Purification and Characterization of Extracellular Glucosyltransferase from *Streptococcus mutans* Serotype *b* (Subspecies *rattus*)

By HIDEFUMI KUMADA, TOSHIO UMEMOTO, MASAO ONISI, HIDEAKI TSUMORI, ATSUNARI SHIMAMURA, AND HIDEHIKO MUKASA

1 Department of Oral Microbiology, Kanagawa Dental College, 82, Inaoka-cho, Yokosuka, Kanagawa 238, Japan
2 Department of Oral Microbiology, School of Dentistry, Showa University, 1-5-8, Hatanodai, Shinagawa, Tokyo 142, Japan
3 Department of Chemistry, National Defense Medical College, 2, Namiki 3-chome, Tokoroza, Saitama 359, Japan

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An extracellular glucosyltransferase (GT-S) synthesizing water-soluble glucan was purified from the culture supernatant of *Streptococcus mutans* BHT (serotype *b*, subsp. *rattus*) by DEAE-Sepharose chromatography and preparative isoelectric focusing. The *M*<sub>r</sub> of the enzyme was 155000 and the pI was 4.5. The GT-S had a specific activity of 10.2 i.u. (mg protein)<sup>-1</sup>, an optimum pH of 6.0 and a *K*<sub>m</sub> value of 0.8 mM for sucrose, and was activated twofold by dextran T10. The GT-S was immunologically partially identical with the corresponding enzymes in crude preparations from serotypes *c*, *e* and *f*. The glucan synthesized *de novo* from sucrose by the GT-S was water-soluble and consisted of 29 mol% of non-reducing terminal, 49 mol% of 1,6-α-linked, 11 mol% of 1,3-α-linked and 11 mol% of 1,3,6-α-branched glucose residues.

INTRODUCTION

Glycosyltransferases from *Streptococcus mutans* convert dietary sucrose into adhesive water-insoluble polysaccharides which subsequently produce dental plaque (Gibbons & Nygaard, 1968; Mukasa & Slade, 1973). These polysaccharides are formed by cooperative action of water-soluble glucan-synthesizing glucosyltransferase (GT-S), insoluble glucan-synthesizing glucosyltransferase (GT-I), polysaccharide, and, in some *S. mutans* serotypes, fructosyltransferase (Carlsson, 1970; Mukasa & Slade, 1974; Mukasa et al., 1979; Ciardi et al., 1977; Fukushima et al., 1981). Strains of *S. mutans* serotype *a* (subsp. *cricetus*) and serotypes *d* and *g* (subsp. *sobrinus*) secrete three kinds of glucosyltransferase, while strains of serotype *b* (subsp. *rattus*) and serotypes *c*, *e* and *f* (subsp. *mutans*) secrete glucosyl- and fructosyltransferases (Ciardi et al., 1976; Shimamura et al., 1983; Tsumori et al., 1983; Mukasa, 1986). The glycosyltransferases have been purified from serotypes *a* to *g* except for serotype *b* (Mukasa et al., 1982a, 1985; Fukui et al., 1982; Shimamura et al., 1982, 1983; Koga et al., 1983; Kuramitsu & Wondrack, 1983; Sato et al., 1984; Tsumori et al., 1985a, b). Glycosyltransferase complexes have been obtained from a serotype *b* strain, which contained both glucosyl- and fructosyltransferase activities (Scales et al., 1975). It was essential, therefore, to purify and characterize the glycosyltransferases from a serotype *b* strain to clarify the differences among glycosyltransferases from all the *S. mutans* serotypes. In this paper, we describe purification procedures for the

Abbreviations: GT-S, glucosyltransferase synthesizing water-soluble glucan; GT-S1, glucosyltransferase synthesizing highly branched 1,6-α-glucan; GT-S2, glucosyltransferase synthesizing 1,6-α-glucan with lesser amounts of 1,3-α-linked and 1,3,6-α-branched glucose residues; GT-I, glucosyltransferase synthesizing water-insoluble glucan (1,3-α-glucan); IEF, isoelectric focusing; PAS, periodic acid-Schiff; PMSF, phenylmethanesulphonyl fluoride; APMSF, (p-aminophenyl)methanesulphonyl fluoride hydrochloride.

