Buckinghamshire, UK), pH 6.8, 20% glycerol (Wako Pure Chemical Industries Ltd, Osaka, Japan), 1% (wt/vol) SDS (Wako), 1% 2-mercaptoethanol (2-ME, Sigma Chemical), and 0.0012% bromphenol blue (Wako)] and native PAGE sample buffer without SDS and 2-ME. The SDS-PAGE samples were heated at 100°C for 3 min just prior to applying to the gel. The samples were then subjected to a 10% polyacrylamide gel (PAGEL-compact, ATTO Corp., Tokyo, Japan) in the presence of 0.025 M of Tris-HCl, 192 mM of glycine (Wako), 0.1% (wt/vol) SDS, and 1% 2-ME for SDS-PAGE, as well as to 5-20% gradient polyacrylamide gels (PAGEL-compact, ATTO Corp.) without SDS and 2-ME for native PAGE. Electrophoretic separation of the proteins was carried out in SDS and native PAGE for 60 min at 100 mA. For molecular weight markers, low-molecular (phosphorylase b 94.0 kDa, bovine serum albumin 66.0 kDa, ovalbumin 45.0 kDa, carbonic anhydrase 30.0 kDa, trypsin inhibitor 20.1 kDa, and *α*-lactalbumin 14.4 kDa) and high-molecular (thyroglobulin 669 kDa, ferritin 440 kDa, catalase 232 kDa, lactate dehydrogenase 140 kDa, and albumin 66 kDa) weight markers were

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Table 1. Comparison of physiological and biological changes in saliva between E2F-1<sup>-/-</sup>, C57BL/6, NOD/LtJ and 129SV mice

| Mice ( <i>n</i> )    | Body weight (g)                  | Total saliva volume (µl)                              | Saliva volume<br>(µl/100 g BW)                     | Saliva amylase<br>activity (IU/I)                    | Saliva protein<br>concentration (µg/µl) | Saliva protein/min<br>(µg/min)      |
|----------------------|----------------------------------|---|--|--|---|-------------------------------------|
| $F2E 1^{-/-} (10)$   | $22.7 \pm 1.6^{*,**,***}$        | $40.4 \pm 17.5 * * * * * * * * * * * * * * * * * * *$ | $7100 \pm 752************************************$ | $1027.4 \pm 280.8*$                                  | $75 \pm 0.8$                            | $\frac{10}{220 \pm 0.5* **}$        |
| C57BL/6 (10)         | $22.7 \pm 1.0$                   | $49.4 \pm 17.3$                                       | $218.2 \pm 75.3$<br>$438.2 \pm 84.0$               | $1037.4 \pm 380.8^{\circ}$<br>$435.4 \pm 178.3^{**}$ | $7.3 \pm 0.8$<br>64 + 18                | $23.0 \pm 9.5$<br>$45.0 \pm 13.5**$ |
| Nondiabetic NOD (10) | $25.2 \pm 2.0$<br>$25.9 \pm 1.9$ | $138.0 \pm 39.2^{***}$                                | $538.8 \pm 160.5$                                  | $1198.5 \pm 350.0^{*}$                               | $7.7 \pm 1.3$                           | $69.5 \pm 5.4^*$                    |
| Diabetic NOD (6)     | $25.0 \pm 1.5$                   | $94.5 \pm 65.7 **$                                    | $377.7 \pm 265.8$                                  |  |   |                                     |
| 129SV (4)            | $19.9 \pm 0.7^{***}$             | $176.0 \pm 20.0^{***}$                                | $883.5\pm89.9^{***}$                               | _  | _                                       | _                                   |

All values are mean  $\pm$  SD.

\*P < 0.05, vs C57BL/6 and 129SV. \*\* P < 0.05, vs nondiabetic NOD. \*\*\* P <0.05, vs diabetic NOD.

-: Not detected. BW, body weight.

used (Amersham Pharmacia Biotech). When electrophoresis was completed, proteins in the gels were stained with 0.1% (wt/vol) Coomassie Brilliant Blue. For Western blot analysis, proteins in the gels were transferred to nitrocellulose membranes (ATTO Corp.), which were washed in 25 mM Tris-HCl (pH 7.4), 0.5 M NaCl, and 0.1% (v/v) Tween-20 (TTBS) buffer, and specific binding was blocked by



*Fig. 1.* Detection of mucin 1 in C57BL/6, E2F-1<sup>-/-</sup>, and nondiabetic NOD mice. (A) Salivary components, as detected by Coomassie Blue staining in native PAGE. Whole saliva samples diluted by 1/2, 1/4, and 1/8 were subjected to 5–20% gradient polyacrylamide gels. M: High-molecular markers. (B) Western blot detection of mucin 1. Positive smear bands were detected in some lanes. Data are representative of three independent experiments, with similar results obtained in each.



