

1 Exploration of the Pharmacodynamics for *Pseudomonas aeruginosa* Biofilm Eradication by
2 Tobramycin

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13 Running Head: Biofilm Eradication

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23 **Abstract**

24 *Pseudomonas aeruginosa* is a Gram-negative, opportunistic pathogen which is involved in
25 numerous infections. It is of growing concern within the field of antibiotic resistant and tolerance
26 and often exhibits multi-drug resistance. Previous studies have shown the emergence of antibiotic
27 resistant and tolerant variants within the zone of clearance of a biofilm lawn after exposure to
28 aminoglycosides. As concerning as the tolerant variant emergence is, there was also a zone of
29 killing (ZOK) immediately surrounding the antibiotic source from which no detectable bacteria
30 emerged or were cultured. In this study, the ZOK was analyzed using both *in vitro* and *in silico*
31 methods to determine if there was a consistent antibiotic concentration versus time constraint (area
32 under the curve, (AUC)) which is able to completely kill all bacteria in the lawn biofilms in our *in*
33 *vitro* model. Our studies revealed that by achieving an average AUC of 4,372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$,
34 complete eradication of biofilms grown on both agar and hydroxyapatite was possible. These
35 findings show that appropriate antibiotic concentrations and treatment duration may be able to
36 treat antibiotic resistant and tolerant biofilm infections.

37

38 **Introduction**

39 *Pseudomonas aeruginosa* is a Gram-negative bacterium and opportunistic pathogen. Most
40 commonly, it is associated with cystic fibrosis (CF) related lung infections but is also implicated
41 in chronic wounds and post-surgical site infections (1-3). Antimicrobial tolerance and resistance
42 are also major concerns with *P. aeruginosa*, as the formation of biofilms, variant populations, and
43 multidrug resistance mechanisms are also prevalent (4-8). A recent study on the antibiotic
44 resistance rates of *P. aeruginosa* have shown a minimum 20% rate of resistance for carbapenems,
45 cephalosporins, aminoglycosides, and piperacillin/tazobactam (9). In addition, multi-drug
46 resistance was also found in 20% of infections (9).

47

48 In a previous study, variant colony phenotypes of *P. aeruginosa* emerging within the region
49 cleared by a tobramycin-loaded calcium sulfate bead were identified (8). This region of bacterial
50 lawn clearance is referred to as the zone of clearance (ZOC, Figure 1). These variant colonies
51 included classical resistance, persister cells, viable but non-culturable like colonies (VBNC-like),
52 and newly identified, tolerant, phoenix colonies (8). While the significance of these emergent
53 phenotypes may be of concern from a clinical standpoint, it is important to note that there was also
54 a smaller, consistent region within the ZOC, immediately adjacent to the antibiotic source, from
55 which no variants emerged or were cultured (8, 14). This previously reported zone, referred to as
56 the zone of killing (ZOK), represents a region of complete biofilm killing including antibiotic
57 tolerant and resistant variants (Figure 1, (8, 14)).

58

59 INSERT FIGURE 1 HERE

60

61 While pharmacodynamic studies have been done using planktonic *P. aeruginosa* exposed to
62 antibiotics (15, 16), further research into the pharmacodynamics necessary to eliminate biofilms is
63 needed. Currently, the primary method for measuring pharmacokinetics and pharmacodynamics
64 (PK/PD) *in vitro* is the modified Calgary biofilm device method (17). This method is used to grow
65 biofilms on pegs which are suspended in wells of a 96-well plate before exposing the biofilms to
66 antibiotics (17). Use of this method can be complicated by contamination risks, variation in
67 recovered biofilm after antibiotic exposure due to the rinsing steps necessary, and residual
68 antibiotics left after rinses (17). In this study, the ZOK from which no bacteria could be cultured
69 was further analyzed using a new method for exploring PK/PD *in vitro*. We hypothesized that the
70 junction between the outer edge of the ZOK and the region containing the emergent variants
71 represents an antibiotic concentration versus treatment time constraint in which complete bacterial
72 biofilm eradication can be achieved. In order to evaluate our hypothesis, a combination of a biofilm
73 plate *in vitro* model and an *in silico* approach was used to identify the concentration and time
74 (AUC) necessary to eliminate a *P. aeruginosa* biofilm, including variants which can typically
75 survive antibiotic therapy, using tobramycin. Additionally, substrates using both a tissue mimic
76 (agar, (18, 19)) and a bone mimic (hydroxyapatite (HA), (20)) were used to determine if the biofilm
77 substrate would affect the ability of an AUC to completely eradicate the biofilm.