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GT-S from *S. mutans* BHT (serotype b, subsp. rattus), its physicochemical and kinetic characteristics, and the structure of the glucan synthesized by this enzyme from sucrose.

**METHODS**

**Strains and culture conditions.** *Streptococcus mutans* BHT (serotype b, subsp. rattus), originally isolated from human dental plaque by Zinner *et al.* (1965), was provided by Dr S. Hamada (Department of Dental Research, National Institute of Health, Shinagawa-ku, Tokyo, Japan). Cells were cultured in 5:1 chemically defined medium (Terleckyj *et al.*, 1975) containing 0.05% Tween 80 at 37°C for 10 h. The OD$_{550}$ of the culture was 4.5 and the pH was 4.7. The culture supernatant obtained by centrifugation at 5500 g at 5°C was made 0.1 mM and 0.01% with respect to PMSF and Merthiolate, respectively.

**Purification of glucosyltransferase.** (1) Ethanol fractionation. The culture supernatant (5:3 l) was treated with 50% (v/v) ethanol below 0°C. The precipitate collected by centrifugation at 5500 g for 30 min was dissolved in 52 ml distilled water and then used as the crude enzyme preparation.

(2) DEAE-Sepharose chromatography. The ethanol fraction (52 ml) obtained above was mixed with 270 ml 10 mM-sodium maleate buffer, pH 6:0, containing 1 mM-APMSF, 0.1 mM-PMSF and 0.1 µg α2-macroglobulin ml$^{-1}$, and then applied to a DEAE-Sepharose (Pharmacia) column (1.8 x 3 cm) previously equilibrated with the same buffer containing 50 mM-NaCl and the enzyme was eluted with a linear gradient of 50 to 300 mM-NaCl in this buffer (1 litre) at room temperature (approximately 20°C). The column flow rate was 73 ml h$^{-1}$ and fraction volumes of 5 ml per tube were collected. The active fractions (nos 145–165) were combined and concentrated by adsorption to a DEAE-Sepharose column (1 x 3 cm) previously equilibrated in 150 mM-NaCl, 0.1 mM-APMSF, 1 mM-AMPSF and 0.1 µg α2-macroglobulin ml$^{-1}$ and by elution with 350 mM-NaCl in the same buffer. The concentrated enzyme solution was dialysed against 5 mM-sodium maleate buffer, pH 6.0, containing 1 µM-APMSF at 0°C.

(3) Preparative IEF. The dialysed enzyme solution (4:1 ml) was subjected to preparative IEF. The density gradient was formed manually (Vesterberg, 1971) with glycerol. Ampholine pH 4–6, Ampholine pH 3.5–10 and Triton X-100 were used at concentrations of 0.95, 0.05 and 0.1%, respectively. The focusing was carried out at a constant wattage of 4 W for 3 h and then 8 W for 15.3 h at approximately 0°C. The active fractions (nos 57–65) were pooled and stored at –40°C.

**Enzyme assay.** The enzyme activity was measured as previously reported (Mukasa *et al.*, 1979) in 1 ml 0.1 M-sodium phosphate buffer (pH 6.5) or 0.1 M-sodium maleate buffer (pH 6.0) containing 41.8 mM-sucrose and 0.01% Merthiolate with or without 0.34 mg dextran T10. The reaction was linear for at least 12 h under these assay conditions.

The reducing sugar released from sucrose was measured by the method of Somogyi (1945), using fructose as a standard. One unit (i.u.) of total glucosyltransferase activity is defined as the amount of enzyme that releases 1 µmol reducing sugar from sucrose min$^{-1}$ at 37°C.

Polysaccharides were collected, washed as described by Mukasa *et al.* (1979) and measured by the phenol/sulphuric acid method (Dubois *et al.*, 1956), using glucose as a standard. Fructan was measured by the method of Van Handel (1967) with some modifications as follows: 0.2 g anthrone was dissolved in a mixture of 95 ml concentrated sulphuric acid (18 M) and 5 ml distilled water, and subsequently added with 20 ml distilled water at 4°C. A sample solution, adjusted to 0.5 ml with distilled water, and 3 ml of the anthrone reagent were mixed at 4°C, and kept at 45°C for 20 min. After the mixture had been cooled to room temperature, the OD$_{550}$ was measured, using fructose as a standard. The amount of fructan was subtracted from the total amount of polysaccharides to give the amount of glucan. One unit (i.u.) of GT-S activity was the amount of enzyme catalysing the incorporation of 1 µmol glucose from sucrose into glucan min$^{-1}$.

