

1 Evaluation of Various Culture Media for Detection of Rapidly-  
2 Growing Mycobacteria from Patients with Cystic Fibrosis.

3 Clair L. Preece,<sup>a,b</sup> Thomas A. Wichelhaus,<sup>c</sup> Audrey Perry,<sup>a</sup> Amanda L. Jones,<sup>b</sup> Stephen P.  
4 Cummings,<sup>b</sup> John D. Perry,<sup>a,b</sup>#Michael Hogardt<sup>c</sup>

5  
6 Microbiology Department, Freeman Hospital, Newcastle upon Tyne, UK<sup>a</sup>; Faculty of Health  
7 and Life Sciences, Northumbria University, Newcastle upon Tyne, UK<sup>b</sup>; Institute of Medical  
8 Microbiology and Infection Control, University Hospital Frankfurt, Frankfurt am Main,  
9 Germany<sup>c</sup>

10 #Address correspondence to John. D. Perry, [john.perry@nuth.nhs.uk](mailto:john.perry@nuth.nhs.uk)

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13 **Running title:** Recovery of mycobacteria from cystic fibrosis patients

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21 **ABSTRACT**

22 Isolation of non-tuberculous mycobacteria (NTM) from the sputum of patients with cystic  
23 fibrosis (CF) is challenging due to overgrowth by rapidly-growing species that colonise the  
24 lungs of patients with CF. Extended incubation of *Burkholderia cepacia* selective agar  
25 (BCSA) has been recommended as an expedient culture method for isolation of rapidly-  
26 growing NTM in this setting. The aim of this study was to assess the efficacy of five selective  
27 media designed for the isolation of *Burkholderia cepacia* complex (BCC) along with two  
28 media designed for the isolation of mycobacteria (RGM medium and Middlebrook 7H11  
29 agar) for their ability to isolate NTM. All seven media were challenged with 147 isolates of  
30 rapidly-growing mycobacteria and 185 isolates belonging to other species. RGM medium  
31 was then compared with the most selective brand of BCSA for the isolation of NTM from  
32 224 sputum samples from patients with CF. Different agars designed for isolation of *B.*  
33 *cepacia* complex varied considerably in their inhibition of other bacteria and fungi. RGM  
34 medium supported the growth of all isolates of mycobacteria and was more selective than any  
35 other medium. NTM were recovered from 17 of 224 sputum samples using RGM medium  
36 compared with only seven samples using the most selective brand of BCSA ( $P = 0.023$ ).  
37 RGM medium offers a superior option to other selective agars for isolation of rapidly-  
38 growing mycobacteria from the sputum of patients with CF. Furthermore, the convenience of  
39 using RGM medium enables routine screening for rapidly-growing NTM in all sputum  
40 samples submitted by patients with CF.

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45 **INTRODUCTION**

46 *Burkholderia cepacia* complex (BCC) and non-tuberculous mycobacteria (NTM) are both  
47 recognized as potentially important pathogens when isolated from the lungs of patients with  
48 cystic fibrosis (CF). The use of a selective culture medium is recommended for isolation of  
49 BCC (1) and several brands of such media are commercially available. Rapidly-growing  
50 mycobacteria represent a subset of NTM that generate colonies on solid culture media within  
51 seven days of incubation (2). The predominant species of NTM within the CF population in  
52 Europe is *Mycobacterium abscessus* complex (MABSC) (3-5) and there is convincing  
53 evidence that the prevalence of infection by MABSC is increasing in the CF population (3, 5-  
54 7). This rapidly-growing species comprises three subspecies: *Mycobacterium abscessus*  
55 subsp. *abscessus*, *Mycobacterium abscessus* subsp. *bolletii* and *Mycobacterium abscessus*  
56 subsp. *massiliense*.

57 Culture of mycobacteria from sputum samples traditionally relies upon methods that  
58 were designed to accommodate slow growing mycobacteria – in particular *Mycobacterium*  
59 *tuberculosis*, which is comparatively rare in cystic fibrosis patients. Such methods are  
60 laborious and expensive as they involve chemical decontamination of samples and  
61 subsequent culture on both liquid and solid media (8, 9). Furthermore, contamination of  
62 cultures by faster-growing microorganisms may mean that cultures have to be abandoned (9),  
63 and decontamination protocols may reduce the yield of mycobacteria (8).

64 Esther *et al.* (2011) demonstrated that extended incubation of *Burkholderia cepacia*  
65 selective agar (BCSA) afforded an increased recovery rate of NTM from 0.7% to 2.8% using  
66 routine microbiological culture methods and recommended this as an expedient method for  
67 culture of rapidly-growing mycobacteria from patients with CF (10). The aim of this study  
68 was to assess five commercially-available media designed for the isolation of BCC for their

69 ability to support the growth of BCC ( $n = 43$ ) and rapidly-growing mycobacteria ( $n = 147$ ).  
70 We also assessed the selectivity of these culture media against 142 other bacteria and fungi,  
71 focussing on the inclusion of species frequently recovered from the sputum of patients with  
72 CF. Two agar-based media designed for the isolation of mycobacteria (RGM medium and  
73 Middlebrook 7H11 agar) were included for comparison. The two media with the most  
74 potential to recover mycobacteria were compared for their ability to isolate rapidly-growing  
75 NTM from 224 sputum samples from patients with CF.

## 76 MATERIALS AND METHODS

77 **Materials.** All five media for isolation of BCC were purchased as pre-poured plates from  
78 their respective manufacturers. *Burkholderia cepacia* selective agar (BCSA; Product Ref:  
79 33631) and Cepacia selective agar (Product Ref: 44347) were purchased from bioMérieux,  
80 Basingstoke, UK or bioMérieux, Nürtingen, Germany. *Burkholderia cepacia* agar (Product  
81 Ref: PO0938) was purchased from Oxoid Ltd, Basingstoke, UK. BD Cepacia medium  
82 (Product Ref: 256180) and BD OFPBL (oxidation-fermentation polymyxin-bacitracin-  
83 lactose) medium (Product Ref: 254481) were purchased from BD Diagnostic Systems,  
84 Oxford, UK. Middlebrook 7H11 agar (Product Ref: PP4080) was obtained from E&O  
85 Laboratories, Bonnybridge, UK, as pre-poured plates. Blood agar was prepared from  
86 Columbia agar powder (Oxoid, Basingstoke, UK) and supplemented with 5% defibrinated  
87 horse blood (TSC Biosciences, Buckingham, UK). Sabouraud agar and bacteriological agar  
88 were purchased from Oxoid, Basingstoke, UK. A sample of 9-chloro-9-[4-  
89 (diethylamino)phenyl]-9,10-dihydro-10-phenylacridine hydrochloride (C-390) was kindly  
90 synthesized by Dr. Annette Johnston of Northumbria University, Newcastle upon Tyne, UK  
91 but is also available commercially from Biosynth, Staad, Switzerland. Colistin  
92 methanesulfonate, yeast extract and amphotericin B were kindly provided by bioMérieux,