78

79 **Results**

80 *Zones of Clearance of Biofilms Exhibit Dose-dependency*

81 In order to begin assessing the possible correlation between the ZOK and an antibiotic
82 concentration versus time constraint, twenty-four hour biofilm lawns of *P. aeruginosa* Xen41 were
83 generated as reported previously (8, 18) before being exposed to various weight quantities of

84 tobramycin (Figure 2). *In vitro* Imaging System (IVIS) evaluation of the biofilm lawns post
85 tobramycin exposure showed the emergence of dose-dependent ZOCs which continued to expand
86 over time. Within the ZOC, variant colonies began to emerge and encroach on the ZOC which had
87 formed. However, there was also a dose-dependent ZOK within the ZOC from which no
88 discernable bacterial activity was detected (Figure 2). It should also be noted that the luminescence
89 which appears to overly the antibiotic disks is likely due to surface associated biofilm growth. The
90 presence of the disk likely provides an additional substrate for biofilm growth and allows for initial
91 levels of high bioluminescence before the tobramycin ultimately is able to clear the formed
92 biofilm.

93

94 INSERT FIGURE 2 HERE

95

96 *Profiling of the ZOK Confirms Dose-dependency*

97 After exposure of the biofilm lawns to tobramycin, the plates were further analyzed to obtain
98 measurements of the ZOK over time. These measurements were used to generate plots (Figure 3)
99 which show the linear generation of the ZOC, followed by a ZOC peak right before variant
100 colonies began to emerge. The peak of the plot for each weight quantity of tobramycin shifts to a
101 later time point as the weight quantity increases. This is likely due to the time necessary for the
102 antibiotic to diffuse to low enough concentrations to allow variant colonies to begin to emerge. As
103 variant colonies began to emerge and grow within the ZOC, this allows the ZOK to be visualized.
104 The plots then plateaued as the ZOK became more apparent in which no detectable bacteria
105 emerged or grew. Once the ZOK stabilized, the radius of the edge of the ZOK was noted, as this

106 is the point likely to represent the minimum antibiotic concentration away from the antibiotic
107 source versus time constraint necessary for complete biofilm eradication including the killing of
108 any antibiotic tolerant or resistant variants. These radii values and time points were then used to
109 generate antibiotic diffusion plots using numerical modeling. In addition, a time-kill curve was
110 generated for the ZOK of a 1 mg tobramycin loaded disk. At 40 hours post tobramycin exposure,
111 a slight decrease in CFUs/cm² can be seen which further decreases and becomes significant
112 (p=0.017) by 48 hours. At 72 hours of tobramycin exposure and thereafter, no CFUs are able to be
113 recovered from within the ZOK (Figure 4).

114

115 INSERT FIGURE 3 HERE

116

117 INSERT FIGURE 4 HERE

118

119 *Computational Modeling Identified an Area Under the Curve Value for Biofilm Eradication*

