

## Effects of formulation on microbicide potency and mitigation of the development of bacterial insusceptibility

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1                   **The Effect of Formulation on Microbicide**  
2                   **Potency and Mitigation of the Development of**  
3                   **Bacterial Insusceptibility**

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29 Risk assessments into the potential for microbicides to select for reduced bacterial susceptibility  
30 have been based largely on data generated through the exposure of bacteria to microbicides in  
31 aqueous solution. Since microbicides are normally formulated with multiple excipients, we have  
32 investigated the effect of formulation on antimicrobial activity and the induction of bacterial  
33 insusceptibility. The susceptibilities of 9 species of bacteria (7 genera) were determined before  
34 and after repeated exposure (14 passages) using a previously validated gradient plating system,  
35 to the microbicides benzalkonium chloride, benzisothiazolinone, chlorhexidine, didecylmethyl  
36 ammonium chloride, DMDM-hydantoin, polyhexamethylene biguanide, thymol and triclosan in  
37 aqueous solution (non-formulated) and in formulation with excipients often deployed in  
38 consumer products. Susceptibilities were also assessed following an additional 14 passages  
39 without microbicide to determine the stability of any susceptibility changes. Minimum  
40 inhibitory concentrations (MIC) and minimum bactericidal concentrations (MBC) were on  
41 average 11-fold lower for formulated *vs.* non-formulated microbicides. After antimicrobial  
42 exposure, of 72 combinations of microbicide and bacterium, there were 19  $\geq 4$ -fold (mean 8-  
43 fold) increases in MIC for non-formulated and 8  $\geq 4$ -fold (mean 2-fold) increases in MIC for  
44 formulated microbicides. Furthermore, there were 20  $\geq 4$ -fold increases in MBC (mean 8-fold)  
45 for non-formulated and 10  $\geq 4$ -fold (mean 2-fold) increases in MBC for formulated  
46 microbicides. Susceptibility decreases fully or partially reverted back to pre-exposure values for  
47 49% of MICs and 72% of MBCs after further passage. In summary, formulated microbicides  
48 exhibited greater antibacterial potency than unformulated actives and susceptibility decreases  
49 following repeated exposure were lower in frequency and extent.

50

## 51 **INTRODUCTION**

52 Microbicides are broad-spectrum chemical agents that inactivate microorganisms (1-3). They  
53 are widely deployed throughout healthcare (4-6), domestic (7, 8) and industrial environments (9-  
54 11) where their application includes antiseptics (12), hard surface disinfection (13) and  
55 pharmaceutical product preservation (14). They may also be incorporated into medical device  
56 coatings, for instance in sutures (15), wound dressings (16) and urinary catheters (17) to inhibit  
57 bacterial adhesion and subsequent biofilm formation.

58 It has been hypothesized that the use of microbicides could select for bacterial adaptation,  
59 resulting in reduced efficacy of the primary agent as well as potentially decreasing bacterial  
60 susceptibility to chemically-unrelated agents such as other microbicides and antibiotics (18).

61 Whilst there have been reports documenting the laboratory selection of bacteria with decreased

62 microbicide sensitivity following repeated exposure to microbicides in highly selective  
63 conditions, it remains unclear whether this commonly occurs in the environment (19-24).

64 The majority of studies reporting reductions in microbicide susceptibility have used the active  
65 compound in aqueous solution with or without the addition of co-solvents such as DMSO (25)  
66 or ethanol (26, 27). In real use however, microbicides are deployed in formulated products with  
67 multiple excipients that may enhance potency. The potential effect of the formulation of  
68 microbicides on reducing the development of bacterial insusceptibility has received little  
69 research attention. Furthermore, despite the research effort that has been directed towards the  
70 possible risk of induced microbicide insusceptibility, the stability of such susceptibility changes  
71 has been investigated infrequently (24).

72

73 With the ultimate aim of developing realism-based approaches to risk assessment, the current  
74 investigation evaluates the frequency, magnitude and reversibility of susceptibility changes that  
75 may be induced by the repeated exposure of a range of bacteria to microbicides in aqueous  
76 solution or in formulation. The microbicides selected reflect those frequently used in consumer  
77 products such as laundry detergents, hard surface disinfectants and personal care products.  
78 Planktonic susceptibilities (MIC, MBC) and minimum biofilm eradication concentrations  
79 (MBEC) were determined before and after repeated exposure to sub-lethal concentrations of the  
80 microbicides benzalkonium chloride (BAC), benzisothiazolinone (BIT), chlorhexidine (CHX),  
81 didecyldimethyl ammonium chloride (DDAC), glydant (DMDM hydantoin),  
82 polyhexamethylene biguanide (PHMB), thymol, and triclosan in aqueous solution and in  
83 formulation with commonly used sequestrants and surfactants. Bacteria were also passaged  
84 further in the absence of any antimicrobial to determine the stability of any observed change in  
85 susceptibility.

## 86 METHODS

87 **Bacteria.** *Pseudomonas aeruginosa* ATCC 9027, *Staphylococcus aureus* ATCC 6538,  
88 and *Escherichia coli* ATCC 25922 were obtained from Oxoid (Basingstoke, UK). *Acinetobacter*  
89 *baumanii* (Accession number: JX966428.1), *Pseudomonas putida* (Accession number:  
90 JQ968690.1), *Moraxella osloensis* (Accession number: AB643597.1), *Escherichia coli*  
91 (Accession number: CP003034.1) and *Cronobacter sakazakii* (Accession number: HQ880381.1)  
92 were isolated from a domestic kitchen drain biofilm. *Enterococcus faecalis* (Accession number  
93 KJ818115.1) was provided by Angela Oates, The University of Manchester.

94 **Chemical Reagents and Growth Media.** Bacteriological growth media was purchased  
95 from Oxoid (Basingstoke, UK). All other chemical reagents were purchased from Sigma-  
96 Aldrich (Dorset, UK) unless otherwise stated. Bacterial growth media was sterilized at 121°C  
97 and 15 psi for 15 min prior to use. *Pseudomonas aeruginosa*, *Staphylococcus aureus*,  
98 *Escherichia coli* and *Enterococcus faecalis* were cultured on Tryptone Soya Agar and Broth.  
99 *Acinetobacter baumannii*, *Pseudomonas putida*, *Moraxella osloensis* and *Cronobacter sakazakii*  
100 were grown on Wilkins Chalgren agar and broth containing 2% sucrose. All bacteria were  
101 incubated aerobically at 37°C for 18h unless stated otherwise.

