ARTICLE IN PRESS

Journal of Microbiological Methods xxx (2014) xxx-xxx



Contents lists available at ScienceDirect

Journal of Microbiological Methods

journal homepage: www.elsevier.com/locate/jmicmeth



Mutans streptococci enumeration and genotype selection using different

- bacitracin-containing media
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9 ARTICLE INFO

- 10 Article history:
- 11 Received 22 April 2014
- 12 Received in revised form 5 May 2014
- 13 Accepted 6 May 2014
- 14 Available online xxxx
- 15 Keywords:
- 16 Genotypes

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- 17 rep-PCR18 Selective media
- 19 Streptococcus mutans
- 20 Streptococcus sobrinus
- 21 Total streptococci

ABSTRACT

The primary etiological agents associated with dental caries include the mutans streptococci (MS) comprised of 22 *Streptococcus mutans* and *Streptococcus sobrinus*. The effective cultivation and isolation of MS are necessary for the 23 study of MS, including their proper clinical assessment in the epidemiological study of dental caries. Several 24 selective media have been developed for the isolation, enumeration, and characterization of MS. However, 25 inhibition of MS may occur, reducing counts and perhaps limiting selection of some strains. The purpose of 26 this study was to compare five culture media containing bacitracin recommended for the isolation of MS.

Five commonly used bacitracin-containing media (MSB, MSKB, GTSB, TYS20B, and TYCSB) used for MS isolation 28 were quantitatively evaluated. Standard plate counts were performed in duplicate for 2 prototype MS strains 29 (*S. mutans* UA159 and *S. sobrinus* 6715) and for MS isolates from clinical saliva samples obtained from 16 children 30 (approximate age 5 years) to determine total plate counts, and total *S. mutans* count. Selected isolates (*n* = 249) 31 from all of the five media from 5 saliva samples were further confirmed as *S. mutans* with real-time PCR then 32 subsequently evaluated qualitatively with rep-PCR for genotype determination.

All media resulted in variable enumeration with no significant difference in MS counts. MS prototype strains 34 grew well on all five media; clinical isolates demonstrated more variability in counts but no overall significant 35 differences were found. MSB demonstrated comparable ability to grow *S. mutans* but allowed for more non- 36 *S. mutans* growth. All 5 media identified a consistent predominant genotype by rep-PCR. Recovery of minor 37 genotypes was not inhibited by media type.

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1. Introduction

The mutans streptococci (MS) are primary etiological agents associated with the initiation of dental caries (Loesche, 1986). *Streptococcus mutans* and *Streptococcus sobrinus* are the two clinically relevant species

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Abbreviations: DL, DiversiLab™; GSTB, glucose, sucrose, tellurite, bacitracin media; MS, mutans streptococci; MSB, mitis salivarius-bacitracin media; MSKB, mitis salivarius, kanamycin, bacitracin media; OD, optical density; rep-PCR, repetitive extragenic palindromic polymerase chain reaction; SmTC, S. mutans total plate count; THB, Todd Hewitt Broth; TPC, total plate count; TYCSB, tryptone, yeast, cysteine, sucrose, bacitracin media; TYS20B, trypticase soy, sucrose, bacitracin media.

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of MS found in the oral cavity. These two species are typically identified 48 and quantified in "caries epidemiological and early intervention studies 49 (Saravia et al., 2013).

Numerous selective media have been introduced for the isolation, 51 quantitation, and characterization of MS. Typically these media are 52 supplemented with bacitracin and sucrose as selective agents for 53 clinically relevant streptococci (Gold et al., 1973; Hildebrandt and 54 Bretz, 2006). However, unintentional inhibition of MS may occur, 55 reducing counts and perhaps limiting the quantity of detection and 56 isolation of some strains (Hildebrandt and Bretz, 2006; Tanzer et al., 57 1984). Initially, mitis salivarius with bacitracin (MSB) medium was 58 considered a reliable selective media for isolation of MS (Gold et al., 59 1973). As such, MSB medium was selected as the primary selective 60 medium for a large-scale epidemiological study of S. mutans by our 61 laboratory. However, several studies have suggested that MSB medium 62 results in lower MS counts than other available selective media and can 63 possibly result in false-negatives (Saravia et al., 2013; Schaeken, 1986; 64 Tanzer et al., 1984; Van Palenstein Helderman et al., 1983; Wan et al., 65 2002).

http://dx.doi.org/10.1016/j.mimet.2014.05.010 0167-7012/© 2014 Published by Elsevier B.V.

