# **RESEARCH ARTICLE**







Longitudinal study of dental caries incidence associated with *Streptococcus mutans* and *Streptococcus sobrinus* in patients with intellectual disabilities

Yuki Oda, Fumiko Hayashi and Mitsugi Okada<sup>\*</sup>

# Abstract

**Background:** Mutans streptococci (*Streptococcus mutans* and *S. sobrinus*) are considered to be major etiologic agents of dental caries. Using a polymerase chain reaction method, we detected those bacteria from 145 outpatients (6–30 years old) with intellectual disabilities (ID) and their presence was compared with the incidence of dental caries.

**Methods:** Plaque samples were collected from all erupted tooth sites in subjects with a sterile toothbrush. A dental examination was performed to determine the number of decayed and filled teeth (DFT score) in permanent dentition using the WHO caries diagnostic criteria. A Mann–Whitney U-test was employed to compare the caries scores between combinations of the bacteria, and with a Wilcoxon rank test used to compare caries scores between the baseline and after 1 year.

**Results:** Among all subjects, *S. mutans* and *S. sobrinus* were possessed by 78.7 and 83.5 %, respectively, while 13.1 % were positive for *S. mutans* alone, 17.9 % for *S. sobrinus* alone, and 65.6 % for both organisms, with 3.4 % were negative for both. The mean DFT score of subjects positive for both *S. mutans* and *S. sobrinus* at after 1 year was significantly higher than that of those positive for *S. mutans* alone (P < 0.01). The increase in caries increment was also significantly greater in subjects with both bacteria detected (P < 0.001).

**Conclusion:** Our results indicate that patients with ID harboring both *S. mutans* and *S. sobrinus* have a significantly higher incidence of dental caries than those with *S. mutans* alone.

# Background

Mutans streptococci (*Streptococcus mutans* and *S. sobrinus*) are considered to be major etiologic agents of dental caries in humans [1-3]. These bacteria are the most common putative pathogens isolated from human dental plaque and their prevalence has been reported in epidemiological studies [4-7]. Various methods have been used for the detection of putative pathogens, including direct microscopy, cultivation, enzyme tests, enzyme-linked immunosorbent assays and species-specific DNA probes. Several investigators have also developed polymerase chain reaction (PCR) methods and reported them to be more sensitive for detection as

\* Correspondence: mitsugi@hiroshima-u.ac.jp

Department of Special Care Dentistry, Hiroshima University Hospital, 1-2-3, Kasumi, Minami-ku, Hiroshima 734-8553, Japan

compared to conventional culture techniques [8, 9], as they have been shown to be capable of detecting low numbers (5–100) of bacterial cells [9, 10], while they are also quick and relatively simple to perform. Furthermore, PCR assays have been found suitable for specific detection and identification of human cariogenic bacteria, including *S. mutans* and *S. sobrinus* [10, 11].

In previous cross-sectional and longitudinal studies, we reported that preschool children with primary dentition harboring both *S. mutans* and *S. sobrinus* had a significantly higher incidence of dental caries than those with *S. mutans* alone [6, 12]. Recently, we also noted that schoolchildren harboring both *S. mutans* and *S. sobrinus* had a significantly greater dental caries experience in both permanent and primary teeth as compared to those with *S. mutans* alone [13]. Thus, identification and determination of the prevalence of those pathogens



© 2015 Oda et al. **Open Access** This article is distributed under the terms of the Creative Commons Attribution 4.0 International License (http://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated. are of fundamental importance for understanding the initiation and development of the dental caries, and for determining better forms of treatment and prevention. However, few longitudinal studies of the relationship between these two species and caries activities in children [12, 14] or individuals with intellectual disabilities (ID) [15] and Down syndrome [16] have been performed.

Individuals with ID that are cared for at home by family members may also suffer from poor oral hygiene, as higher bacterial counts were found in subjects with ID as compared to those with no disabilities [17]. The methods available to treat and prevent dental caries, and improve oral hygiene individuals with severe disabilities are limited. However, it is important for clinicians to choose from among the various techniques available for prevention of dental caries in their patients with ID according to risk. In the present study, we detected *S. mutans* and *S. sobrinus* in Japanese patients with ID using a PCR method, and then compared their presence with the incidence of dental caries over a 1-year period.