93 Craponne, France. All other chemicals and antibiotics were purchased from Sigma-Aldrich,  
94 Poole, UK.

95         **Preparation of RGM medium.** RGM (rapidly-growing mycobacteria) medium was  
96 designed and prepared by staff at the Freeman Hospital Microbiology Department (Newcastle  
97 upon Tyne, UK) and is an adaptation of Middlebrook agar described by Middlebrook and  
98 Cohn in 1958 (11). A tenfold-strength solution of Middlebrook 7H9 broth was prepared by  
99 dissolving the following ingredients in 960 ml of deionized water: ammonium sulphate (5 g),  
100 L-glutamic acid (5 g), di-sodium phosphate (25 g), mono-potassium phosphate (10 g),  
101 sodium citrate (1 g), magnesium sulfate (0.5 g), calcium chloride (0.005 g), biotin (0.005 g),  
102 copper sulfate (0.01 g), zinc sulfate (0.01 g), pyridoxine (0.01 g), ferric ammonium citrate  
103 (0.4 g) and 40 ml of glycerol. The pH of the solution was adjusted to 6.6 and it was sterilized  
104 by autoclaving at 116°C for 20 minutes. One hundred milliliters of this solution was added to  
105 750 ml of deionized water with 10 g of agar and 4 g of yeast extract and this was autoclaved  
106 as before and allowed to cool to 50°C in a waterbath. OADC supplement was prepared by  
107 dissolving 5 g bovine serum albumin, 2 g glucose, 0.004 g catalase and 63 µl oleic acid in  
108 100 ml deionized water. The OADC supplement was filter sterilized and added to the molten  
109 agar. Finally, the following additives were each dissolved in 10 ml sterile deionized water  
110 and added to make 1 liter of medium: 32 mg colistin (as colistin methanesulfonate), 400 mg  
111 fosfomycin, 25 mg glucose-6-phosphate, 5 mg amphotericin and 32 mg C-390. Amphotericin  
112 required initial dissolution in 200 µl of N-methyl-2-pyrrolidinone followed by addition of 9.8  
113 ml sterile deionized water. The agar was immediately poured into 90 mm Petri dishes.

114

115         **Microbial strains.** A collection of 147 isolates of rapidly-growing mycobacteria  
116 previously isolated by standard methods from sputum samples from patients with CF was  
117 used for evaluation of all media. These included *Mycobacterium abscessus* subsp. *abscessus*

118 ( $n = 79$ ), *Mycobacterium chelonae* ( $n = 43$ ), *Mycobacterium abscessus* subsp. *massiliense* ( $n =$   
119 12), *Mycobacterium abscessus* subsp. *bolletii* ( $n = 3$ ), *Mycobacterium fortuitum* ( $n = 3$ ),  
120 *Mycobacterium salmoniphilum* ( $n = 3$ ), *Mycobacterium llutzerense* ( $n = 2$ ), *Mycobacterium*  
121 *immunogenum* ( $n = 1$ ) and *Mycobacterium mucogenicum* ( $n = 1$ ). Seventy three of these  
122 isolates were obtained from the Microbiology Department, Freeman Hospital, Newcastle  
123 upon Tyne, UK and all were from distinct patients. Seventeen were kindly provided by St.  
124 Vincent's University Hospital, Dublin, Ireland and were also from distinct patients. The  
125 remaining 57 were consecutive clinical isolates kindly supplied by Public Health England,  
126 Newcastle upon Tyne, UK. The species and subspecies identity of all strains had been  
127 previously confirmed by sequencing of at least two of three housekeeping genes (*rpoB*, *hsp65*  
128 and *sodA*) as previously described (12).

129

130 The collection of non-mycobacteria was selected to represent a variety of species  
131 frequently recovered from the sputa of patients with CF. Non-mycobacterial strains ( $n = 185$ )  
132 were obtained from national culture collections ( $n = 23$ ) or from the culture collection of the  
133 Microbiology Department, Freeman Hospital, Newcastle upon Tyne ( $n = 162$ ) and included  
134 an international *Pseudomonas aeruginosa* reference panel ( $n = 43$ ) (13) and a BCC  
135 experimental strain panel ( $n = 26$ ) (14-16) as well as clinical isolates of both species. In total,  
136 the collection comprised: *Pseudomonas aeruginosa* ( $n = 55$ ), BCC ( $n = 43$ ), *Staphylococcus*  
137 *aureus* ( $n = 28$ ), various species of *Enterobacteriaceae* ( $n = 11$ ), *Achromobacter*  
138 *xylooxidans* ( $n = 8$ ), *Ralstonia mannitolilytica* ( $n = 7$ ), *Stenotrophomonas maltophilia* ( $n =$   
139 4), *Streptococcus* spp. ( $n = 4$ ), *Aspergillus* spp. ( $n = 3$ ), *Candida* spp. ( $n = 3$ ), *Pandoraea* spp.  
140 ( $n = 3$ ), *Acinetobacter* spp. ( $n = 2$ ), *Enterococcus* spp. ( $n = 2$ ), *Inquilinus limosus* ( $n = 2$ ),  
141 *Scedosporium* spp. ( $n = 2$ ), *Bacillus subtilis* ( $n = 1$ ), *Delftia acidovorans* ( $n = 1$ ),

142 *Elizabethkingia miricola* ( $n = 1$ ), *Geosmithia argillacea* ( $n = 1$ ), *Haemophilus influenzae* ( $n =$   
143 1), *Moraxella catarrhalis* ( $n = 1$ ), *Neisseria flavescens* ( $n = 1$ ) and *Ochrobactrum* sp. ( $n = 1$ ).