Please cite this article as: Momeni, S.S., et al., Mutans streptococci enumeration and genotype selection using different bacitracin-containing media, J. Microbiol. Methods (2014), http://dx.doi.org/10.1016/j.mimet.2014.05.010

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In this study, five selective media are evaluated: mitis salivariusbacitracin (MSB), mitis salivarius-kanamycin-bacitracin (MSKB), glucose-sucrose-tellurite-bacitracin (GSTB), trypticase soy-sucrosebacitracin (TYS20B), and tryptone-yeast-cysteine-sucrose-bacitracin (TYCSB) (Gold et al., 1973; Kimmel and Tinanoff, 1991; Schaeken, 1986; Tanzer et al., 1984; Van Palenstein Helderman et al., 1983). Detailed descriptions of these media can be found in the literature, but briefly, these media typically are supplemented with bacitracin and high concentrations of sucrose that act as selective agents for clinically relevant MS. Sucrose allows MS to be distinguished while inhibiting other organisms and bacitracin is known to inhibit other oral streptococci (Gold et al., 1973; Hildebrandt and Bretz, 2006; Schaeken, 1986). Mitis salivarius media (in MSB, MSKB) is commonly used as a base medium since it is selective for streptococci and because MS colony morphology is easily distinguished on this media (Gold et al., 1973). Sucrose is replaced by sorbitol in MSKB and the addition of kanamycin was incorporated to enhance the activity of bacitracin (Kimmel and Tinanoff, 1991).

The purpose of this study is to evaluate these five media quantitatively and qualitatively using MS prototype strains and clinical isolates. In addition, evaluation of MSB in comparison to the other four media will be assessed.

2. Material and methods

2.1. MS prototype strains

S. mutans UA159 and *S. sobrinus* 6715 were selected as the MS prototype strains.

2.2. S. mutans clinical strains

For clinical isolates, saliva was collected from pre-school children (n = 16, approximate age 5 years) as part of an ongoing longitudinal epidemiological study of a high caries risk population located in Uniontown, AL, a small rural city. Sample collection was performed at a community health center during a scheduled follow-up visit. Inclusion criteria were that children were free of systemic disease. Human use approval was obtained from the University of Alabama at Birmingham (UAB) Institutional Review Board, with parents of participants providing informed consent, while children gave assent. Whole saliva samples were collected using sterile cotton swabs, which were then stored in 4.5 ml of reduced transport fluid (Syed and Loesche, 1972). Samples were shipped on ice to our laboratory at the University of Alabama at Birmingham School of Dentistry and stored at 4 °C until processed (within 24 h of collection).

2.3. Media & plating

All five media were prepared as originally describe in original publications (Gold et al., 1973; Kimmel and Tinanoff, 1991; Schaeken, 1986; Tanzer et al., 1984; Van Palenstein Helderman et al., 1983).

Prototype strains were inoculated from $-80\,^{\circ}\text{C}$ frozen cultures into 5 ml Todd Hewitt Broth (THB) (Becton Dickinson, Sparks, MD, USA) and incubated anaerobically (10% CO₂, 10% H₂, and 80% N₂) at 37 $^{\circ}\text{C}$ for 24 h. A 500 μ l aliquot of fresh culture was transferred to 4.5 ml of THB and grown to late log phase. Optical density was determined at 600 nm (OD₆₀₀) using a Bio-Rad SmartSpec Plus spectrophotometer (Bio-Rad, Hercules, CA, USA), then adjusted to an OD₆₀₀ of approximately 1.0 with THB, which corresponds to 1×10^9 cells/ml using electronic enumeration (Childers et al., 2011). Bacterial samples were serially diluted to 1×10^{-5} and plated in duplicate on each of the five media using a spiral plater (Spiral Systems Inc., Cincinnati, Ohio, USA) dispensing a 50 μ l volume. This process was repeated in four independent tests.