#### Methods

One hundred forty-five outpatients with ID (an Intelligence Quotient (IQ)  $\leq$  70) aged 6 to 30 years and each with mixed or permanent dentition, who visited Hiroshima University Hospital, Hiroshima City were enrolled. The demographic details of the participating outpatients and data on their disability status, IQ, systemic diseases and history of regular medications (if any) were collected from the medical records. Subjects were comprised of 135 with ID and 10 with Down syndrome, and a total of 106 (73.1 %) males and 39 (26.9 %) females were examined (Table 1). Consent for participation was obtained from at least one of their parents prior to the study according to the ethical guidelines of the Declaration of Helsinki (1975) and ethical clearance was obtained from the Ethical Committee of Hiroshima University (Epidemiology-No. 34). Each subject underwent a dental examination performed in the Special Care Dental Clinic by a single well-trained dentist (M.O.) while seated in a dental chair in a supine position, using the WHO caries diagnostic criteria to

Table 1 Study population by age and sex

Age	Male		Female		Total
(years)	No.	%	No.	%	
6–10	12	92.3	1	7.7	13
11-15	19	86.4	3	13.6	22
16-20	21	77.8	6	22.2	27
21-25	32	69.6	14	30.4	46
26-30	24	64.9	13	35.1	37

determine the decayed and filled teeth (DFT) index [18]. Those who had received antibiotics within the previous 3 months or with systemic diseases were excluded.

Dental plaque was collected from all erupted teeth by brushing with a sterile toothbrush for 1 min using a previously described method [19]. During toothbrushing, plaque adhering to the toothbrush was removed by washing several times in a tube of sterile distilled water. The plaque samples in the tube were immediately transported to our research laboratory and stored at -20 °C, prior to extraction of genomic DNA.

Streptococcus mutans JCM5175 and S. sobrinus ATCC 27607 were used as control species. PCR detection of the target species was performed using primers described by Igarashi et al. [8, 10]. Oligonucleotide primers were designed to the *dex* DNA sequence of S. mutans (GenBank accession no. D49430) and S. sobrinus (GenBank accession no. M96978). For S. mutans, the forward primer, 5' TAT GCT GCT ATT GGA GGT TC 3', is complementary to the sequence 973–992, and the reversed primer, 5' AAG GTT GAG CAA TTG AAT CG 3', is complementary to the sequence 2225–2244. For S. sobrinus, the forward primer, 5' TGC TAT CTT TCC CTA GCA TG 3', is complementary to the sequence 134–153, and the reversed primer, 5' GGT ATT CGG TTT GAC TGC 3', is complementary to the sequence 1726–1743.

The primers for eubacteria 16S ribosomal RNA sequence (GenBank accession number M75035) were used to confirm the presence of bacteria in plaque samples as positive control [20]. The forward primer, 5' CAG GAT TAG ATA CCC TGG TAG TCC ACG C 3', is complementary to the sequence 783–810, and the reversed primer, 5' GAC GGG CGG TGT GTA CAA GGC CCG GGA ACG 3', is complementary to the sequence 1378–1407. The size of the expected PCR product was 625 bp.

Plaque samples were first subjected to centrifugation at 1600 × g for 20 min. Next, supernatant was discarded, and individual cell pellets were stored at -20 °C until DNA isolation, for which the pellets were resuspended in 180 µl of enzymatic lysis buffer (20 mM Tris–HCl, pH 8.0; 2 mM EDTA; 1.2 % Triton X-100, 20 mg/ml lysozyme). A genomic DNA preparation from each plaque sample was obtained using a DNeasy<sup>®</sup> Blood and Tissue Kit (Qiagen, Austin, TX, USA) for DNA extraction of Gram-positive bacteria, to which we added an RNase treatment [21]. DNA concentrations in the dental plaque samples were determined by measuring  $A_{260}$ , while quality was estimated using the  $A_{260}/A_{280}$  ratio [22].