The effect of dextran T10 on glucosyltransferase activity was tested with 0–0.5 mg dextran ml$^{-1}$ in 0.1 M-sodium maleate buffer (pH 6.0) containing 41.8 mM-sucrose.

**Analytical IEF.** This was done by the method of Mukasa *et al.* (1982b) in the presence of 0.1% Triton X-100 by using the low pi calibration kit (Pharmacia) as pi marker proteins. After focusing, protein was stained with Coomassie brilliant blue R-250. GT-S activity and reducing-sugar-release activity were directly detected in the gel as previously described (Mukasa *et al.*, 1982b).

**SDS-PAGE.** This was done by the method of Weber & Osborn (1969) in a 5% (w/v) gel for 5.5 h at 5 mA per gel. Protein was stained with Coomassie brilliant blue R-250 and $M_r$ was determined, using Bio-Rad high $M_r$ protein standards. The enzyme activity in the gel was detected as previously reported (Russell, 1979b; Mukasa *et al.*, 1982b).

**Immunological procedure.** The crude enzyme preparations from serotypes a to g were obtained by ethanol fractionation as described by Mukasa *et al.* (1985). Antisera against the crude and purified glucosyltransferases were prepared from male New Zealand White rabbits as previously described (Mukasa & Slade, 1973). A double immunodiffusion test was done by the Ouchterlony technique (Ouchterlony, 1958) using 0.5% agarose gel.
Glucosyltransferase of S. mutans serotype b

Determination of optimum pH and $K_m$ value. The optimum pH was estimated by the method of Tsumori et al. (1985a). The $K_m$ value for sucrose was estimated by measuring the activity in 0.1 M-sodium maleate buffer (pH 6.0) containing 0.5-50 mM-sucrose by the direct linear plot method (Eisenthal & Cornish-Bowden, 1974). The values quoted represent the mean of values from triplicate experiments.

Linkage analysis of glucan. The water-soluble glucan was synthesized by the purified enzyme (14 µg) in 3.2 ml 0.1 M-sodium maleate buffer (pH 6.0) containing 5% (w/v) sucrose and 0.01% Merthiolate at 37 °C for 8 h. The synthesized glucan was washed (Mukasa et al., 1979), dialysed against distilled water and lyophilized. The glucan (2.5 mg) was permethylated by the method of Hakomori (1964), hydrolysed, reduced with sodium borohydride, and acetylated. The resultant partially methylated alditol acetates were analysed by GLC (Shimamura et al., 1982).

Protein assay. Protein was determined by the Lowry method using bovine serum albumin as a standard. The protein content of the purified enzyme was also estimated by staining the gel after analytical IEF with Coomassie brilliant blue R-250 and by scanning with a densitometer (Chromatoscanner CS-910, Shimadzu) using bovine serum albumin as a standard.

Materials. DEAE-Sepharose CL-6B and dextran T10 were purchased from Pharmacia, PMSF from Calbiochem-Behring, APMSF from Wako Pure Chemicals, and Ampholines from LKB. All other chemicals were of analytical grade.

RESULTS

Purification of GT-S

The GT-S from S. mutans BHT, serotype b, which synthesized water-soluble glucan from sucrose, was purified 16-fold with a 23% recovery as summarized in Table 1. The GT-S fraction eluted from the DEAE-Sepharose column was contaminated by trace amounts of a minor GT-S species and fructosyltransferase (Figs 1 and 2). Further purification of the GT-S by preparative IEF gave a single active peak. This preparation synthesized more than 99.7% (w/w) of glucan from sucrose. Dextranase and inulinase activities were not detected in the preparation, as measured by using dextran T10 and inulin as substrates.