102 Antimicrobial actives: benzalkonium chloride, chlorhexidine, thymol and triclosan were  
103 purchased from Sigma-Aldrich (Dorset, UK). Didecyltrimethyl ammonium chloride (50% v/v)  
104 was purchased from Merck Millipore (Durham, UK). Vantocil (a 20% v/v aqueous solution of  
105 PHMB) was obtained from Arch Chemicals Inc. (Manchester, UK). Glydant (DMDM  
106 hydantoin) was obtained from Lonza (Bishop's Stortford, UK). All microbicides were tested in  
107 aqueous solution as previously described (27) and in formulation, at concentrations reflective of  
108 their normal deployment in consumer products. BAC, CHX, DDAC, DMDM hydantoin, PHMB  
109 and thymol were prepared at 1% (v/v) in a general purpose cleaner. Triclosan was formulated

110 into a laundry detergent at 0.0066% (w/v). Benzisothiazolinone was formulated into a laundry  
111 detergent at 0.02% (v/v).

112 **Exposure of Bacteria to Sub-lethal Concentrations of Microbicides as active and**  
113 **formulation.** A previously validated system (20, 25) was used to generate reproducible *c.* 100-  
114 fold antimicrobial concentration gradients on Tryptone Soya Agar plates using a spiral plater  
115 (Whitley Automated Spiral Plater, Don Whitley Scientific, Shipley, UK). Initial MIC  
116 antimicrobial stock solutions (50µl) were deposited on the agar surface. Plates were dried for 1h  
117 at room temperature prior to radial deposition of bacterial pure cultures and then incubated (4d;  
118 37°C) in a static aerobic incubator. After incubation, growth observed at the highest microbicide  
119 concentration was aseptically removed and streaked onto a fresh plate containing the same  
120 antimicrobial concentration gradient. Where growth was observed across the whole  
121 antimicrobial gradient, a new plate produced with a five times higher microbicide concentration  
122 was used<sup>25</sup>. This process was repeated until 14 passages had occurred (P14). Bacteria that  
123 exhibited  $\geq 4$ -fold changes in MIC, MBC or MBEC were then passaged a further 14 times in the  
124 absence of any antimicrobial (X14) to ascertain the stability of adaptation. Bacteria at P0, P14  
125 and X14 were archived for subsequent MIC and MBC testing. Susceptibility testing (MIC,  
126 MBC, MBEC) was performed in two separate experiments each with three technical replicates.

127 **Determination of bacterial Minimum Inhibitory Concentrations (MIC) and**  
128 **Minimum Bactericidal Concentrations (MBC).** MIC values were determined using the  
129 microdilution method as described previously (28). Briefly, overnight bacterial cultures were  
130 adjusted to an OD<sub>600</sub> of 0.8 and diluted 1 in 100 in Tryptone Soya Both or Wilkins Chalgren  
131 Broth with 2% sucrose in a 96-well microtiter plate containing doubling dilutions of the relevant  
132 microbicide. Plates were incubated at 37°C (24h) with agitation (100rpm). The MIC was defined  
133 as the lowest concentration for which bacterial growth did not occur. Growth was viewed as  
134 turbidity (600nm) in comparison to an uninoculated well (negative control) and was detected

135 using a microtiter plate reader (Anthos HTII; Anthos-Labtec Instruments. Salzburg. Austria).  
136 MBCs were determined as stated previously (25), in brief aliquots (10µl) from wells exhibiting  
137 no turbidity were transferred to sterile Tryptone Soya Agar or Wilkins Chalgren Agar prior to  
138 4d incubation at 37°C to determine the minimum bactericidal concentration (MBC) (25). The  
139 MBC was defined as the lowest concentration of microbicide at which no growth occurred after  
140 4d of incubation.

141 **Determination of Minimum Biofilm Eradication Concentrations.** Single species  
142 biofilms were grown on the pegs of a Calgary Biofilm Device (CBD) (29). To produce inocula  
143 for biofilm susceptibility testing, single colonies of test bacteria were inoculated into 10ml of  
144 sterile Tryptone Soya Broth or Wilkins Chalgren Broth with 2% sucrose and incubated at 37°C  
145 in a shaking aerobic incubator (100rpm) for 18h. Cultures were diluted to an OD<sub>600</sub> of 0.8, then  
146 further diluted 1:100 using fresh growth medium. 100µl of bacterial inoculum was added to  
147 each well of the CBD base, plates were then incubated at 37°C and 30 rpm for 48h to allow  
148 biofilm formation on the pegs. Doubling dilutions for microbicides (150µl) were prepared in  
149 sterile broth across a 96 well microtiter plate. Biofilms were exposed to antimicrobials and  
150 incubated for 24h at 37°C and 100rpm. After incubation the lid was transferred to a 96-well  
151 plate containing 200µl of sterile broth and was incubated for 24h at 37°C and 100rpm. Minimum  
152 biofilm eradication concentrations (MBECs) were determined as the lowest concentration for  
153 which bacterial growth did not occur after 18h of incubation. Growth was viewed as turbidity in  
154 comparison to an uninoculated well (negative control) and was detected using a microtiter plate  
155 reader (BioTek, Bedfordshire, UK).

156

## 157 **RESULTS**

158 Two main variables describe data associated with the selection of decreased susceptibility by  
159 exposure to microbicides in the current study; i) the frequency of susceptibility decreases greater

160 than two-fold (25) for multiple test bacteria and microbicides and ii) the extent of susceptibility  
161 changes for each combination of bacterium and microbicide.

162

163 Repeated exposure to the microbicide-containing formulations resulted in a lower frequency of  
164 susceptibility reductions than did exposure to the same microbicide in aqueous solutions and,  
165 where decreases in susceptibility did occur; these were generally smaller for formulated  
166 microbicides. All individual values for bacterial susceptibility before, during and after  
167 microbicide exposure have been given in Tables 1-8. However, due to the large number of  
168 combinations of bacterium and antimicrobial that were tested, the extent of susceptibility has  
169 also been expressed as mean values in the following section.