*Fig.* 2. Detection of IgA in C57BL/6, E2F-1<sup>-/-</sup>, and nondiabetic NOD mice. (A) Salivary components, as detected by Coomassie Blue staining in SDS-PAGE. Whole saliva samples diluted by 1/2, 1/4, and 1/8 were subjected to 10% polyacrylamide gels. M: Low-molecular markers. (B) Western blot detection of IgA. Molecular weight markers are indicated on the left. Data are representative of three independent experiments, with similar results obtained in each.

incubating the blots for 1 h in 0.3% skim milk/TTBS. The transferred proteins were incubated for 1 h in 1/1000 rabbit polyclonal antibody mouse IgA (Nordic Immunological Laboratories, Tilburg, Netherlands) or goat polyclonal antibody mouse mucin 1 (Santa Cruz biotechnology, Santa Cruz, CA), and then washed in TTBS three times for 5 min. The membranes were then incubated for 1 h in 1/ 500 antirabbit and goat IgG antibodyhorseradish peroxidase conjugate (Amersham Pharmacia Biotech), and washed again in TTBS three times for 5 min. Specific bindings were finally detected using ECL Plus, a Western blot detection system (Amersham Pharmacia Biotech), with a chemiluminescent reaction and a Lumishoot (ATTO Corp.) to produce the image.

#### Statistical analysis

Comparative analyses were performed by ANOVA. *P*-values less than 0.01 or 0.05 were considered statistically significant for two-tailed comparisons. All statistical analyses were performed using StatView for the Macintosh operating system.

#### Results

# Measurements of salivary flow and components

Table 1 summarizes the relative differences in body weight, saliva volume, amylase activity, protein concentration, protein production/minute and in E2F-1<sup>-/-</sup>, nondiabetic and diabetic NOD/ LtJ, C57BL/6, and 129SV mice. Total volume and volume per 100 g of body weight of secreted saliva were significantly smaller in  $E2F-1^{-/-}$  than in the others. However, salivary amylase activity in E2f1-deficient mice was similar to that in nondiabetic NOD mice and higher than in C57BL/6 mice. There were no significant differences in C57BL/6, E2F-1<sup>-/-</sup>, and nondiabetic NOD mice in salivary protein concentration. However, salivary protein production/minute in E2f1-deficient mice  $(23.0\pm9.5)$  was lower than that in non-diabetic NOD (69.5  $\pm$  5.4), and C57BL/6 (45.0  $\pm$  13.5) mice.

Strong mucin 1 smear expressions were observed in 1/2 dilution of whole saliva samples from all mice by Western blot analysis (Fig. 1). However, the expressions were stronger in 1/4 and 1/8 dilutions of whole saliva samples from E2F-1<sup>-/-</sup> and nondiabetic NOD mice than in C57BL/6 mice. Further, distinct IgA expressions were observed in 1/2, 1/4, and 1/8 dilutions of whole saliva samples from  $E2F-1^{-/-}$  mice, and in 1/2 and 1/4 dilutions from NOD mice, but not in those from C57BL/6 mice (Fig. 2). Thus, amylase, mucin 1, and IgA were found in higher concentrations in the saliva of E2F-17 and NOD mice than in the control C57BL/ 6 mice.

# Colonization of oral streptococci on tooth surfaces

*S. mutans* adhesion results revealed that bacteria were colonized on the tooth surfaces of  $E2F-1^{-/-}$  mice in higher num-