120 Using the data collected from the profiling of the ZOC and ZOK, a model was used (21) to predict
121 the tobramycin concentration over time at the ZOK radius identified for each mass of tobramycin
122 used (Figure 5). The model calculates the concentration of tobramycin over time assuming Fickian
123 diffusion in a finite space (21, 22). The plots in Figure 3 show a dose-dependent increase in the
124 antibiotic concentration after antibiotic placement until concentrations for each weight quantity of
125 antibiotic peaked (120.9, 241.9, 483.8, and 967.5 $\mu\text{g}/\text{mL}$, respectively). The concentration then
126 gradually begins to decrease and plateau as the system continues to approach equilibrium (12.5,
127 25, 50, 100 mg/mL , respectively) for each weight quantity (250, 500, 1000, 2000 μg , respectively).
128 An AUC ($\mu\text{g}\cdot\text{hr}/\text{mL}$) was calculated for each plot to determine the necessary minimum value

129 necessary for ZOK generation for each of the amounts of tobramycin (Table 1). Interestingly,
130 despite a 4-fold change in the starting tobramycin weight quantity, the AUCs for each were
131 relatively similar. This observation highlights the importance of an AUC being taken into
132 consideration during antibiotic dosing as opposed to purely relying on the antibiotic concentration
133 alone. The mean of the AUC values was also calculated to determine that the average minimum
134 value necessary for biofilm eradication, including antibiotic tolerant and resistant variants, was
135 approximately 4,372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$. The concentration of antibiotic needed here is much higher than
136 standard minimum inhibitory concentrations (MIC) for tobramycin and *P. aeruginosa* PAO1 (4
137 $\mu\text{g}/\text{mL}$). Additionally, by dividing the AUC by the MIC, an AUC/MIC ratio of 46 can be calculated
138 when normalizing for 24 hours. This higher concentration ratio is likely responsible for the ability
139 to kill even antibiotic resistant and tolerant variants.

140

141 INSERT FIGURE 5 HERE

142

143 INSERT TABLE 1 HERE

144

145 *Identified AUC Can Eradicate Biofilms Grown on an Alternate Substrate*

146 In order to test the AUC value associated with complete *P. aeruginosa* biofilm eradication by
147 tobramycin, biofilms were grown on LB agar coated pegs, as well as HA coupons and plastic pegs
148 as alternate biofilm substrates. Biofilm colony forming units (CFUs) were measured for biofilms
149 exposed to tobramycin at the approximate 4,372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$ AUC. LB agar coated peg, HA
150 coupon, and plastic peg biofilms were exposed to tobramycin for 24 hours at a stable concentration
151 of 182 $\mu\text{g}/\text{mL}$ (Figure 6) which equals an approximate AUC value of 4372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$. In all

152 tobramycin exposure samples, no CFUs were recovered, indicating a complete eradication of the
153 biofilms. Additionally, a control was run at a 0.1 AUC equivalent on agar grown biofilms. The
154 control samples showed full lawns of growth after tobramycin treatment, indicating minimal
155 effectiveness of biofilm clearance. These results confirm that by obtaining a high concentration of
156 antibiotics for at least 24 hours, complete eradication of a *P. aeruginosa* lawn biofilm including
157 any antibiotic resistant or tolerant variants was possible and, substrate differences were not
158 relevant.

159

160 INSERT FIGURE 6 HERE

161

162 **Discussion**

163 *P. aeruginosa* is an opportunistic, bacterial pathogen which has been associated with numerous
164 infection sites. It has also been shown to readily develop tolerance and resistance to multiple
165 antibacterial drugs leading to an increased concern for complicated and difficult to treat infections
166 (23-25). Previously, the antibiotic tolerant phoenix and the VBNC-like phenotypes were identified,
167 in addition to classically resistant colonies and persister cells, within the ZOC of produced by
168 antibiotics released from tobramycin loaded Kirby-Bauer filter disks and CaSO₄ beads (8). While
169 the emergence of these variant colonies is concerning, it was also shown that there was a consistent
170 ZOK immediately adjacent to the antibiotic source from which no bacteria emerged or were
171 cultured. Replica plating was also performed on these plates and showed no growth within the
172 ZOK, indicating that this region is sterile (8). In these studies, both the ZOC and ZOK were
173 measured since it is not possible to differentiate between the two zones at early time points before
174 variant colonies begin to emerge. The distinct ZOK within the ZOC indicates a smaller region

175 where we may be able to eliminate antibiotic resistance and tolerance with high enough
176 concentrations of antibiotics over extended time frames. By understanding how antibiotic kinetics
177 effect biofilms and antibiotic resistant and tolerant variants, we will be better prepared to control
178 and prevent the rise of these dangerous biofilm infections.