Polymerase chain reaction amplification was performed in a reaction mixture (25  $\mu$ l) consisting of PCR beads (GE Healthcare UK Limited, Little Chalfont, Buckinghamshire, UK) that contained an enzyme (two units of *Taq* DNA polymerase) and the required reagents, as well as 25 pmol of each primer, and 20 to 50 ng of the template DNA solution in a thermal cycler (DNA Engine PTC-220 DYAD<sup>TM</sup>, MJ Research, Hatoboro, PA, USA). Each set of PCR analyses included a negative control (water blank) in addition to the positive control. The reaction mixture was denatured at 95 °C for 3 min, followed by 26 cycles of denaturation at 95 °C for 1 min, annealing at 55 °C for 1 min, and extension at 72 °C for 1 min, with a final cycle of 94 °C for 1 min, 55 °C for 1 min, and 72 °C for 5 min [8]. Following amplification, 15  $\mu$ l of the PCR products was analyzed by electrophoresis on a 1.0 % agarose gel and, after staining with ethidium bromide, the newly synthesized DNA fragments were visualized under ultraviolet light at 302 nm. The sizes of the PCR products were estimated from the electrophoretic migration of products relative to a 100-base ladder marker (Amersham Pharmacia Biotech, AB, Uppsala, Sweden). The sensitivity of the PCR method was tested by using known amounts of purified S. mutans JCM5175 and S. sobrinus ATCC27607 as a template. Serially diluted DNA was also used as a template, and the detection limit was 1 pg of DNA for S. mutans (12 cells) and 100 fg of template DNA for S. sobrinus (9 cells) (data not shown).

Descriptive statistics and statistical analyses were performed using a software statistical package (SPSS 14.0, Inc., Chicago, IL, USA). A Mann–Whitney U-test was employed to compare the caries scores between subjects possessing the various combinations of bacteria, and a Wilcoxon rank test to compare caries scores between the baseline and 1 year later.

## Results

Table 2 shows caries prevalence in subjects with ID possessing *S. mutans* alone or in combination with *S. sobrinus* at baseline and after 1 year. *S. mutans* and *S. sobrinus* were found in 78.7 and 83.5 %, respectively, of our subjects with ID. In addition, 13.1 % were positive for *S. mutans* alone, 17.9 % for *S. sobrinus* alone, and 65.6 % for both, while 3.4 % were negative for both *S. mutans* and *S. sobrinus*. The DFT score in those positive

for both organisms after 1 year was significantly higher than that of those positive for *S. mutans* alone (P < 0.01). Furthermore, that score in subjects positive for both after 1 year was significantly higher than at baseline (P < 0.001) and the increase of DFT score in those positive for both was significantly greater than in subjects positive for *S. mutans* alone (P < 0.01).

Table 3 shows subjects with previous caries experience at baseline and caries incremental increases after 1 year, along with the combination of mutans streptococci detected. Seventeen (89.5 %) with *S. mutans* alone, and 86 (90.5 %) with both *S. mutans* and *S. sobrinus* had past caries experiences. Four (21.1 %) of the subjects with *S. mutans* alone, 9 (34.6 %) with *S. sobrinus* alone, and 54 (55.7 %) with both had increases in caries increment, while none of the subjects possessing neither organism showed an increase in increment. Also, approximately 60 % of the subjects with both had an increase in increment caries over the 1-year study period, whereas approximately 20 % of those with *S. mutans* alone demonstrated such an increase.

PCR analysis with 16S rRNA primers confirmed the presence of bacterial DNA in all plaque samples (data not shown).

# Discussion

Individuals who are intellectually disabled and cared for at home by family members often suffer from poor oral hygiene, as they have been reported to have higher bacterial counts as compared to individuals with no disabilities [17]. The methods available to treat and prevent dental caries, and also improve oral hygiene for patients with severe disabilities are limited, though it is important for clinicians to choose a suitable technique for prevention of dental caries based on the risk of their development in their patients with ID.