144

145 **Inoculation of isolates onto culture media.** Strains were previously stored at  $-20^{\circ}\text{C}$   
146 in glycerol/skimmed milk and frozen isolates were subcultured onto Columbia agar with 5%  
147 horse blood prior to testing. Each isolate was suspended in 1 ml of saline (0.85%) to a  
148 turbidity equivalent to a McFarland 0.5 standard (approximately  $1.5 \times 10^8$  CFU/ml) using a  
149 densitometer. For NTM rough colony-types, where clumping occurred, vortexing with three  
150 sterile 3 mm glass beads for 10 min effectively dispersed all clumps. A 1  $\mu\text{l}$  aliquot of each  
151 suspension of mycobacteria was inoculated onto each medium type and the inoculum was  
152 spread using a loop. Filamentous fungi were inoculated in the same way. Suspensions of all  
153 other isolates were inoculated onto media using a multipoint inoculator to deliver inocula of  
154 approximately 1  $\mu\text{l}$  per spot (i.e. approximately  $1.5 \times 10^5$  CFU/spot). All plates were  
155 incubated at  $30^{\circ}\text{C}$  and growth was recorded after four, seven days and ten days of incubation.  
156 To demonstrate the viability of isolates, Columbia blood agar for bacterial isolates and  
157 Sabouraud agar for fungal isolates were used as controls. All tests were performed in  
158 duplicate on separate occasions.

159

160 **Comparison of RGM medium with BCSA (Ref: 33631) for isolation of**  
161 **mycobacteria from sputum samples.** From the data generated in the experiments detailed  
162 above, the most selective brand of BCSA (Ref: 33631) was evaluated against RGM medium  
163 for isolation of mycobacteria from sputum samples. A total of 224 consecutive sputum  
164 samples were prospectively collected from 133 adults and children with CF attending the  
165 Christiane Herzog CF-Centre, University Hospital Frankfurt, Frankfurt am Main, Germany,  
166 between July 2015 and January 2016. Samples were digested using Copan Sputum

167 Liquefying solution in accordance with manufacturer's instructions. After vortexing for 30 s  
168 samples were left for 15 minutes. A 100  $\mu$ L aliquot was then cultured onto RGM and BCSA  
169 (bioMérieux Ref: 33631) and the inoculum was spread to obtain isolated colonies. Both  
170 media were incubated for 10 days at 30°C and examined for growth after 4, 7 and 10 days of  
171 incubation. A minority of samples were read after 11-12 days of incubation if the day of the  
172 final reading fell on a weekend.

173 Colonies were identified by matrix-assisted laser desorption/ionization time-of-flight  
174 mass spectrometry using VITEK-MS IVD system with knowledge database version 2.0  
175 (bioMérieux, Nürtingen, Germany). Suspected isolates of mycobacteria were confirmed as  
176 acid-fast bacilli using a Ziehl-Neelsen stain and identified to species level by sequencing of  
177 the internal transcribed spacer (ITS) region. The ITS region was amplified with primers ITS1  
178 5'-GATTGGGACGAAGTCGTAAC-3' and ITS2 5'-AGCCTGCCACGTCCTTCATC-3'  
179 (TIB MOLBIOL, Berlin, Germany) as previously described (17). PCR was performed in a 50  
180  $\mu$ l reaction mixture with 0.4 pmol/ $\mu$ l of each primer, 1.5 mM MgCl<sub>2</sub>, 0.2 mM of dATP,  
181 dGTP, dCTP, dTTP (Roche, Mannheim, Germany), 1.25 U of Taq polymerase (Invitrogen,  
182 Darmstadt, Germany) and an annealing temperature of 64°C for 40 cycles. The PCR product  
183 was sequenced in both directions using ITS1 and ITS2 and the resulting DNA sequence was  
184 aligned with NCBI sequence database.

185

186 **Statistical analysis.** Any difference in performance of the two media for isolation of  
187 NTM from sputum samples was investigated for statistical significance using McNemar's test  
188 with the continuity correction applied. Statistical significance was assigned to a probability  
189 (*P*) value of  $\leq 0.05$ .

190

191



192 **RESULTS**

193 **Evaluation of seven selective agars for supporting the growth of mycobacteria.** Clear  
194 differences were revealed between the five different brands of BCSA in terms of their ability  
195 to support the growth of mycobacteria. For example, on Cepacia selective agar (bioMérieux)  
196 95.9% of mycobacteria generated growth within 4 days of incubation compared with only  
197 40.1% of isolates on Oxoid *B. cepacia* agar (Table 1). After 10 days of incubation, ten  
198 isolates had still not grown on Oxoid *B. cepacia* agar including MABSC ( $n = 4$ ), *M. chelonae*  
199 ( $n = 3$ ), *M. llatzerense* ( $n = 2$ ) and *M. mucogenicum* ( $n = 1$ ). All isolates were recovered on  
200 Cepacia selective agar (bioMérieux; Ref: 44347) whereas other selective agars for BCC  
201 failed to support the growth of between four and eight isolates. All isolates were recovered on  
202 Middlebrook 7H11 agar and RGM medium.

203

204 **Evaluation of seven selective agars for inhibition of non-mycobacteria.** Table 2  
205 provides insights into the selectivity of the seven selective media with 185 non-mycobacteria.  
206 All five media for BCC showed effective inhibition of *P. aeruginosa*, which is an essential  
207 attribute of such media. Inhibition of other species was more variable however. For example,  
208 of 28 isolates of *S. aureus* (mainly methicillin-resistant strains), 21 (75%) were able to grow  
209 on BD OFPBL medium whereas only three isolates were able to grow on Oxoid *B. cepacia*  
210 agar and bioMérieux BCSA (Ref: 33631). All brands of media for isolation of BCC showed a  
211 poor ability to inhibit the growth of fungi – particularly *Aspergillus* spp. and yeasts. Overall,  
212 bioMérieux BCSA showed the greatest selectivity and BD OFPBL showed the weakest  
213 selectivity against non-mycobacteria among the five brands tested.

214

215 Although Middlebrook selective medium is designed specifically for the isolation of  
216 mycobacteria from clinical samples, growth of non-mycobacterial species was common with

217 75 out of 186 (40.3%) isolates able to grow. Overall, its selectivity was inferior to the two  
218 most selective media for BCC, although it was able to inhibit the growth of *Aspergillus*  
219 *fumigatus*. RGM medium was by far the most selective of all of the agars tested, with 90% of  
220 non-mycobacteria inhibited including all fungi and Gram-positive bacteria (Table 2).