bers than on those in the control C57BL/6 mice at various time points (Fig. 3A). After extending the adhesion phase from 0 to 30 min, there was a decrease in biofilm growth; however, a large amount of bacteria remained on the tooth surfaces in E2F-1<sup>-/-</sup> mice. A significant number of bacteria (1471.5  $\pm$  1177.6) was colonized on the tooth surfaces of  $E2F-1^{-/-}$  mice at 90 min after inoculation, as compared to the control B10.D2 mice  $(44.0 \pm 72.0)$ (Fig. 3A), and the colonization of S. mutans in E2F-1<sup>-/-</sup> mice was approximately 10-fold higher than that in nondiabetic NOD mice (Fig. 3B). S. mutans, S. sanguis, S. sobrinus, S. salivarius, and S. mitis were inoculated into the oral cavities of E2F-1<sup>-/-</sup> and C57BL/6 mice, after which we measured the amount of bacterial colonization on the tooth surfaces. The CFU/ml of S. mitis  $(1928.7 \pm 1553.9)$  had a higher association than the others (less than 1200) (Fig. 4A) in E2F-1<sup>-/-</sup> mice, while that of each streptococci was less than 200 in C57BL/6 mice (data not shown). Furthermore, the CFU/ml of S. mutans



*Fig.* 3. Adhesion of *S. mutans* to tooth surfaces of experimental mice. (A) Asterisks denote significant differences (vs. C57BL/6, \*P < 0.01). Data are representative of three independent experiments, with similar results obtained in each. (B) Adhesion of *S. mutans* in nondiabetic NOD, E2F-1<sup>-/-</sup>, and B10.D2 female mice, 4 months of age, 90 min after inoculation. CFU were determined in MSB after 48 h of anaerobic incubation at 37°C. One dot was expressed as the number of *S. mutans* CFU per mouse. Seven NOD, five E2F-1<sup>-/-</sup>, and seven B10.D2 mice were used for the *S. mutans* adherence assay. Data are representative of three independent experiments, with similar results obtained in each.

 $(1478.1 \pm 770.9)$  was found to be higher than that of *S. sanguis*, *S. sobrinus*, *S. salivarius*, and *S. mitis* in the E2F-1<sup>-/-</sup> mice that ingested 1% sucrose in water in the 24 h prior to inoculation (Fig. 4B).

#### Discussion

Recent studies have found that E2F-1 functions primarily as a suppressor of cell proliferation (7, 20). Along with those findings, others have observed that E2F-1 DNA binding sites in the promoters of several all-cycle genes function primarily to suppress transcription on these genes during the G0 and G1 phases (21, 40, 41), presumably by binding to the E2F-1-Rb complex. Further, some studies have suggested that suppression of transcription of the E2F-1-Rb complex suppresses entry into the S phase (7, 20). Thus, E2f1deficient cells may be unable to tether Rb to E2F-1 DNA binding sites during the G0 or G1 phase, which may lead to an aberrant expression of particular cell cvcle regulatory genes and inappropriate cell proliferation (20). It has been speculated that a direct role of E2F-1 as a suppressor of cell cycle progression may explain the phenotypes that some investigators have observed in *E2f1*-deficient mice, such as testicular atrophy, hyposalivation or exocrine gland dysplasia, increased mortality, hair loss, and sporadic skin tumor formation (7, 16, 20, 42).

In the present study, we examined the role of *E2f1* in susceptibility to adherence of oral streptococci in model animals with dry mouth and compared the results with NOD mice, who naturally develop Sjögren's syndrome and dry mouth. The prominent features of these diseases are manifested in the E2f1-deficient mice by dry mouth and hyposalivation (4, 35, 37). Further, we found a decreased production in the volume of saliva, along with increased amylase activity and IgA and mucin 1 concentrations as compared with wild type mice (C57BL/6). However, distinct differences in IgA and mucin 1 concentrations between E2F-1<sup>-/-</sup> and NOD mice were not observed. In addition, salivary protein production/minute in E2f1-deficient mice was lower than in nondiabetic NOD and C57BL/6 mice, though the differences in salivary protein concentration were not significant. Collectively, decreased levels of saliva were seen in both E2F-1<sup>-/-</sup> and NOD mice; however, lower levels and slower production of both saliva liquid and protein were found in  $E2F-1^{-/-}$  mice than in the NOD mice. In previous histologic experiments, destruc-



*Fig.* 4. (A) Adhesion of *S. mutans, S. sanguis, S. sobrinus, S. salivarius, and S. mitis* in E2F-1<sup>-/-</sup> female mice, 4 months of age, 90 min after inoculation. (B) Adhesion of the same bacteria after oral ingestion of 1% sucrose in water. CFU were determined in MS after 48 h of anaerobic incubation at 37°C. Data were obtained from three independent experiments and are expressed as the mean  $\pm$  SD. \**P* < 0.05.

tion of salivary gland tissues by infiltration of lymphocytes in NOD mice was noted in one report (27), whereas in another, salivary gland section showed no infiltration of lymphocytes and large or doubled nuclei in  $E2F-1^{-/-}$  mice (42). It is therefore suggested that the  $E2F-1^{-/-}$  mouse has

different phenotypes for decreasing saliva and dry mouth than the NOD mouse.