We performed the present longitudinal study to compare the presence of mutans streptococci with the incidence of dental caries in Japanese patients with ID over a 1-year period using a PCR method. The average concentration of DNA recovered from brushing plaque samples was approximately 700  $\mu$ g/ml, which was deemed

Table 2 Caries prevalence in patients with S. mutans alone or in combination with S. sobrinus at baseline and after 1 year

Oraganism pre	esent		Mean (SD) DFT		Increase in DFT	Range
S. mutans	S. sobrinus	No. of subjects (%)	Baseline	After 1 year	Mean (SD)	
+	-	19 (13.1)	6.47 (4.64)	6.95 (4.61)*	0.47 (1.22)*	0–5
-	+	26 (17.9)	5.96 (5.03)	6.42 (5.33)	0.46 (0.71)	0-2
+	+	95 (65.6)	8.65 (7.60)†	10.31 (8.13)* †	1.65 (2.04)*	0-9
-	-	5 (3.4)	0.60 (1.34)	0.60 (1.34)	0.00 (0.00)	0

 $\label{eq:statistical significance between groups: *P<0.01, Mann-Whitney U-test; +P<0.001, Wilcoxon rank test.$ 

Data for DFT scores are presented as mean (SD)

Organism present		Previous caries experience		Caries incremental increase	
S. mutans	S. sobrinus	No.	%	No.	%
+	-	17	89.5	4	21.1
-	+	22	84.6	9	34.6
+	+	86	90.5	54	55.7
-	-	1	20.0	0	0.0

**Table 3** Patients with previous caries experience at baseline and incremental increases after 1 year, and the combination of mutans streptococci detected

sufficient for performing this PCR-based survey. To ensure presence of a representative bacterial sample in all cases and absence of PCR inhibiting substances, we performed a broad-range PCR assays applying eubacterial 16S rRNA-based primers and subjecting all samples obtained by the toothbrushing method. This confirmed the presence of bacteria and bacterial DNA in all plaque samples (data not shown). This method has been shown to be a more sensitive means of detection of cariogenic microorganisms as compared with conventional culture techniques [8–11].

The present study results indicate that the prevalence of mutans streptococci in patients with ID aged 6 to 30 years old is 96.6 %, which is in agreement with similar surveys conducted with school children in other parts of the world [23-26], while the prevalence of mutans streptococci in adults has been reported to be greater than 60 % [27] and 82.7 % in subjects with Down syndrome aged 1-48 years old [16]. The prevalence of S. sobrinus alone in the present study was higher than in those studies. However, those studies employed cultural methods and previous findings have shown that mitissalivarius bacitracin agar inhibits the growth of S. sobrinus to a greater degree than that of S. mutans [28, 29]. Thus, the contrasting results may have been due to the more sensitive isolation and detection method used for S. sobrinus in our study as compared to those employed in the past, as well as differences in type and stage of carious lesions in the subjects. Nevertheless, since the present cohort was limited, additional studies are required.

The DFT score in those positive for both organisms at after 1 year was significantly higher than that of those positive for *S. mutans* alone. Furthermore, that score in subjects positive for both after 1 year was significantly higher than at baseline and the increase of DFT score in those positive for both was significantly greater than in subjects positive for *S. mutans* alone. The caries incremental increase for subjects possessing both organisms was 3.5 times higher than that for those with *S. mutans* alone, indicating that individuals with ID possessing both have higher caries activities that those with only *S. mutans*. In our previous study, the increase in dft score

for primary teeth in preschool children possessing both was four times greater than for those possessing *S. mutans* alone [12]. The present PCR results showed that subjects with ID possessing both *S. mutans* and *S. sobrinus* had a significantly higher caries incidence and increment increase as compared to those with only *S. mutans* at after 1 year, which also agrees with the results of previous studies of schoolchildren aged 11–12 years old [30, 31], as well as studies of preschool children aged 3–5 years old [6, 12, 32]. In contrast, there was no significant difference in baseline score for DFT between subjects positive for both *S. mutans* and *S. sobrinus* and those with *S. mutans* alone, though the DFT score of those positive for both *S. mutans* and *S. sobrinus* showed a trend to be higher than that of those with *S. mutans* alone.

The present subjects were being treated as outpatients and were referred from local dental clinics because of their poor dental cooperation and already had untreated decayed teeth. All decayed teeth were immediately treated in the Special Care Dental Clinic, thus the caries incremental increase in subjects positive with both S. mutans and S. sobrinus was significantly higher than in those with S. mutans alone. In previous studies, subjects who possessed both bacteria had increases in caries increment over a 1-year period [14, 33]. The results of that former study suggested that the prevalence of S. sobrinus was more closely associated with future caries activity, especially with an increase in smooth-surface caries increment, than the prevalence of S. mutans. However, the subject population in the present study was limited, thus further studies are required. In addition, it remains unknown how mixed colonization increases the cariogenicity of bacteria, while quantitative studies using a real-time quantitative PCR are required to understand the mechanisms of increased caries risk for future studies.