221

222 **Performance of selective agars for supporting growth of BCC.** Of the five brands  
223 of media for BCC, none was able to support the growth of every isolate of BCC within the  
224 standard incubation period of 5 days (Table 3). Cepacia selective agar (bioMérieux; 44347)  
225 showed the highest sensitivity (93%) with only three isolates inhibited (1 *B. stabilis* and 2 *B.*  
226 *multivorans*) whereas the growth of seven isolates was inhibited on Oxoid *B. cepacia* agar (*B.*  
227 *cenocepacia* ( $n = 1$ ), *B. multivorans* ( $n = 5$ ), *B. stabilis* ( $n = 1$ )). Extended incubation up to  
228 ten days resulted in three additional isolates recovered on BD OFPBL medium but had no  
229 impact on other brands of BCSA.

230

231 **Comparison of RGM medium with BCSA (Ref: 33631) for recovery of**  
232 **mycobacteria from sputum samples.** A total of 17 isolates of mycobacteria were recovered  
233 from 224 sputum samples (Table 4). These 17 isolates were derived from a total of 12  
234 patients (prevalence: 12/133; 9%). For four of these 12 patients, the same species was  
235 recovered from more than one sample.

236

237 All 17 isolates of mycobacteria were recovered on RGM medium compared with only  
238 seven (41%) recovered on BCSA ( $P = 0.023$ ). For seven of 12 patients, mycobacteria were  
239 only detected using RGM medium ( $P = 0.023$ ). The calculation of sensitivity in Table 4 is for  
240 comparative purposes only and assumes that all mycobacteria were recovered by at least one  
241 of the two methods. Clearly this cannot be proven and moreover might be considered

242 unlikely for slower-growing species such as *M. avium* complex. The average time-to-  
243 detection for mycobacteria was 7.9 days using RGM medium (range 4-11 days) and 7 days  
244 using BCSA (range 4-11 days).

245

246 Table 4 shows that RGM medium was much more selective than BCSA for the  
247 inhibition of non-mycobacteria with only 17 isolates of non-mycobacteria recovered on RGM  
248 medium from 224 sputum samples (compared with 59 on BCSA). This is highly likely to  
249 have had an impact on the recovery of mycobacteria on BCSA. For example, for the ten  
250 sputum samples shown to contain mycobacteria that were not recovered on BCSA, five of  
251 these showed growth of other bacterial species ( $n = 3$ ) or fungal species ( $n = 2$ ) on BCSA. All  
252 17 isolates of mycobacteria recovered on RGM medium were isolated in pure culture.

253

## 254 DISCUSSION

255

256 The accurate and prompt detection of rapidly-growing NTM for patients with CF is important  
257 for treatment management and for infection control purposes. In the only previously reported  
258 study with RGM medium, a comparison was performed with Cepacia selective agar  
259 (bioMérieux; 44347) for the isolation of mycobacteria from 502 sputum samples.  
260 Mycobacteria were detected in 54 samples using RGM medium and from only 17 samples  
261 using Cepacia selective agar ( $P \leq 0.0001$ ) (18). As media for isolation of *B. cepacia* have  
262 been recommended for isolation of mycobacteria, this prompted us to examine different  
263 commercial brands of such media to compare their ability to support the growth of rapidly-  
264 growing mycobacteria and their selectivity against other flora associated with CF sputum  
265 samples. Cepacia selective agar (bioMérieux; 44347) was at least as effective for the recovery  
266 of pure strains of mycobacteria as any other selective agar for *B. cepacia*. It was less selective

267 than some other agars and much of this could be attributed to lack of inhibition of methicillin-  
268 resistant *S. aureus*. Cepacia selective agar was less selective than bioMérieux BSCA but  
269 more selective than BD OFPBL.

270

271 In 1985, Gilligan *et al.* were the first to report the design of a selective culture  
272 medium for *B. cepacia* (PC medium) for use with sputum samples from patients with CF  
273 (19). Their medium included polymyxin B, ticarcillin, crystal violet and bile salts as selective  
274 agents and such agents are commonly exploited in commercial brands. At around the same  
275 time, Welch *et al.* evaluated the use of OFPBL medium, exploiting the use of polymyxin B  
276 and bacitracin as selective agents (20). Finally, in 1997, Henry *et al.* described *B. cepacia*  
277 selective agar (BCSA) and showed it to have greater selectivity than PC agar and OFPBL  
278 medium. In this medium, polymyxin B and crystal violet were retained as selective agents  
279 with the addition of gentamicin and vancomycin (21). In a large trial with 656 clinical  
280 samples, Henry *et al.* concluded that BCSA was superior to OFPBL and PC medium for  
281 supporting the growth of *B. cepacia* and suppressing the growth of other flora (22). In this  
282 study we re-affirm the high selectivity of BCSA, which was much more selective than  
283 OFPBL, however, six isolates of BCC were inhibited using BCSA. The selective agents  
284 exploited by various pre-poured media commercially available for isolation of BCC are  
285 detailed in Table 5.

286

287 Mycobacteria grow more slowly than most if not all of the other bacterial and fungal  
288 isolates commonly recovered from sputum samples from patients with CF; this means that  
289 high selectivity is extremely important to inhibit or restrict the growth of non-mycobacteria  
290 so that they do not remain undetected due to overgrowth by other species. Although BCSA  
291 was the most selective of the agars designed for recovery of BCC, it was much less selective

292 than RGM medium. If BCC is excluded (as BCSA is designed to detect this species), 25 non-  
293 mycobacteria were able to grow on BCSA compared with only six on RGM medium (Table  
294 1). A particular drawback of selective agars for BCC is their failure to inhibit fungi, and  
295 particularly *Aspergillus* species. On extended incubation of these media, the growth of  
296 *Aspergillus* can overwhelm the entire culture plate severely compromising the isolation of  
297 mycobacteria. This is particularly problematic with sputum samples from CF patients where  
298 infection of mycobacteria has been associated with concomitant isolation of *Aspergillus* sp.  
299 (23, 24).