170

171 After repeated exposure to unformulated microbicides there were 19  $\geq$ 4-fold increases in MIC  
172 (1 of which fully reverted back to pre-exposure values after subsequent passage in the absence  
173 of microbicide, 13 of which partially reverted and 5 which did not revert; average increase in  
174 MIC (P0 to P14) was 11-fold across the test panel of bacteria and microbicides). There were 20  
175 increases in MBC (2 fully, 11 partially and 7 non-revertible; average 8-fold increase) and 17  
176 increases in MBEC (7 fully, 6 partially and 4 non-revertible; average 4-fold increase) after  
177 microbicide exposure (Tables 1-8). After exposure to microbicide containing formulations there  
178 were 8  $\geq$ 4-fold increases in MIC (2 fully and 6 non-revertible; average 2-fold increase), 10  
179 increases in MBC (3 fully, 5 partially and 2 non-revertible; average 2-fold increase) and 16  
180 increases in MBEC (5 fully, 8 partially and 3 non-revertible; average 3-fold increase) (Tables 1-  
181 8). In terms of antimicrobial potency, when comparing the formulated to non-formulated  
182 microbicides across the test panel of bacteria we saw an approximately 11-fold lower MIC/  
183 MBC and 3-fold lower MBEC for the unexposed (P0) bacterial isolates. For the P14 isolates we



184 observed an approximately 35-fold lower MIC, 36-fold lower MBC and 4-fold lower MBEC  
185 (Tables 1-8).

186 **Benzalkonium Chloride.** All test bacteria, with the exception of *M. osloensis*, *C.*  
187 *sakazakii* and the *E. coli* drain isolate exhibited a  $\geq 4$  fold increase in MIC after exposure to BAC  
188 (Table 1). Increases in MBC, whilst generally smaller than those in MIC, were also observed at  
189  $\geq 4$  fold for *S. aureus*, *E. coli* and *P. aeruginosa*. Furthermore  $\geq 4$  fold increases in MBEC  
190 occurred for *S. aureus* and *E. faecalis* after BAC exposure. After growth in the absence of BAC,  
191 subsequent full or partial reversion in MIC, MBC or MBEC occurred for all test bacteria with  
192 the exception of *E. coli* and *P. aeruginosa* (MIC and MBC). In contrast, after exposure to the  
193 BAC formulation only *S. aureus*, *E. coli*, *P. aeruginosa* and *A. baumannii* showed a  $\geq 4$  fold  
194 increase in MIC with *S. aureus* and *E. coli* also demonstrating a  $\geq 4$  fold increase in MBC. *S.*  
195 *aureus*, *E. faecalis* and *P. aeruginosa* also exhibited a  $\geq 4$  fold increase in MBEC after exposure  
196 to BAC formulation. After recovery in the absence of BAC formulation only *S. aureus*  
197 demonstrated any reversion in susceptibility (MBEC).

198 **Benzisothiazolinone (BIT).** No bacterium displayed a substantial change in  
199 susceptibility ( $\geq 4$  fold MIC, MBC or MBEC) to BIT or to BIT formulation after long-term  
200 exposure to the respective agent (Table 2).

201 **Chlorhexidine.** After repeated exposure to chlorhexidine both *S. aureus* and *E. coli*  
202 showed  $\geq 4$  fold increases in MIC and MBC which partially reverted in the absence of the  
203 microbicide (Table 3). *P. aeruginosa* demonstrated a  $\geq 4$  fold increase in MIC which did not  
204 revert after regrowth in a chlorhexidine free environment. *E. faecalis* and *M. osloensis* exhibited  
205  $\geq 4$  fold increases in MBEC, which partially and fully reverted in the absence of chlorhexidine  
206 respectively. In contrast, after exposure to chlorhexidine formulation no bacterium exhibited a  
207  $\geq 4$  fold decrease in susceptibility at MIC, MBC or MBEC level.

208           **Didecyldimethyl Ammonium Chloride.** After repeated DDAC exposure *P. aeruginosa*,  
209 *A. baumannii* and the *E. coli* drain isolate exhibited a  $\geq 4$  fold increase in MBC, of which *P.*  
210 *aeruginosa* fully reverted whilst *A. baumannii* and *E. coli* partially reverted following repeated  
211 growth the absence of DDAC. *S. aureus*, *E. coli*, *E. faecalis* and the *E. coli* drain isolate all  
212 exhibited a  $\geq 4$  fold increase in MBEC, out of which *E. faecalis* and the *E. coli* drain isolate  
213 partially reverted, *E. coli* fully reverted and *S. aureus* did not revert back to pre-exposure values  
214 following growth in the absence of the microbicide (Table 4). After exposure to the DDAC-  
215 containing formulation, *P. aeruginosa* and the *E. coli* drain isolate exhibited a  $\geq 4$  fold increase  
216 in MBC, out of which *E. coli* partially reverted and *P. aeruginosa* fully reverted after passage in  
217 an antimicrobial free environment. In agreement with the changes in MBEC observed after  
218 exposure to DDAC active, *S. aureus*, *E. coli*, *E. faecalis* and the *E. coli* drain isolate also  
219 showed a  $\geq 4$  fold increase in MBEC after exposure to DDAC formulation. MBEC values  
220 partially reverted for both *E. coli* isolates and for *E. faecalis* but did not revert for *S. aureus* after  
221 recovery in the absence of DDAC.

222           **Glydant (DMDM Hydantoin).** The *E. coli* drain isolate exhibited a  $\geq 4$  fold increase in  
223 MBC after repeated exposure to DMDM hydantoin; this susceptibility decrease fully reverted in  
224 the absence of the microbicide (Table 5). Comparatively after exposure to DMDM hydantoin  
225 formulation both *E. coli* isolates as well as *C. sakazakii* showed a  $\geq 4$  fold increase in MBEC, all  
226 of which fully reverted in an antimicrobial free environment.