It has been reported that salivary components play important roles in controlling microbial colonization in the oral cavity of individuals with Sjögren's syndrome or dry mouth (37), and dry mouth may lead to qualitative and quantitative changes in the protective salivary films or pellicles that coat hard and soft tissues (3, 4). This loss of protective activities in the oral cavity could result in an increase in dental caries and gingivitis. Streptococcal adhesion is often the result of specific interactions between the carbohydrate portions of receptor glycoproteins and protein complexes, which are known as adhesins. Further, alterations in the mucosal pellicle may make oral soft tissues more susceptible to colonization by opportunistic microflora (17-19). The S. mutans adhesion results seen in the present experiments demonstrated that a significant number of bacteria were colonized on the tooth surfaces of E2F-1<sup>-/-</sup> mice, as compared to C57BL/6 and B10.D2 mice (Fig. 3A, B). In addition, in E2F-1<sup>-/-</sup> mice a larger number of S. mutans organisms adhered to tooth surfaces than in the NOD mice (Fig. 3B). NOD mice are not suitable for studies of long-term infection by oral bacteria as the average lifespan is short compared with other mice. Therefore, E2F-1<sup>-/-</sup> mice have benefits for in vivo experiments regarding oral streptococcus adhesion to tooth surfaces as compared to wild type and NOD mice. Our present results with those mice showed that the E2F-1 gene is involved in susceptibility not only to dry mouth but also to bacterial adherence of oral streptococci to tooth surfaces.

A problem facing in vivo oral biofilm research is the lack of a naturalistic, reproducible, longitudinal monitoring system that permits the assessment of dry mouth and oral bacterial infection in the same animals throughout the duration of the study. Studies of S. mutans infection in mouse oral cavities have been performed by feeding the animals with powdered diet 301 and diet 2000, which contain unnatural amounts of sucrose (1% and 56%, respectively) (10, 38, 43). In other studies of infection, mice were provided with either a 5% sucrose or sucrose-free diet. in which the 56% of the sucrose in diet 2000 was replaced with wheat flour, and the presence of sucrose facilitated the colonization of S. mutans, even when smaller S. mutans inocula were used (12, 25). However, continuous ingestion of food containing such excess amounts of sucrose is unusual (34). Since S. mutans adherence to tooth surfaces may depend on a balance between physical adherence and synthesis of insoluble glucan in a natural environment, those previous methods may be inappropriate for investigation of the natural biofilm formation by streptococci, including S. mutans and S. sanguis.

In the present study, the importance of salivary flow for controlling the initial adhesion of oral streptococci in the oral cavity was demonstrated in mouse models of Sjögren's syndrome and dry mouth. Recently, we established recombinant NOD.B10.D2 mice by replacing major histocompatibility complex (MHC) class II  $(A^{g7}, E^{g7})$  and class ID<sup>b</sup> with MHC class II  $(A^d, E^d)$  and class I D<sup>d</sup>, and found them to be nondiabetic and suitable for studies of long-term infection by oral streptococci, as the average lifespan is long compared with that of NOD mice (32). However, susceptibility to bacterial adherence to tooth surfaces in the E2F-1<sup>-/-</sup> mice was seen to be higher than that in NOD.B10.D2 and NOD mice (data not shown). Therefore, we concluded that  $E2F-1^{-/-}$  mice, which have a high sensitivity for initial adhesion, may be more useful than other types previously used for investigation of the initial adherence of streptococci onto tooth surfaces without synthesis of insoluble glucan. S. mutans colonization increased after oral ingestion of 1% sucrose and the affinities of the tested streptococci were considered to be close to those found on tooth surfaces in humans. Therefore, the E2F-1<sup>-/-</sup> mouse is proposed as a useful animal model, as it showed initial adherence activities of oral streptococci on the tooth surfaces in a manner similar to humans with dry mouth.

#### Acknowledgments

This work was supported in part by a Grant-in-Aid for Developmental Scientific Research (1539057) from the Ministry of Education, Science, Sports and Culture of Japan.

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