300

301 Middlebrook 7H11 agar, designed for isolation of mycobacteria, was better at  
302 inhibiting fungi, due to the inclusion of amphotericin (Table 5). However, other species, such  
303 as *Aspergillus terreus* and *Scedosporium apiospermum* remained uninhibited and overall the  
304 selectivity of Middlebrook 7H11 agar was inferior to that of bioMérieux BCSA and Oxoid *B.*  
305 *cepacia* agar (Table 1). In contrast, no yeasts or fungi were able to grow on RGM medium.

306

307 As BCSA (Ref: 33631) was found to be the most selective of the five media designed for  
308 isolation of BCC (and more selective than Middlebrook 7H11 agar), it was compared with  
309 RGM for further evaluation with 224 sputum samples. The study with sputum samples  
310 confirmed the superior selectivity of RGM medium (Table 4) and it is likely that this  
311 facilitated the significantly greater yield of mycobacteria recovered on RGM medium ( $P =$   
312 0.023). We believe that the use of RGM medium constitutes a simple, convenient method for  
313 culture of mycobacteria that can be embedded within routine diagnostic methods allowing the  
314 culture of all submitted sputum samples from patients with CF. A dedicated culture method  
315 for detection of BCC is accepted practice for sputum samples from patients with CF (1) and it  
316 is noteworthy that NTM were recovered in almost three times as many samples as BCC in

317 this study. From our analysis we conclude that RGM medium offers a superior option  
318 compared with any of the other selective agars for screening and monitoring of rapidly-  
319 growing mycobacteria from the sputum of patients with CF. Further studies are required to  
320 compare the sensitivity of RGM medium with formal culture methods for acid-fast bacilli  
321 (AFB), (e.g. automated liquid culture). It would also be of interest to examine the utility of  
322 RGM medium in locations where slower-growing species of mycobacteria, such as *M. avium*  
323 complex, may predominate. Until such data are available, formal AFB culture methods  
324 remain essential in order to detect slow-growing species of NTM (25).

325

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340 **REFERENCES**

- 341 1. **Cystic Fibrosis Trust**. 2010. Laboratory standards for processing microbiological samples  
342 from people with cystic fibrosis. Report of the UK Cystic Fibrosis Trust Microbiology  
343 Laboratory Standards Working Group. ISBN 0-9548511-4-5.
- 344 2. **Brown-Elliott BA, Wallace RJ**. 2011. *Mycobacterium*: clinical and laboratory  
345 characteristics of rapidly-growing mycobacteria, p 525-538. In Versalovic J, Carroll KC,  
346 Jorgensen JH, Funke G, Landry ML, Warnock DW (ed), Manual of Clinical Microbiology,  
347 10th ed, ASM Press, Washington, DC.
- 348 3. **Qvist T, Gilljam M, Jönsson B, Taylor-Robinson D, Jensen-Fangel S, Wang M, Svahn**  
349 **A, Kötz K, Hansson L, Hollsing A, Hansen CR, Finstad PL, Pressler T, Høiby N,**  
350 **Katzenstein TL; Scandinavian Cystic Fibrosis Study Consortium (SCFSC)**. 2015.  
351 Epidemiology of nontuberculous mycobacteria among patients with cystic fibrosis in  
352 Scandinavia. *J Cyst Fibros* **14**:46-52.
- 353 4. **Roux AL, Catherinot E, Ripoll F, Soismier N, Macheras E, Ravilly S, Bellis G, Vibet**  
354 **MA, Le Roux E, Lemonnier L, Gutierrez C, Vincent V, Fauroux B, Rottman M,**  
355 **Guillemot D, Gaillard JL; Jean-Louis Herrmann for the OMA Group**. 2009. Multicenter  
356 study of prevalence of nontuberculous mycobacteria in patients with cystic fibrosis in France.  
357 *J Clin Microbiol* **47**:4124-4128.
- 358 5. **Seddon P, Fidler K, Raman S, Wyatt H, Ruiz G, Elston C, Perrin F, Gyi K, Bilton D,**  
359 **Drobniewski F, Newport M**. 2013. Prevalence of nontuberculous mycobacteria in cystic  
360 fibrosis clinics, United Kingdom, 2009. *Emerg Infect Dis* **19**:1128-1130.
- 361 6. **Bar-On O, Mussaffi H, Mei-Zahav M, Prais D, Steuer G, Stafler P, Hananya S, Blau**  
362 **H**. 2015. Increasing nontuberculous mycobacteria infection in cystic fibrosis. *J Cyst Fibros*  
363 **14**:53-62.

- 364 7. **Cystic Fibrosis Foundation**. Cystic Fibrosis Foundation Patient Registry Annual Data  
365 Report; 2010.
- 366 8. **Burns JL, Rolain JM**. 2013. Culture-based diagnostic microbiology in cystic fibrosis:  
367 Can we simplify the complexity? *J Cyst Fibros* **13**:1-9.
- 368 9. **Whittier S, Hopfer RL, Knowles MR, Gilligan PH**. 1993. Improved recovery of  
369 mycobacteria from respiratory secretions of patients with cystic fibrosis. *J Clin Microbiol*  
370 **31**:861–864.
- 371 10. **Esther CR Jr, Hoberman S, Fine J, Allen S, Culbreath K, Rodino K, Kerr A,**  
372 **Gilligan P**. 2011. Detection of rapidly-growing mycobacteria in routine cultures of samples  
373 from patients with cystic fibrosis. *J Clin Microbiol* **49**:1421-1425.
- 374 11. **Middlebrook G., Cohn, ML**. 1958. Bacteriology of tuberculosis: laboratory methods.  
375 *Am J Public Health Nations Health* **48**:844–853.
- 376 12. **Blauwendraat C, Dixon GL, Hartley JC, Foweraker J, Harris KA**. 2012. The use of a  
377 two-gene sequencing approach to accurately distinguish between the species within the  
378 *Mycobacterium abscessus* complex and *Mycobacterium chelonae*. *Eur J Clin Microbiol*  
379 *Infect Dis* **31**:1847-1853.
- 380 13. **De Soyza A, Hall AJ, Mahenthalingam E, Drevinek P, Kaca W, Drulis-Kawa Z,**  
381 **Stoitsova SR, Toth V, Coenye T, Zlosnik JE, Burns JL, Sa-Correia I, De Vos D, Pirnay**  
382 **JP, T JK, Reid D, Manos J, Klockgether J, Wiehlmann L, Tummler B, McClean S,**  
383 **Winstanley C**. 2013. Developing an international *Pseudomonas aeruginosa* reference panel.  
384 *Microbiologyopen* **2**:1010-1023.