![Fig. 1. IEF patterns of the glycosyltransferase preparations from S. mutans BHT (serotype b). Protein was stained with Coomassie brilliant blue R-250 (a) and the polysaccharides synthesized by the glycosyltransferases were stained with PAS reagent (b). Lanes: 1, pl marker proteins (10 µg); 2, culture supernatant (80 µg for activity); 3, crude enzyme preparation (4 µg for protein, 23 µg for activity); 4, DEAE-Sepharose fraction (0.9 µg, 0.2 µg); 5, purified GT-S (0.2 µg, 0.02 µg). The pl values were determined by reference to the following pl markers: human carbonic anhydrase B (6.55); bovine carbonic anhydrase B (5.85); β-lactoglobulin A (5.20); soybean trypsin inhibitor (4.55); glucose oxidase (4.15); amyloglucosidase (3.50); and pepsinogen (2.80).]
Fig. 2. SDS-PAGE patterns of the glycosyltransferase preparations from S. mutans BHT (serotype b). Protein was stained with Coomassie brilliant blue R-250 (a), and glycosyltransferase activity was stained with PAS reagent (b). 1, M, markers (2.5 μg); 2, culture supernatant (160 μg for activity); 3, crude enzyme preparation (8 μg for protein and activity); 4, DEAE-Sepharose fraction (3 μg); 5, purified GT-S (0.7 μg). The M, was estimated by using the following proteins: myosin (200000), β-galactosidase (1 16 300), phosphorylase b (92 500), and bovine serum albumin (66 200).

Fig. 3. (a) Immunodiffusion of the various glucosyltransferase preparations against anti-GT-S(b) serum (centre well; 15 μl). 1, Purified GT-S(b) (1 μg); 2–6, crude enzyme preparation from strains: BHT (b), Ingbritt (c), LM7 (e), OMZ 175 (f) and HS6 (a), respectively (30–100 μg protein). (b) Immunodiffusion of the purified GT-S(b) (centre well; 1 μg) to various antisera (15 μl each). 1, Anti-GT-S(b); 2, anti-crude enzyme(c); 3, anti-GT-S(e); 4, anti-GT-I(c); 5, anti-GT-S1(a); 6, anti-GT-S2(a).
**Glucosyltransferase of *S. mutans* serotype b**

### Table 1. Purification of GT-S from *S. mutans* BHT (serotype b, subsp. rattus)

<table>
<thead>
<tr>
<th>Step</th>
<th>Total protein (mg)</th>
<th>Activity* (i.u.)</th>
<th>Specific activity (i.u. mg⁻¹)</th>
<th>Recovery (%)</th>
<th>Purification (fold)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ethanol fractionation</td>
<td>40.0</td>
<td>25.1</td>
<td>0.63</td>
<td>100</td>
<td>1</td>
</tr>
<tr>
<td>DEAE-Sepharose</td>
<td>1.2</td>
<td>6.3</td>
<td>5.3</td>
<td>25.1</td>
<td>8.4</td>
</tr>
<tr>
<td>Preparative IEF</td>
<td>0.56</td>
<td>5.7</td>
<td>10.2</td>
<td>22.7</td>
<td>16.2</td>
</tr>
</tbody>
</table>

* GT-S activity was determined by measuring the amount of glucan synthesized.

### Properties of GT-S

**Electrophoretic properties.** The final GT-S preparation (preparative IEF preparation) consisted of a main GT-S species with a negligible amount of other GT-S species as found by analytical IEF (Fig. 1). Minor amounts of Coomassie-brilliant-blue-staining materials were apparent on SDS-PAGE gels (Fig. 2); these materials may have been derived from the Ampholine mixture used, since the broad bands disappeared when the gel was dialysed before staining (data not shown). The $M_r$ of the major GT-S was 155000 ± 2000 (mean ± SD of five experiments) by SDS-PAGE and the pI was 4.5 (mean of three experiments) (Figs 1 and 2). The GT-S activity stained with PAS coincided in position with reducing-sugar-release activity stained with 2,3,5-triphenyltetrazolium chloride (data not shown).