227           **Polyhexamethylene Biguanide.** *S. aureus*, *E. faecalis* *M. osloensis* and *A. baumannii*  
228 exhibited a  $\geq 4$  fold increase in MIC after PHMB exposure out of which *M. osloensis* and *A.*  
229 *baumannii* fully reverted and *S. aureus* and *E. faecalis* partially reverted after growth in the  
230 absence of PHMB (Table 6). *S. aureus*, *E. coli*, *P. aeruginosa*, *E. faecalis*, and the *E. coli* drain  
231 isolate demonstrated a  $\geq 4$  fold increase in MBC out of which *S. aureus*, *E. faecalis* and the *E.*  
232 *coli* drain isolate showed partial reversion and *E. coli* and *P. aeruginosa* showed no reversion to

233 pre-exposure values in the absence of PHMB. After PHMB exposure, *S. aureus*, *E. faecalis*, *A.*  
234 *baumanii*, *C. sakazakii*, and the *E. coli* drain isolate also displayed a  $\geq 4$  fold increase in MBEC,  
235 which fully reverted for *S. aureus*, *A. baumannii* and *E. coli* drain isolate, and partially reverted  
236 for *E. faecalis* and *C. sakazakii* after re-growth in the absence of PHMB. After exposure to  
237 PHMB formulation *S. aureus*, *E. faecalis* and *P. aeruginosa* showed substantial changes in their  
238 PHMB susceptibility displaying  $\geq 4$  fold increases in MBC all of which fully or partially  
239 reverted in the absence of the antimicrobial formulation. *S. aureus* and *E. faecalis* also exhibited  
240 a  $\geq 4$  fold increase in MBEC after exposure to PHMB formulation, all of which partially reverted  
241 back to pre-exposure values after regrowth in the absence of the formulation.

242 **Thymol.** After long-term thymol exposure none of the bacterial isolates showed a  $\geq 4$   
243 fold decrease in thymol susceptibility at MIC, MBC or MBEC level (Table 7). After exposure to  
244 the thymol-containing formulation, *E. coli* and *A. baumannii* both underwent  $\geq 4$  fold increases in  
245 MBC whilst *P. putida* demonstrated a  $\geq 4$  fold increase in MIC and MBC, all of which partially  
246 reverted in the absence of thymol formulation. Furthermore, both *E. coli* isolates showed a  $\geq 4$   
247 fold increase in MBEC, which partially reverted after growth in the absence of thymol  
248 formulation.

249 **Triclosan.** All bacterial isolates, with the exception of *E. faecalis*, *A. baumannii* and *P.*  
250 *aeruginosa*, which is non-susceptible to triclosan, demonstrated an increase in MIC after  
251 repeated triclosan exposure, none of which fully reverted back to pre-exposure levels after  
252 regrowth in the absence of triclosan (Table 8). All isolates apart from *P. aeruginosa*, *A.*  
253 *baumanii* and *P. putida* showed a  $\geq 4$  fold increase in MBC out of which *C. sakazakii* and the *E.*  
254 *coli* drain isolate showed partial reversion, whilst the others showed no reversion after passage  
255 in the absence of triclosan. Both *E. coli* isolates in addition to *C. sakazakii*, *E. faecalis* and *A.*  
256 *baumanii* showed  $\geq 4$  fold increase in MBEC after repeated triclosan exposure out of which *C.*  
257 *sakazakii* and *E. faecalis* did not revert and both *E. coli* isolates completely reverted in the

258 absence of the microbicide. In comparison after exposure to triclosan formulation only the *E.*  
259 *coli* isolates and *P. aeruginosa* showed  $\geq 4$  fold increase in MIC, which fully reverted for *P.*  
260 *aeruginosa* but did not revert for either *E. coli* strain in the absence of triclosan formulation.  
261 MBECs increased  $\geq 4$  fold for *S. aureus* and *E. faecalis* but fully reverted for both bacteria after  
262 regrowth in the absence of triclosan formulation.

263

## 264 **DISCUSSION**

265         The majority of investigations into the potential of microbicides to select for changes in  
266 bacterial susceptibility have been conducted by exposing pure cultures of bacteria to  
267 microbicides as pure actives in aqueous solution or in simple formulations (aqueous solutions  
268 containing the active and in some studies, cosolvents such as DMSO (25) or ethanol (27)). It has  
269 been hypothesized that formulated products may interact with bacteria in a manner that is  
270 distinct from aqueous solutions (28, 30) potentially reducing the frequency and extent of  
271 susceptibility reductions. Whilst numerous studies have evaluated the antimicrobial potency of  
272 formulated microbicides (3, 31, 32), to our knowledge there are no studies in the literature that  
273 have compared the effects of repeated bacterial exposure to microbicides in aqueous solution  
274 and in complex formulation, for a range of bacteria and microbicides. In the current  
275 investigation therefore, we have evaluated the effect of the formulation of microbicides on  
276 antimicrobial potency and on the mitigation of bacterial insusceptibility for a selection of  
277 bacterial isolates and microbicides encompassing biguanides, quaternary ammonium  
278 compounds, phenolics, isothiazolinones, formaldehyde releasers and essential oils. Microbicides  
279 were tested as aqueous solutions of the active compounds and in complex formulations with  
280 sequestrants and ionic/non-ionic surfactants to mimic their real world use as hard-surface  
281 disinfectants (for BAC, chlorhexidine, DDAC, DMDM hydantoin, PHMB and thymol), and

282 laundry detergents (for BIT and triclosan). The reversibility of any induced susceptibility  
283 changes was also investigated to ascertain the stability of adaptation.

284

285 Reductions in bacterial susceptibility to an antimicrobial agent can be influenced by several  
286 factors related to the antimicrobial or the microorganism. Bacterial susceptibility may be  
287 affected by the structural integrity of the bacterial cell envelope and its ability to function as an  
288 effective permeability barrier (33-35). Innate bacterial non-susceptibility towards an  
289 antimicrobial agent may occur due to effective barrier components of the bacterial cell, such as  
290 an outer membrane in Gram-negative bacteria (36) or the spore coat in bacterial endospores  
291 (37). Changes in cell envelope permeability may therefore affect bacterial susceptibility which  
292 can include alterations in lipopolysaccharide expression and structure<sup>33</sup>, reduction in the  
293 number of outer membrane porins (23) and alterations in membrane fatty acid composition (38).  
294 The expression of efflux pumps has also been linked to decreases in microbicide susceptibility  
295 in bacteria, particularly towards membrane-active compounds such as biguanides (39) (CHX  
296 and PHMB) and quaternary ammonium compounds<sup>40</sup> (BAC and DDAC in the current  
297 investigation). The increased expression of efflux pumps may therefore also provide a plausible  
298 explanation for some of the susceptibility changes observed in many of our bacterial isolates.