Seventeen (89.5 %) with *S. mutans* alone, and 86 (90.5 %) with both *S. mutans* and *S. sobrinus* had past caries experiences. Four (21.1 %) of the subjects with *S. mutans* alone, 9 (34.6 %) with *S. sobrinus* alone, and 54 (55.7 %) with both had increases in caries increment, while none of the subjects possessing neither organism

showed an increase in increment. Also, approximately 60 % of the subjects with both had a caries incremental increase over the 1-year study period, whereas approximately 20 % of those with *S. mutans* alone demonstrated such an increase.

## Conclusions

In conclusion, the present longitudinal study results indicate that individuals with ID harboring both *S. mutans* and *S. sobrinus* have a significantly higher incidence of dental caries than those positive for *S. mutans* alone.

#### **Competing interests**

The authors declare that they have no competing interests.

#### Authors' contributions

YO, FH and MO desighned the study. MO performed the dental examination. YO and MO collected plaque samples. YO and FH performed the conventional PCR assay. YO and MO were responsible for data analysis. All authors participated in the manuscript processing and approved the final version.

#### Received: 1 May 2015 Accepted: 24 August 2015 Published online: 02 September 2015

#### References

- Masuda N, Tsutsumi N, Sobue S, Hamada S. Longitudinal survey of the distribution of various serotypes of *Streptococcus mutans* in infants. J Clin Microbiol. 1979;10:497–502.
- Loesche WJ. Role of *Streptococcus mutans* in human dental decay. Microbiol Rev. 1986;50:353–80.
- Marsh PD, Featherstone A, McKee AS, Hallsworth AS, Robinson C, Weatherell JA, et al. A microbiological study of early caries of approximal surfaces in schoolchildren. J Dent Res. 1989;68:1151–4.
- Hamada S, Slade HD. Biology, immunology, and cariogenicity of Streptococcus mutans. Microbiol Rev. 1980;44:331–84.
- Whiley RA, Beighton D. Current classification of the oral streptococci. Oral Microbiol Immunol. 1998;13:195–216.
- Okada M, Soda Y, Hayashi F, Doi T, Suzuki J, Miura K, et al. PCR detection of *Streptococcus mutans* and *S. sobrinus* in dental plaque samples from Japanese pre-school children. J Med Microbiol. 2002;51:443–7.
- Takahashi N, Nyvad B. The role of bacteria in the caries process: Ecological perspectives. J Dent Res. 2011;90:294–303.
- Igarashi T, Yamamoto A, Goto N. PCR for detection and identification of Streptococcus sobrinus. J Med Microbiol. 2000;49:1069–74.
- Ono T, Hirota K, Nemoto K, Fernandez EJ, Ota F, Fukui K. Detection of Streptococcus mutans by PCR amplification of spaP gene. J Med Microbiol. 1994;41:231–5.
- Igarashi T, Yamamoto A, Goto N. Direct detection of *Streptococcus mutans* in human dental plaque by polymerase chain reaction. Oral Microbiol Immunol. 1996;11:294–8.
- Shiroza T, Shinozaki N, Watanabe T, Ikemi T, Fukushima K, Abiko Y. Rapid isolation of chromosomal DNA from oral streptococci and polymerase chain reaction-oriented restriction fragment-length polymorphism analysis for genetic heterogeneity. Oral Microbiol Immunol. 1998;13:11–6.
- Okada M, Soda Y, Hayashi F, Doi T, Suzuki J, Miura K, et al. Longitudinal study of dental caries incidence associated with *Streptococcus mutans* and *Streptococcus sobrinus* in pre-school children. J Med Microbiol. 2005;54:661–5.
- Okada M, Kawamura M, Oda Y, Yasuda R, Kojima T, Kurihara H. Caries prevalence associated with *Streptococcus mutans* and *Streptococcus sobrinus* in Japanese schoolchildren. Int J Paediatr Dent. 2012;22:342–8.
- Hirose H, Hirose K, Isogai E, Miura H, Ueda I. Close association between Streptococcus sobrinus in the saliva of young children and smooth-surface caries increment. Caries Res. 1993;27:292–7.
- Mäkinen KK, Isotupa KP, Kivilompolo T, Mäkinen PL, Murtomaa S, Petäjä J, et al. The effect of polyol-combinant saliva stimulants on *S. mutans* levels in plaque and saliva of patients with mental retardation. Spec Care Dentist. 2002;22:187–93.