**Immunological analysis.** The purified GT-S and crude enzyme from strain BHT (serotype b) formed a single precipitin band with an anti-GT-S(b) serum (Fig. 3a). Crude enzymes from strains Ingbritbt (c), LM7 (e) and OMZ 175 (f) also formed a single precipitin band with the anti-GT-S(b) serum, but these enzymes were partially identical with the GT-S(b). Crude enzymes from HS6 (a) (Fig. 3a), B13 (d) and 6715 (g) (not shown) did not react with the anti-GT-S(b) serum. The purified GT-S(b) also reacted with an anti-GT-S(c) serum, but did not react with anti-GT-I(c), anti-GT-S1(a), anti-GT-S2(a) (Fig. 3b) and anti-GT-I(a) sera (not shown).

**Kinetic study.** The optimum pH was 6.0 and the $K_m$ value for sucrose was 0.8 ± 0.2 mM (mean ± SD) at pH 6.0. The enzyme activity was stimulated twofold by dextran T10 (0.5 mg ml⁻¹).

**Linkage analysis of glucan.** The glucan synthesized was converted into partially methylated alditol acetates and analysed by GLC. The acetate derivatives of 2,3,4,6-tetra-, 2,4,6-tri-, 2,3,4-tri- and 2,4-di-O-methyl-D-glucitol were found in the molar ratio of 1.0:0.4:1.7:0.4, indicating that the glucan was composed of 49 mol% of 1,6- and 11 mol% of 1,3-α-linked glucose with 11 mol% of 1,3,6-α-branch points.

### DISCUSSION

In our preliminary experiments, the attempts to obtain a homogeneous preparation of glucosyltransferase from a culture supernatant of *S. mutans* serotype b (subsp. rattus) always resulted in a low yield of the activity and in multiple forms of the enzyme different in size and charge, as reported by Ciardi et al. (1976). This was probably due to the action of extracellular proteases present during the purification procedures (Grahame & Mayer, 1984). These problems were largely overcome by the addition, after culture, of PMSF, a serine protease inhibitor (Russell et al., 1983). In the present study, APMSF and $\alpha_2$-macroglobulin were added, in addition to PMSF, in order to further inhibit the protease actions after culture and during the purification procedures, and a GT-S was finally obtained in a high state of purity (Figs 1 and 2) from the culture supernatant of *S. mutans* BHT (serotype b, subsp. rattus) by DEAE-Sepharose chromatography and preparative IEF, although a trace amount of the other GT-S species, probably derived from the major GT-S species (pI 4.5) by the extracellular proteases, was present in the purified GT-S preparation (Figs 1 and 2).

The GT-S from serotype b was immunologically partially identical with the GT-Ss from serotype c, e and f strains, and immunologically unrelated to the GT-I from serotype c strain and to the GT-S1, GT-S2 and GT-I from serotype a, d and g strains. These results were compatible with those in the inhibition test of enzyme activity by antisera (Smith & Taubman, 1977) and
double immunodiffusion tests (Russell, 1979a; Kuramitsu & Wondrak, 1983), although the GT-S from serotype c was reported to be immunologically unrelated to the crude enzyme preparation from serotype b (Kuramitsu & Ingersoll, 1976).

The glucan synthesized from sucrose by the GT-S(b) was water-soluble and mostly consisted of 1,6-α-linked glucose residues with 1,3-α-linked and 1,3,6-α-branched glucose residues. This glucan closely resembled the glucans synthesized by GT-S2s purified from serotype a, d and g strains (Shimamura et al., 1983; Koga et al., 1983; Tsumori et al., 1985b) and the glucans by GT-Ss from serotype c, e and f strains (Sato et al., 1984). It may be concluded, therefore, that all the S. mutans serotypes secrete enzymes which synthesize soluble glucans mostly consisting of 1,6-α-linked glucose residues with lesser amounts of 1,3-α-linked and 1,3,6-α-branched glucose residues.

Serotype b strains, which are predominantly found in rat oral cavities, may be the most suitable strains for use in studies on preventive dental caries using rats as experimental animals. Therefore, in addition to the GT-S now purified, characterizations of the other glycosyltransferases secreted from serotype b strains, GT-I and fructosyltransferase, would contribute to the elucidation of the caries induction mechanism of S. mutans on tooth surfaces.

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**REFERENCES**


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