299 Reversible susceptibility changes to microbicides may result from temporary phenotypic  
300 adaptations in bacteria, such as the induction of stress responses that revert once the bacteria  
301 recover in an antimicrobial-free environment (41, 42). Equally, the development of microbicide  
302 insusceptibility may be attributable to the selection of insusceptible mutants, for instance  
303 mutations in FabI are reportedly render some bacteria insusceptible to triclosan (43, 44).  
304 However, the inherent stability of a particular mutation largely depends upon the overall fitness  
305 cost that it exerts on the host microorganism versus the competitive advantage that it provides in  
306 a particular environment (45). Hence, any mutation that renders a bacterium less susceptible

307 towards an antimicrobial agent may eventually be lost once the selective pressure is removed if  
308 the mutation results in a biologically significant reduction in the fitness of the microorganism  
309 (46).

310  
311 Whilst previous studies have reported the induction of microbicide insusceptibility in bacteria, it  
312 should be noted that adapted bacterial isolates often remain susceptible to the microbicide at  
313 concentrations used in consumer products, and that true microbicide resistance is likely to be  
314 uncommon (25). In the current investigation, the only test bacterium that was refractory to a  
315 microbicide was *P. aeruginosa* to triclosan. This was apparent before microbicide exposure and  
316 has previously been attributed to the expression of efflux pumps 47. Interestingly this bacterium  
317 was comparatively susceptible to the triclosan formulation, illustrating marked differences in  
318 potency for the microbicide in aqueous solution compared to the formulated product.

319  
320 Out of all the microbicides in unformulated form, BAC and triclosan induced the highest  
321 frequency of  $\geq 4$ -fold increases in MIC with 6/9 bacterial isolates showing a reduction in  
322 susceptibility to both antimicrobials at this level. This was followed by PHMB (4 isolates) and  
323 CHX (3 isolates). Triclosan exposure resulted in the highest frequency of  $\geq 4$ -fold increases in  
324 MBC (6 isolates) followed by PHMB (5 isolates), DDAC and BAC (3 isolates), then CHX (2  
325 isolates) and DMDM hydantoin (1 isolate). In terms of the susceptibility of bacteria when grown  
326 as biofilms, PHMB adaptation resulted in the highest number of isolates showing  $\geq 4$ -fold  
327 increases in MBEC (5 isolates) followed by triclosan and DDAC (4 isolates each) then BAC and  
328 CHX (2 isolates).

329  
330 With respect to the formulated microbicides, BAC induced the highest number of  $\geq 4$ -fold  
331 increases in MIC (4 isolates) followed by triclosan (3 isolates) and thymol (1 isolate). DMDM

332 hydantoin, thymol and PHMB containing formulations induced the largest number of  $\geq 4$ -fold  
333 increases in MBC (3 isolates each) followed by BAC and DDAC (2 isolates each). Exposure to  
334 the DDAC containing formulations resulted in the highest numbers of bacterial isolates  
335 exhibiting a  $\geq 4$ -fold increase in MBEC (4 isolates), followed by BAC and DMDM hydantoin (3  
336 isolates) then PHMB, thymol and triclosan formulations (2 isolates).

337

338 Whilst the current investigation demonstrates that induced reductions in susceptibility towards  
339 both microbicides and microbicide-containing formulations may occur, a substantially higher  
340 number of bacterial isolates underwent  $\geq 4$ -fold increases in MIC, MBC or MBEC when exposed  
341 to microbicides in aqueous solution, in comparison to those in formulation. The only exception  
342 to this was thymol, for which changes in susceptibility were more frequent in bacteria exposed  
343 to the compound in formulation. Thymol is poorly soluble in water and formulation may  
344 therefore have substantially improved solubility, increasing bacterial exposure and thus  
345 selectivity. Furthermore, since incorporating microbicides into formulations frequently  
346 enhanced antimicrobial potency, the formulated microbicides often maintained higher  
347 antimicrobial activity in comparison to microbicides in aqueous solution, even after repeated  
348 exposure. The incorporation of non-ionic surfactants and sequestrants into microbicide-  
349 containing formulations therefore appears to increase antimicrobial potency as well as  
350 mitigating the development of antimicrobial insusceptibility both in terms of frequency and  
351 magnitude of susceptibility change. Since excipients can interact with different cellular targets  
352 to the accompanying microbicide, formulations may have a cumulative antimicrobial effect  
353 which would require multiple further physiological adaptations to render the microorganism  
354 insusceptible.

355

356 Alcohol ethoxylates are a major class of non-ionic surfactants which are often used in household  
357 detergents, cleaners and personal care products and have previously shown bacteriostatic effects  
358 due to their direct impact on the bacterial cell membrane leading to the leakage of cytoplasmic  
359 components, indicating an increase in membrane permeability (48). An increase in membrane  
360 permeability would allow microbicides to more readily transverse the cytoplasmic membrane  
361 increasing their access to intracellular target sites. Therefore combining microbicides and  
362 alcohol ethoxylates in formulation may enhance overall antimicrobial potency, when compared  
363 to the pure active. Sodium tripolyphosphate, a chelating agent commonly used in domestic  
364 detergents, has previously shown antibacterial activity against several bacteria often found as  
365 food contaminants (49). Since sodium tripolyphosphate is a chelating agent it is plausible, as  
366 with other chelators such as EDTA, which this antibacterial activity occurs by disruption of the  
367 bacterial cell envelope through the sequestration of stabilising divalent cations. Such cations  
368 normally link bacterial lipopolysaccharides to the outer membrane and interference with this  
369 process can destabilise the outer membrane in Gram negative bacteria, impairing barrier  
370 function (50-52). Furthermore, strong chelating agents may inhibit bacterial growth by  
371 sequestering trace minerals required for bacterial metabolism (51, 53).

372

373 Essential oils such as thymol are often incorporated into antimicrobial formulation due to their  
374 inhibitory effects on bacterial growth. The antimicrobial activity of essential oils reportedly  
375 occurs through interaction with the bacterial cytoplasmic membrane, resulting in increased cell  
376 permeability and the disruption of energy generation (54, 55). Compensatory adaptations may  
377 occur, but whether these would result in outcome-changing effects during deployment depends  
378 on the extent of any susceptibility decreases, the concentration used in the product and the



379 antimicrobial potency of the formulation (i.e. the active compound and excipients in  
380 combination).