- 385 14. **Mahenthiralingam E, Coenye T, Chung JW, Speert DP, Govan JR, Taylor P,**  
386 **Vandamme P.** 2000. Diagnostically and experimentally useful panel of strains from the  
387 *Burkholderia cepacia* complex. J Clin Microbiol **38**:910-913.
- 388 15. **Coenye T, Vandamme P, LiPuma JJ, Govan JR, Mahenthiralingam E.** 2003.  
389 Updated version of the *Burkholderia cepacia* complex experimental strain panel. J Clin  
390 Microbiol **41**:2797-2798.
- 391 16. **Vermis K, Coenye T, LiPuma JJ, Mahenthiralingam E, Nelis HJ, Vandamme P.**  
392 2004. Proposal to accommodate *Burkholderia cepacia* genomovar VI as *Burkholderia dolosa*  
393 sp. nov. Int J Syst Evol Microbiol **54**:689-691.
- 394 17. **Richter E, Niemann S, Rüsç-Gerdes S, Hoffner S.** 1999. Identification of  
395 *Mycobacterium kansasii* by using a DNA probe (AccuProbe) and molecular techniques. J  
396 Clin Microbiol **37**:964-970.
- 397 18. **Preece CL, Perry A, Gray B, Kenna DT, Jones AL, Cummings SP, Robb A, Thomas**  
398 **MF, Brodlie M, O'Brien CJ, Bourke SJ, Perry JD.** 2015. A novel culture medium for  
399 isolation of rapidly-growing mycobacteria from the sputum of patients with cystic fibrosis. J  
400 Cyst Fibros May 20. pii: S1569-1993(15)00117-4. doi: 10.1016/j.jcf.2015.05.002.
- 401 19. **Gilligan PH, Gage PA, Bradshaw LM, Schidlow DV, DeCicco BT.** 1985. Isolation  
402 medium for the recovery of *Pseudomonas cepacia* from respiratory secretions of patients  
403 with cystic fibrosis. J Clin Microbiol **22**:5-8.
- 404 20. **Welch DF, Muszynski MJ, Pai CH, Marcon MJ, Hribar MM, Gilligan PH, Matsen**  
405 **JM, Ahlin PA, Hilman BC, Chartrand SA.** 1987. Selective and differential medium for  
406 recovery of *Pseudomonas cepacia* from the respiratory tracts of patients with cystic fibrosis. J  
407 Clin Microbiol **25**:1730-1734.

- 408 21. **Henry DA, Campbell ME, LiPuma JJ, Speert DP.** 1997. Identification of *Burkholderia*  
409 *cepacia* isolates from patients with cystic fibrosis and use of a simple new selective medium.  
410 J Clin Microbiol **35**:614-619.
- 411 22. **Henry D, Campbell M, McGimpsey C, Clarke A, Loudon L, Burns JL, Roe MH,**  
412 **Vandamme P, Speert D.** 1999. Comparison of isolation media for recovery of *Burkholderia*  
413 *cepacia* complex from respiratory secretions of patients with cystic fibrosis. J Clin Microbiol  
414 **37**:1004-1007.
- 415 23. **Verregghen M, Heijerman HG, Reijers M, van Ingen J, van der Ent CK.** 2012. Risk  
416 factors for *Mycobacterium abscessus* infection in cystic fibrosis patients; a case-control  
417 study. J Cyst Fibros **11**:340-343.
- 418 24. **Esther CR Jr, Esserman DA, Gilligan P, Kerr A, Noone PG.** 2010. Chronic  
419 *Mycobacterium abscessus* infection and lung function decline in cystic fibrosis. J Cyst Fibros  
420 **9**:117-123.
- 421 25. **Floto RA, Olivier KN, Saiman L, Daley CL, Herrmann JL, Nick JA, Noone PG,**  
422 **Bilton D, Corris P, Gibson RL, Hempstead SE, Koetz K, Sabadosa KA, Sermet-**  
423 **Gaudelus I, Smyth AR, van Ingen J, Wallace RJ, Winthrop KL, Marshall BC, Haworth**  
424 **CS.** 2016. US Cystic Fibrosis Foundation and European Cystic Fibrosis Society consensus  
425 recommendations for the management of non-tuberculous mycobacteria in individuals with  
426 cystic fibrosis: executive summary. Thorax **71**:88-90
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**TABLE 1** Percentage of rapidly-growing mycobacteria recovered on various selective agars at 30°C.

|                               |          | BCSA         | Cepacia<br>selective<br>agar | <i>B.</i><br><i>cepacia</i><br>agar | Cepacia<br>medium | OFPBL         | RGM<br>medium | Middlebrook<br>7H11 agar |
|-------------------------------|----------|--------------|------------------------------|-------------------------------------|-------------------|---------------|---------------|--------------------------|
|                               | <i>n</i> | bioMérieux   | bioMérieux                   | Oxoid                               | BD                | BD            | -             | E&O<br>Laboratories      |
|                               |          | <b>33631</b> | <b>44347</b>                 | <b>PO0938</b>                       | <b>256180</b>     | <b>254481</b> | <b>-</b>      | <b>PP4080</b>            |
| <b>MABSC</b>                  | 94       |              |                              |                                     |                   |               |               |                          |
| Day 4                         |          | 92.6         | 96.8                         | 57.4                                | 96.8              | 93.6          | 98.9          | 98.9                     |
| Day 7                         |          | 98.9         | 98.9                         | 91.5                                | 98.9              | 98.9          | 98.9          | 98.9                     |
| Day 10                        |          | 98.9         | 100                          | 95.7                                | 100               | 98.9          | 100           | 100                      |
| <b><i>M. chelonae</i></b>     | 43       |              |                              |                                     |                   |               |               |                          |
| Day 4                         |          | 97.7         | 100                          | 9.3                                 | 95.3              | 100           | 100           | 100                      |
| Day 7                         |          | 100          | 100                          | 69.8                                | 95.3              | 100           | 100           | 100                      |
| Day 10                        |          | 100          | 100                          | 93                                  | 97.7              | 100           | 100           | 100                      |
| <b>Other species</b>          | 10       |              |                              |                                     |                   |               |               |                          |
| Day 4                         |          | 10           | 70                           | 10                                  | 70                | 40            | 90            | 70                       |
| Day 7                         |          | 30           | 80                           | 60                                  | 70                | 40            | 90            | 70                       |
| Day 10                        |          | 30           | 100                          | 70                                  | 70                | 70            | 100           | 100                      |
| <b>Total<br/>mycobacteria</b> | 147      |              |                              |                                     |                   |               |               |                          |
| Day 4                         |          | 88.4         | 95.9                         | 40.1                                | 94.6              | 91.8          | 98.6          | 97.3                     |
| Day 7                         |          | 94.6         | 98                           | 83                                  | 95.9              | 95.2          | 98.6          | 97.3                     |
| Day 10                        |          | 94.6         | 100                          | 93.2                                | 97.3              | 97.3          | 100           | 100                      |