Clinical saliva samples in reduced transport fluid were vortexed for 20 s on medium speed prior to plating. Undiluted samples were plated

in duplicate on each of the five media using the spiral plater. All plates $\,$ 126 were incubated anaerobically at 37 $^{\circ}\text{C}$ for 48 h. $\,$

2.4. Enumeration 128

After incubation, all plates were counted using a stereomicroscope 129 for total plate counts (TPC) and total *S. mutans* counts (SmTC) by 130 manual counting according to the instructions for the spiral plater 131 (Spiral Systems Inc., Cincinnati, Ohio, USA) to enumerate the colony- 132 forming units per ml (CFUs/ml). Briefly, a quadrant sector-count 133 method was used to enumerate bacteria using a minimum count of 20 134 colonies within a quadrant. Opposite quadrants in the same sector 135 were counted and the bacterial density was found by adding the counts 136 of the two sectors and dividing by the sample volume corresponding to 137 the innermost sector counted.

Plates were randomly selected to perform counts without knowl- 139 edge of counts on other media. The mean plate counts were calculated 140 using duplicate plates. For the clinical isolates, SmTC were recorded 141 based on colony morphology of the prototype strains (Fig. S1) 142 (McGhee et al., 1982; Schaeken, 1986; Tanzer et al., 1984; Van 143 Palenstein Helderman et al., 1983).

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2.5. Genotype evaluation

For clinical samples that demonstrated sufficient *S. mutans* growth 146 on all five selected media (n=5), 10 presumptive *S. mutans* colonies 147 were selected for genotype identification using repetitive extragenic 148 palindromic polymerase chain reaction (rep-PCR) as described previously (Moser et al., 2010). Briefly, colonies were inoculated in 5 ml 150 THB and grown anaerobically at 37 °C for 48 hours. DNA was extracted 151 and checked for purity. All isolates were verified as *S. mutans* using SYBR 152 GreenTM PCR with sequence specific Yoshida primers before performing 153 rep-PCR (Yoshida et al., 2003). Genotypes were determined using the 154 DiversiLabTM (DL) system and cross-referenced with the library of 155 known genotypes from the larger epidemiological study (Cheon et al., 2013).

2.6. Statistical analysis

Raw counts were transformed to approximate normality by taking 159 logarithms of counts (count plus one, i.e. for counts of zero). Ratios of 160 counts of *S. mutans* to total counts were based on raw counts, not on 161 transformed counts. Statistical significance was ascertained using the 162 GLM procedure in SAS version 9.2 using a significance cut-off value of 163 p < 0.05.

3. Results 165

3.1. MS prototype strains

S. mutans UA159 and S. sobrinus 6715 grew equally well on all 167 five media with no significant differences (Tables 1 & S1). For 168 S. mutans UA159, the mean log recovery was marginally better on 169 TYS20B (9.18 \pm 0.10) than on the other media; MSB (9.16 \pm 0.14), 170 GSTB (9.08 \pm 0.19), TYCSB (9.09 \pm 0.12), and MSKB (9.05 \pm 0.16) 171 (p=0.69). In terms of frequency, the highest counts were most 172 often obtained on MSB than other media (2/4 replicates). When 173 MSB was compared to the other media the difference was also non-174 significant (p=0.44).

S. sobrinus 6715 resulted in comparable counts on all five media 176 although at a slightly decreased level as compared to *S. mutans* UA159. 177 The highest counts were most often observed on TYS20B (8.75 \pm 178 0.41) than other media (2/4); MSB (8.79 \pm 0.30), TYCSB (8.67 \pm 179 0.29), MSKB (8.63 \pm 0.25), and GSTB (8.64 \pm 0.35) but not signifiable cantly different (p=0.94) when all five media were compared. In 181

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