## 381 **CONCLUSION**

382 With the ultimate aim of developing realistic approaches to risk assessment, we observed that  
383 repeated exposure of 9 bacteria to 8 microbicides in aqueous solution or within complex  
384 formulations with sequestrants and ionic/non-ionic surfactants, induced reductions in bacterial  
385 susceptibility in a highly selective laboratory exposure system. Susceptibility changes varied in  
386 reversibility, possibly reflecting a range of underlying mechanisms including temporary  
387 phenotypic adaptation, such as the induction of stress responses or the selection of stable  
388 mutations. Importantly, the formulation of microbicides markedly increased overall  
389 antimicrobial potency for the test microbicides against the majority of the bacteria, as well as  
390 reducing the frequency and magnitude of susceptibility changes. Whilst it remains unclear how  
391 observations based on the *in vitro* exposure of bacteria to microbicides can be extrapolated to  
392 their use in the real world, understanding the potential selectivity of microbicide-containing  
393 formulations is likely to better served by testing formulations as well as actives aqueous  
394 solutions. This highlights the need to conduct risk assessments of induced microbicide  
395 susceptibility changes using conditions that more accurately reflect their deployment.

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401

## 402 TRANSPARENCY DECLARATION

403 Alejandro Amézquita is an employee of Unilever. Peter McClure was an employee of Unilever  
404 when this project was initiated. All other authors: none to declare.

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538

539 **Table 1.** Bacterial susceptibility towards benzalkonium chloride in planktonic and biofilm growth modes before, during and after repeated exposure to  
 540 benzalkonium chloride in aqueous solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	0.1	<b>3.9</b>	<b>2.0</b>	0.5	<b>2.0</b>	<b>2.0</b>	2.0	<b>15.6</b>	<b>7.8</b>	2.0	<b>7.8</b>	<b>7.8</b>	2.6 (1)	<b>31.3</b>	<b>15.6</b>	3.9	<b>125</b>	7.8
<i>E. coli</i> †	4.6 (1)	<b>31.3</b>	<b>31.3</b>	3.9	<b>31.3</b>	<b>31.3</b>	7.2 (2)	<b>41.7 (16)</b>	<b>62.5</b>	7.8	<b>31.3</b>	<b>62.5</b>	31.3	31.3	62.5	31.3	62.5	62.5
<i>E. faecalis</i> †	2.0	<b>7.8</b>	3.9	2.0	3.9	3.9	3.3 (1)	7.8	7.8	3.9	7.8	7.8	6.5 (1)	<b>31.3</b>	7.8	6.7 (2)	<b>46.9 (17)</b>	<b>46.9 (17)</b>
<i>P. aeruginosa</i> †	14.3 (2)	<b>62.5</b>	<b>62.5</b>	15.6	<b>62.5</b>	<b>125</b>	23.4 (9)	<b>125</b>	<b>125</b>	31.3	62.5	<b>250</b>	125	250	<b>500</b>	62.5	<b>250</b>	<b>500</b>
<i>M. osloensis</i> *	3.9	2.0	na	1.0	1.0	na	7.8	15.6	na	2.0	2.0	na	7.8	na	na	7.8	2.0	na
<i>A. baumannii</i> *	2.0	<b>62.5</b>	<b>31.3</b>	3.9	<b>31.3</b>	<b>31.3</b>	93.8 (34)	250	125	62.5	62.5	125	125	250	125	125	125	93.8 (34)
<i>P. putida</i> *	15.6	<b>62.5</b>	31.3	15.6	15.6	na	125	125	62.5	62.5	31.3	na	125	na	62.5	125	31.3	na
<i>C. sakazakii</i> *	62.5	52.1 (16)	na	31.3	31.3	na	125	125	na	31.3	31.3	na	31.3	na	na	31.3	62.5	na
<i>E. coli</i> *	18.4 (7)	52.1 (16)	na	15.6	31.3	na	62.5	125	na	31.3	31.3	na	62.5	na	na	62.5	62.5	na

541 MIC, minimum inhibitory concentration; MBC, minimum bactericidal concentration; MBEC, minimum biofilm eradication concentration.  
 542 Before antimicrobial exposure (P0); during antimicrobial exposure (P14) and after passage in the absence of antimicrobial (X14) All values are in mg/L. †,  
 543 non-drain isolates; \*, drain isolates. UF, unformulated (microbicide in aqueous solution); F, formulated (microbicide in formulation). Organisms that  
 544 underwent a  $\geq 4$ -fold increase in MIC, MBC or MBEC (as indicated by bold text) were passaged a further 14 times in the absence of microbicide. na, bacteria  
 545 that did not undergo a  $\geq 4$ -fold change and were not assessed for reversibility. Data represents six replicates. Where data varied between biological replicates,  
 546 standard deviations have been given in parentheses. In controls were bacteria were tested against formulations without microbicide, all bacteria were non-  
 547 susceptible to in-use concentrations.  
 548

549

550 **Table 2.** Bacterial susceptibility towards benzisothiazolinone in planktonic and biofilm growth modes before, during and after repeated exposure to  
 551 benzisothiazolinone in aqueous solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	7.8	15.6	na	1.0	2.0	na	31.3	62.5	na	15.6	15.6	na	62.5	62.5	na	31.3	62.5	na
<i>E. coli</i> †	15.6	15.6	na	7.8	7.8	na	31.3	62.5	na	31.3	31.3	na	250	187.5 (68)	na	125	125	na
<i>E. faecalis</i> †	7.8	15.6	na	0.5	1.0	na	7.8	7.8	na	0.5	1.0	na	250	41.7 (16)	na	125	125	na
<i>P. aeruginosa</i> †	125	250	na	15.6	31.3	na	250	500	na	62.5	125	na	500	500	na	125+	125+	na
<i>M. osloensis</i> *	1.0	1.0	na	0.5	0.5	na	1.0	1.0	na	0.5	0.5	na	2.0	2.0	na	0.5	1.0	na
<i>A. baumannii</i> *	31.3	31.3	na	7.8	15.6	na	31.3	62.5	na	31.3	62.5	na	250	250	na	62.5	125	na
<i>P. putida</i> *	15.6	31.3	na	31.3	31.3	na	62.5	62.5	na	31.3	62.5	na	250	250	na	62.5	125	na
<i>C. sakazakii</i> *	7.8	7.8	na	7.8	7.8	na	31.3	31.3	na	31.3	31.3	na	250	500	na	62.5	125	na
<i>E. coli</i> *	15.6	31.3	na	15.6	15.6	na	62.5	62.5	na	15.6	31.3	na	250	187.5	na	125	125	na