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**TABLE 2** Number of non-mycobacteria isolates recovered on various selective agars after 10 days of incubation at 30°C.

|  | BCSA       |           | Cepacia selective agar | <i>B. cepacia</i> agar | Cepacia medium | OFPBL      | RGM medium | Middlebrook 7H11 agar |
|--|------------|-----------|------------------------|------------------------|----------------|------------|------------|-----------------------|
|  | <i>n</i>   |           | bioMérieux             | Oxoid                  | BD             | BD         | -          | E&O Laboratories      |
|  | 33631      | 44347     | PO0938                 | 256180                 | 254481         | -          | PP4080     |                       |
| <b>Gram Negatives</b>                            | <b>141</b> | <b>54</b> | <b>60</b>              | <b>55</b>              | <b>59</b>      | <b>72</b>  | <b>18</b>  | <b>63</b>             |
| <i>Enterobacteriaceae</i>                        | 11         | 2         | 0                      | 2                      | 2              | 6          | 0          | 1                     |
| <i>A. xylosoxidans</i>                           | 8          | 3         | 3                      | 3                      | 5              | 8          | 2          | 3                     |
| <i>Acinetobacter</i> sp.                         | 2          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <i>B. cepacia</i> complex                        | 43         | 37        | 40                     | 36                     | 37             | 41         | 12         | 39                    |
| <i>D. acidovorans</i>                            | 1          | 1         | 0                      | 0                      | 0              | 1          | 0          | 0                     |
| <i>E. miricola</i>                               | 1          | 1         | 1                      | 1                      | 0              | 1          | 0          | 1                     |
| <i>H. influenzae</i>                             | 1          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <i>I. limosus</i>                                | 2          | 0         | 2                      | 0                      | 2              | 0          | 1          | 2                     |
| <i>M. catarrhalis</i>                            | 1          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <i>Neisseria flavescens</i>                      | 1          | 1         | 1                      | 1                      | 1              | 1          | 1          | 1                     |
| <i>Ochrobactrum</i> sp.                          | 1          | 0         | 1                      | 1                      | 1              | 1          | 0          | 1                     |
| <i>P. aeruginosa</i>                             | 55         | 0         | 2                      | 1                      | 2              | 2          | 0          | 2                     |
| <i>Pandoraea</i> spp.                            | 3          | 3         | 3                      | 3                      | 3              | 3          | 2          | 3                     |
| <i>R. mannitolilytica</i>                        | 7          | 6         | 6                      | 6                      | 6              | 5          | 0          | 7                     |
| <i>S. maltophilia</i>                            | 4          | 0         | 1                      | 1                      | 0              | 3          | 0          | 3                     |
| <b>Gram Positives</b>                            | <b>35</b>  | <b>3</b>  | <b>11</b>              | <b>3</b>               | <b>14</b>      | <b>21</b>  | <b>0</b>   | <b>7</b>              |
| <i>B. subtilis</i>                               | 1          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <i>Enterococcus</i> spp.                         | 2          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <i>S. aureus</i>                                 | 28         | 3         | 11                     | 3                      | 14             | 21         | 0          | 7                     |
| <i>Streptococcus</i> spp.                        | 4          | 0         | 0                      | 0                      | 0              | 0          | 0          | 0                     |
| <b>Yeast and Fungi</b>                           | <b>9</b>   | <b>5</b>  | <b>8</b>               | <b>9</b>               | <b>8</b>       | <b>8</b>   | <b>0</b>   | <b>3</b>              |
| <i>A. fumigatus</i>                              | 2          | 2         | 2                      | 2                      | 2              | 2          | 0          | 0                     |
| <i>A. terreus</i>                                | 1          | 1         | 1                      | 1                      | 1              | 1          | 0          | 1                     |
| <i>Candida</i> spp.                              | 3          | 2         | 3                      | 3                      | 3              | 3          | 0          | 1                     |
| <i>G. argillacea</i>                             | 1          | 0         | 0                      | 1                      | 0              | 0          | 0          | 0                     |
| <i>S. apiospermum</i>                            | 1          | 0         | 1                      | 1                      | 1              | 1          | 0          | 1                     |
| <i>S. prolificans</i>                            | 1          | 0         | 1                      | 1                      | 1              | 1          | 0          | 0                     |
| <b>Total</b>                                     | <b>185</b> | <b>62</b> | <b>79</b>              | <b>67</b>              | <b>81</b>      | <b>101</b> | <b>18</b>  | <b>73</b>             |
| <b>Total excluding <i>B. cepacia</i> complex</b> | <b>142</b> | <b>25</b> | <b>39</b>              | <b>31</b>              | <b>44</b>      | <b>60</b>  | <b>6</b>   | <b>34</b>             |

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**TABLE 3** No. of isolates of *B. cepacia* complex recovered on various selective agars after 5 days of incubation at 30°C.