- 16. de Castilho AR, Pardi V, Pereira CV. Dental caries experience in relation to salivary findings and molecular identification of *S. mutans* and *S. sobrinus* in subjects with Down syndrome. Odontology. 2011;99:162–7.
- 17. Loesche WL. Dental caries: a treatable infection. Sprinfield, Illisois: Charles C Thomas; 1982.
- 18. WHO. Basic Methods. 4th ed. Geneva: World Health Organization; 1997.
- Okada M, Hayashi F, Nagasaka N. Detection of Actinobacillus actinomycetemcomitans and Porphyromonas gingivalis in dental plaque samples from children 2 to 12 years of age. J Clin Periodontol. 2000;27:763–8.
- Goncharoff P, Figurski DH, Stevens RH, Fine DH. Identification of Actinobacillus actinomycetemcomitans: polymerase chain reaction amplification of IktA-specific sequences. Oral Microbiol Immunol. 1993;8:105–10.
- 21. Smith GL, Socransky SS, Smith CM. Rapid method for the purification of DNA from subgingival microorganisms. Oral Microbiol Immunol. 1989;4:47–51.
- 22. Sambrook J, Fritsch E, Maniatis T, editors. Molecular cloning: A Laboratory Manual. New York: Cold Spring Habor Laboratory; 1989.
- Köhler B, Bjarnason S. Mutans streptococci, lactobacilli and caries prevalence in 11- and 12-year-old Icelandic children. Community Dent Oral Epidemiol. 1987;15:332–5.
- 24. del Rio Gl. Dental caries and mutans streptococci in selected groups of urban and native Indian schoolchildren in Mexico. Community Dent Oral Epidemiol. 1991;19:98–100.
- el-Nadeef MA, Adegbembo AO, Adeyinka A. Distribution of mutans streptococci among Nigerian school children. Afr Dent J. 1994;8:11–5.
- Rodis OM, Matsumura S, Kariya N, Okazaki Y, Ogata S, Reissmann DR. Culture-based PCR analysis of plaque samples of Japanese school children to assess the presence of six common cariogenic bacteria and its association with caries risk. Mol Cell Probes. 2009;23:259–63.
- Lindquist B, Emilson CG. Colonization of *Streptococcus mutans* and *Streptococcus sobrinus* genotypes and caries development in children to mothers harboring both species. Caries Res. 2004;38:95–103.
- Jordan HV. Cultural methods for the identification and quantitation of *Streptococcus mutans* and lactobacilli in oral samples. Oral Microbiol Immunol. 1986;1:23–30.
- de Soet JJ, van Dalen PJ, Pavicic MJ, de Graaff J. Enumeration of mutans streptococci in clinical samples by using monoclonal antibodies. J Clin Microbiol. 1990;28:2467–72.
- Ahmady K, Marsh PD, Newman HN, Bulman JS. Distribution of *Streptococcus* mutans and *Streptococcus sobrinus* at sub-sites in human approximal dental plaque. Caries Res. 1993;27:135–9.
- Köhler B, Bjarnason S, Finnbogason SY, Holbrook WP. Mutans streptococci, lactobacilli and caries experience in 12-year-old Icelandic urban children, 1984 and 1991. Community Dent Oral Epidemiol. 1995;23:65–8.
- Seki M, Yamashita Y, Shibata Y, Torigoe H, Tsuda H, Maeno M. Effect of mixed mutans streptococci colonization on caries development. Oral Microbiol Immunol. 2006;21:47–52.
- Straetemans MM, van Loveren C, de Soet JJ, de Graaff J, ten Cate JM. Colonization with mutans streptococci and lactobacilli and the caries experience of children after the age of five. J Dent Res. 1998;77:1851–5.

# Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

) BioMed Central

Submit your manuscript at www.biomedcentral.com/submit