552 See footnote in Table 1

553

554

555

556 **Table 3.** Bacterial susceptibility towards chlorhexidine in planktonic and biofilm growth modes before, during and after repeated exposure to chlorhexidine  
 557 in aqueous solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	1.7 (1)	<b>7.8</b>	3.9	2.0	2.0	na	5.2 (2)	<b>46.9 (17)</b>	<b>31.3</b>	7.8	7.8	na	13 (4)	31.3	31.3	7.8	15.6	na
<i>E. coli</i> †	2.4 (1)	<b>11.7 (4)</b>	7.9	2.0	3.9	na	9.8 (5)	<b>62.5</b>	<b>31.3</b>	15.6	31.3	na	52.1 (16)	62.5	31.3	62.5	31.3	na
<i>E. faecalis</i> †	3.9	7.8	15.6	3.9	7.8	na	14.3 (3)	31.3	31.3	7.8	15.6	na	31.3	<b>125</b>	62.5	31.3	62.5	na
<i>P. aeruginosa</i> †	7.8	<b>31.3</b>	<b>31.3</b>	7.8	15.6	na	68.8 (34)	<b>250</b>	125	125	125	na	250	125	125	250	125	na
<i>M. osloensis</i> *	3.9	2.0	2.0	1.0	1.0	na	31.3	15.6	3.9	1.0	1.0	na	31.3	<b>125</b>	15.6	15.6	31.3	na
<i>A. baumannii</i> *	7.8	7.8	na	3.9	7.8	na	125	62.5	na	15.6	31.3	na	125	125	na	125	31.3	na
<i>P. putida</i> *	7.8	7.8	na	4.6 (2)	3.9	na	93.8 (34)	62.5	na	7.8	7.8	na	62.5	125	na	62.5	62.5	na
<i>C. sakazakii</i> *	7.8	7.8	na	3.9	3.9	na	62.5	125	na	7.8	15.6	na	62.5	125	na	31.3	10.4 (4)	na
<i>E. coli</i> *	7.8	10.4 (4)	15.6	3.9	3.9	na	46.8 (17)	125	125	7.8	15.6	na	125	125	125	62.5	23.4 (9)	na

558 See footnote in Table 1

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564 **Table 4.** Bacterial susceptibility towards didecyldimethyl ammonium chloride in planktonic and biofilm growth modes before, during and after repeated  
 565 exposure to didecyldimethyl ammonium chloride in aqueous solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	
<i>S. aureus</i> †	0.5	1.0	1.0	0.5	0.5	0.5	2.0	3.9	3.9	2.0	0.5	0.5	3.9	<b>31.3</b>	<b>31.3</b>	3.9	62.5	62.5
<i>E. coli</i> †	7.8	11.7 (4)	7.8	3.9	7.8	3.9	3.9	11.7 (4)	15.6	3.9	7.8	3.9	31.3	<b>125</b>	15.6	7.8	36.5 (13)	15.6
<i>E. faecalis</i> †	1.0	2.0	2.0	2.0	2.0	2.0	1.0	2.0	2.0	2.0	3.9	3.9	2.0	<b>125</b>	<b>31.3</b>	2.0	104.2 (32)	62.5
<i>P. aeruginosa</i> †	14.3 (2)	31.3	15.6	15.6	31.3	15.6	31.3	<b>125</b>	31.3	31.3	<b>125</b>	31.3	125	125	250	62.5	125	62.5
<i>M. osloensis</i> *	1.0	1.0	1.0	1.0	1.0	na	1.4 (0.5)	3.9	2.0	2.0	2	na	2.0	3.9	3.9	2.0	2.0	na
<i>A. baumannii</i> *	15.6	31.3	15.6	3.9	7.8	na	15.6	<b>62.5</b>	31.3	62.5	62.5	na	62.5	125	31.3	62.5	62.5	na
<i>P. putida</i> *	47.4 (17)	31.3	na	4.6(1)	3.9	na	62.5	41.7 (17)	na	31.3	62.5	na	62.5	62.5	na	62.5	62.5	na
<i>C. sakazakii</i> *	7.2 (2)	15.6	15.6	7.8	15.6	na	15.6	31.3	31.3	7.8	15.6	na	31.3	62.5	62.5	15.6	31.3	na
<i>E. coli</i> *	4.6 (2)	15.6	15.6	3.9	7.8	3.9	10.4 (4)	<b>41.7 (17)</b>	31.3	3.9	<b>15.6</b>	7.8	15.6	<b>62.5</b>	31.3	15.6	62.5	23.5 (9)

566 See footnote in Table 1

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571 **Table 5.** Bacterial susceptibility towards Glydant (DMDM-hydantoin) in planktonic and biofilm growth modes before, during and after repeated exposure to  
 572 Glydant (DMDM-hydantoin) in aqueous solution or in formulation.

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	187.5	187.5	na	187.5	187.5	na	375	482 (183)	na	375	375	na	3000	3000	na	1500	3000	na
<i>E. coli</i> †	375	375	na	375	375	375	1500	1500	na	375	750	375	6000	6000	na	1500	<b>6000</b>	1500
<i>E. faecalis</i> †	187.5	187.5	na	187.5	187.5	na	1500	1500	na	1500	750	na	3000	3000	na	3000	6000	na
<i>P. aeruginosa</i> †	187.5	187.5	na	187.5	187.5	na	6000	6000	na	1500	1500	na	6000	6000	na	6000	12000	na
<i>M. osloensis</i> *	375	375	na	46.9	62.5	na	325	375	na	187.5	187.5	na	750	1500	na	750	1500	na
<i>A. baumannii</i> *	375	325	na	187.5	187.5	na	750	750	na	375	375	na	6000	6000	na	6000	6000	na
<i>P. putida</i> *	375	375	na	375	375	na	750	750	na	750	375	na	6000	6000	na	3000	6000	na
<i>C. sakazakii</i> *	375	375	na	187.5	187.5	375	3000	3000	na	375	750	375	6000	6000	na	1500	<b>6000</b>	1500
<i>E. coli</i> *	187.5	466 (219)	187.5	187.5	375	187.5	375	<b>1500</b>	375	375	750	375	6000	6000	6000	1500	<b>12000</b>	1500