|                         |          | <b>BCSA</b> | <b>Cepacia<br/>selective<br/>agar</b> | <b><i>B.</i><br/><i>cepacia</i><br/>agar</b> | <b>Cepacia<br/>medium</b> | <b>OFPBL</b> | <b>RGM<br/>medium</b> | <b>Middlebrook<br/>7H11 agar</b> |
|-------------------------|----------|-------------|---------------------------------------|--|---------------------------|--------------|-----------------------|----------------------------------|
|                         | <i>n</i> | bioMérieux  | bioMérieux                            | Oxoid  | BD                        | BD           | -                     | E&O<br>Laboratories              |
|                         |          | 33631       | 44347                                 | PO0938                                       | 256180                    | 254481       | -                     | PP4080                           |
| <i>B. ambifaria</i>     | 2        | 1           | 2                                     | 2  | 2                         | 1            | 0                     | 0                                |
| <i>B. anthina</i>       | 2        | 1           | 2                                     | 2  | 1                         | 1            | 0                     | 1                                |
| <i>B. cenocepacia</i>   | 11       | 11          | 11                                    | 10   | 9                         | 10           | 3                     | 11                               |
| <i>B. cepacia</i>       | 3        | 3           | 3                                     | 3  | 3                         | 3            | 0                     | 3                                |
| <i>B. contaminans</i>   | 1        | 1           | 1                                     | 1  | 1                         | 1            | 0                     | 1                                |
| <i>B. dolosa</i>        | 2        | 2           | 2                                     | 2  | 2                         | 2            | 0                     | 1                                |
| <i>B. multivorans</i>   | 12       | 10          | 10                                    | 7  | 10                        | 11           | 2                     | 10                               |
| <i>B. pyrrocinia</i>    | 2        | 2           | 2                                     | 2  | 2                         | 2            | 1                     | 1                                |
| <i>B. stabilis</i>      | 4        | 2           | 3                                     | 3  | 3                         | 3            | 0                     | 3                                |
| <i>B. vietnamiensis</i> | 4        | 4           | 4                                     | 4  | 4                         | 4            | 0                     | 4                                |
| Total                   | 43       | 37          | 40                                    | 36   | 37                        | 38           | 6                     | 35                               |
| <b>% recovery</b>       |          | <b>86</b>   | <b>93</b>                             | <b>84</b>                                    | <b>86</b>                 | <b>88</b>    | <b>14</b>             | <b>81</b>                        |

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**TABLE 4** Numbers of isolates of mycobacteria and other species recovered on BSCA and RGM medium from culture of 224 sputum samples.

|  | <b>RGM</b> | <b>BCSA (Ref: 33631)</b> |
|--|------------|--------------------------|
| <b>Total mycobacteria</b>                | <b>17</b>  | <b>7</b>                 |
| <i>M. abscessus</i> complex <sup>a</sup> | 9          | 6                        |
| <i>M. avium</i> complex                  | 1          | 0                        |
| <i>M. chelonae</i>                       | 1          | 0                        |
| <i>M. mucogenicum</i>                    | 2          | 0                        |
| <i>M. simiae</i>                         | 3          | 1                        |
| <i>Mycobacterium</i> species             | 1          | 0                        |
| <b>Sensitivity (%)</b>                   | <b>100</b> | <b>41</b>                |
| <b>Total non-mycobacteria</b>            | <b>17</b>  | <b>59</b>                |
| <i>Achromobacter</i> sp.                 | 6          | 13                       |
| <i>Burkholderia multivorans</i>          | 5          | 7                        |
| <i>Chryseobacterium</i> sp.              | 0          | 1                        |
| <i>Cupriavidus</i> sp.                   | 1          | 1                        |
| <i>Proteus mirabilis</i>                 | 0          | 4                        |
| <i>Pseudomonas aeruginosa</i>            | 0          | 7                        |
| <i>Serratia marcescens</i>               | 0          | 2                        |
| <i>Sphingobacterium spiritivorum</i>     | 0          | 1                        |
| <i>Stenotrophomonas maltophilia</i>      | 0          | 2                        |
| <i>Aspergillus fumigatus</i>             | 2          | 9                        |
| <i>Aspergillus terreus</i>               | 0          | 1                        |
| <i>Candida</i> spp.                      | 1          | 7                        |
| <i>Exophiala dermatitidis</i>            | 0          | 1                        |
| <i>Geotrichum</i> sp.                    | 1          | 1                        |
| <i>Trichosporon mycotoxinivorans</i>     | 1          | 1                        |
| Unidentified fungus                      | 0          | 1                        |
| No growth                                | 190        | 160                      |

458 <sup>a</sup>Sub-speciation of the *M. abscessus* complex was not possible using

459 the ITS sequencing method used for identification in Frankfurt.

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**TABLE 5** Selective agents (per liter) included in various culture media as disclosed by manufacturers<sup>a</sup>

|                 | BCSA       | Cepacia selective agar | <i>B. cepacia</i> agar | Cepacia medium | OFPBL    | RGM medium | Middlebrook 7H11 agar <sup>b</sup> |
|-----------------|------------|------------------------|------------------------|----------------|----------|------------|------------------------------------|
|                 | bioMérieux | bioMérieux             | Oxoid                  | BD             | BD       | -          | E&O Laboratories                   |
|                 | 33631      | 44347                  | PO0938                 | 256180         | 254481   | -          | PP4080                             |
| Polmyxyn B      | 600000 U   | 300 000 U              | 150000 U               | 300000 U       | 300000 U | -          | Included                           |
| Colistin        | -          | -                      | -                      | -              | -        | 32 mg      | -                                  |
| Crystal violet  | 2 mg       | 1 mg                   | 1 mg                   | 1 mg           | -        | -          | -                                  |
| Bile salts      | -          | 0.5 g                  | 1.5 g                  | 0.5 g          | -        | -          | -                                  |
| Ticarcillin     | -          | 10 mg                  | 100 mg                 | 100 mg         | -        | -          | Included                           |
| Gentamicin      | 10 mg      | -                      | 5 mg                   | -              | -        | -          | -                                  |
| Vancomycin      | 2.5 mg     | -                      | -                      | -              | -        | -          | -                                  |
| Bacitracin      | -          | -                      | -                      | -              | 200 U    | -          | -                                  |
| Trimethoprim    | -          | -                      | -                      | -              | -        | -          | Included                           |
| Amphotericin B  | -          | -                      | -                      | -              | -        | 5 mg       | Included                           |
| Malachite Green | -          | -                      | -                      | -              | -        | -          | Included                           |
| Fosfomycin      | -          | -                      | -                      | -              | -        | 0.4 g      | -                                  |
| C-390           | -          | -                      | -                      | -              | -        | 32 mg      | -                                  |

465 <sup>a</sup> The composition of these media may be adjusted by manufacturers to meet performance requirements.

466 <sup>b</sup> Concentrations are not published for selective agents in E&O Middlebrook 7H11 agar.

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