573 See footnote in Table 1

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577 **Table 6.** Bacterial susceptibility towards PHMB in planktonic and biofilm growth modes before, during and after repeated exposure to PHMB in aqueous  
 578 solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	3.9	<b>23.5 (9)</b>	<b>15.6</b>	3.9	3.9	3.9	3.9	<b>125</b>	<b>15.6</b>	3.9	<b>15.6</b>	7.8	15.6	<b>125</b>	15.6	15.6	<b>125</b>	31.3
<i>E. coli</i> †	15 (10)	31.3	15.6	7.8	15.6	na	15 (10)	<b>62.5</b>	<b>62.5</b>	15.6	31.3	na	62.5	62.5	62.5	62.5	31.3	na
<i>E. faecalis</i> †	7.8	<b>31.3</b>	15.6	5.9(1)	15.6	7.8	7.8	<b>125</b>	15.6	7.8	<b>31.3</b>	7.8	14.3 (3)	<b>125</b>	31.3	15.6	<b>125</b>	31.3
<i>P. aeruginosa</i> †	22.8 (15)	31.3	62.5	15.6	15.6	15.6	22.8 (15)	<b>125</b>	<b>125</b>	31.3	<b>125</b>	31.3	250	250	250	250	62.5	62.5
<i>M. osloensis</i> *	7.8	<b>31.3</b>	3.9	1.0	1.0	na	62.5	31.3	31.3	7.8	7.8	na	62.5	62.5	31.3	31.3	62.5	na
<i>A. baumannii</i> *	7.8	<b>31.3</b>	7.8	9.1 (3)	15.6	na	62.5	125	62.5	31.3	62.5	na	62.5	<b>250</b>	62.5	62.5	125	na
<i>P. putida</i> *	28.9 (8)	31.3	na	15.6	15.6	na	62.5	62.5	na	31.3	62.5	na	125	125	na	125	125	na
<i>C. sakazakii</i> *	7.8	15.6	15.6	31.2	15.6	na	104 (32)	125	125	15.6	31.3	na	62.5	<b>250</b>	125	62.5	125	na
<i>E. coli</i> *	7.8	7.8	31.3	7.8	15.6	na	15.6	<b>250</b>	31.3	15.6	31.3	na	62.5	<b>250</b>	31.3	62.5	31.3	na

579 See footnote in Table 1

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585 **Table 7.** Bacterial susceptibility towards thymol in planktonic and biofilm growth modes before, during and after repeated exposure to thymol in aqueous  
 586 solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	187.5	187.5	na	187.5	187.5	na	375	375	na	375	750	na	416 (160)	375	na	375	750	na
<i>E. coli</i> †	1500	1500	na	187.5	375	375	1500	1500	na	375	<b>1500</b>	750	1500	1500	na	375	<b>3000</b>	1500
<i>E. faecalis</i> †	375	750	na	187.5	375	na	750	750	na	375	750	na	750	750	na	750	1500	na
<i>P. aeruginosa</i> †	3000	3000	na	1500	3000	na	6000	3000	na	3000	6000	na	6000	6000	na	6000	12000	na
<i>M. osloensis</i> *	750	750	na	187.5	375	na	750	750	na	187.5	375	na	3000	1500	na	3000	375	na
<i>A. baumannii</i> *	750	750	na	375	375	375	1500	3000	na	750	<b>6000</b>	<b>3000</b>	6000	6000	na	6000	6000	6000
<i>P. putida</i> *	750	750	na	375	<b>3000</b>	375	1500	3000	na	1500	<b>6000</b>	3000	6000	6000	na	6000	6000	12000
<i>C. sakazakii</i> *	750	750	na	375	375	na	2250 (822)	3000	na	375	750	na	6000	6000	na	3000	750	na
<i>E. coli</i> *	665 (190)	750	na	187.5	375	na	3000	3000	na	375	750	na	6000	6000	na	750	<b>3000</b>	1500

587 See footnote in Table 1

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593 **Table 8.** Bacterial susceptibility towards triclosan in planktonic and biofilm growth modes before, during and after repeated exposure to triclosan in aqueous  
 594 solution or in formulation

Bacterium	MIC						MBC						MBEC					
	UF			F			UF			F			UF			F		
	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14	P0	P14	X14
<i>S. aureus</i> †	0.2	<b>62.5</b>	<b>31.3</b>	0.1	0.1	0.1	3.9	<b>62.5</b>	<b>62.5</b>	0.1	0.1	0.1	65.1	125	125	2.0	<b>7.8</b>	2.0
<i>E. coli</i> †	2.0	<b>62.5</b>	<b>62.5</b>	0.1	<b>2.0</b>	<b>3.9</b>	2.0	<b>125</b>	<b>125</b>	7.8	7.8	3.9	125	<b>500</b>	125	62.5	15.6	15.6
<i>E. faecalis</i> †	62.5	62.5	62.5	0.1	0.1	0.1	62.5	125	125	0.1	0.1	0.1	15.6	<b>125</b>	<b>125</b>	2.0	<b>7.8</b>	2.0
<i>P. aeruginosa</i> †	ns	ns	ns	7.8	<b>62.5</b>	7.8	ns	ns	ns	62.5	62.5	7.8	ns	ns	ns	62.5	62.5	7.8
<i>M. osloensis</i> *	1.0	<b>15.6</b>	<b>7.8</b>	1.0	1.0	na	7.8	<b>31.3</b>	<b>31.3</b>	3.9	3.9	na	125	125	125	3.9	3.9	na
<i>A. baumannii</i> *	125	125	125	2.0	2.0	na	125	250	125	31.6	15.6	na	125	250	125	62.5	15.6	na
<i>P. putida</i> *	15.6	<b>62.5</b>	<b>62.5</b>	1.0	2.0	na	62.5	125	125	15.6	15.6	na	125	250	500	62.5	15.6	na
<i>C. sakazakii</i> *	7.8	<b>500</b>	<b>188</b>	2.0	2.0	na	7.8	<b>1000</b>	<b>250</b>	31.3	31.3	na	1.3 (0.5)	<b>125</b>	<b>125</b>	62.5	31.3	na
<i>E. coli</i> *	1.0	<b>125</b>	<b>62.5</b>	0.1	<b>2.0</b>	<b>3.9</b>	2.0	<b>250</b>	<b>125</b>	15.6	15.6	15.6	125	<b>500</b>	125	62.5	15.6	15.6

595 See footnote in Table 1. ns, not susceptible (MBC/MIC/MBEC >1000 mg/L)

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