179 In this study, the ZOK of *P. aeruginosa* biofilms was analyzed to determine the antibiotic
180 pharmacodynamics necessary for its generation. For our study, a combination of a biofilm plate
181 model and numerical modeling was used as a simpler alternative to the modified Calgary biofilm
182 plate model (17). The methods presented here could be adapted for further *in vitro* biofilm PK/PD
183 studies including the use of other bacterial species and antibiotics. Previous studies have shown
184 the importance of not only the concentration of antibiotic used but also the exposure time of the
185 biofilm-growing bacteria to the antibiotic. These studies on the antibiotic AUC have shown that
186 as exposure time increases, the minimum concentration needed for biofilm eradication (MBEC)
187 decreases (26). In addition, pharmacodynamic studies on the efficacy of antibiotics against bacteria
188 have shown an importance in the ratio of AUC/MIC (27, 28). The AUC/MIC ratio necessary for
189 bactericidal activity against *P. aeruginosa* by tobramycin has been shown to be approximately 42.
190 For our studies, various weight quantities of tobramycin were used which allowed for a more
191 complete understanding of these required pharmacodynamics. While the varying masses of
192 antibiotic used showed large, dose-dependent variations in the ZOK diameters, it was interesting
193 that the AUCs calculated for each mass did not scale similar to the weight of the antibiotic, ranging
194 from 3560-4870 $\mu\text{g}\cdot\text{hr}/\text{mL}$, although there was a nearly 8-fold increase in antibiotic dosing (250-
195 2000 μg). At an approximate AUC of 4372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$ and an MIC of 4 $\mu\text{g}/\text{mL}$, the AUC/MIC
196 ratio for biofilm eradication by tobramycin would be approximately 46 when normalized for a 24-
197 hour timeframe. Previous studies on local tobramycin concentrations achieved when treating

198 orthopedic infections with tobramycin loaded methylmethacrylate bone cement has shown that the
199 concentrations peak at approximately 40 $\mu\text{g}/\text{mL}$ around 1 hour after beginning exposure. The
200 concentration then drops to approximately 20 $\mu\text{g}/\text{mL}$ through 24 hours (30). These lower
201 concentrations may explain the prevalence of post-surgical infection treatment failure, as antibiotic
202 tolerant or resistant variants may be able to survive. In addition, while $C_{\text{max}}/\text{MIC}$ is the metric
203 typically used in relation to tobramycin (27), it has been shown that by using AUC/MIC, better
204 clinical outcomes are able to be achieved (28). This may be especially true when treating biofilm
205 infections as biofilms are inherently more tolerant of antibiotics due to slow diffusion of the
206 antibiotic into the biofilm matrix as well as heterogeneity in the biofilm metabolism (31),
207 increasing the importance of time-dependence. Our observations suggest that a minimum AUC
208 representing the total treatment effect (concentration x time) may be used to facilitate the complete
209 eradication of *P. aeruginosa* biofilms, including variants that without this treatment could re-seed
210 infection (32). It is also important to note that although VBNC-like colonies may be present in our
211 systems, they would not be apparent as they are unable to be cultured by nature. However, this
212 characteristic also would likely prevent re-emergence in a clinical environment.

213 In addition to using different weight quantities of tobramycin, different substrates were also used
214 to provide breadth to the relevance of the identified 4,372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$ necessary for biofilm
215 eradication. It is possible that biofilms grown on different substrates could have differing
216 phenotypes, including variations in metabolic pathways which may play a role in the efficacy of
217 antibiotics (33, 34). Although biofilm phenotypic variations can occur, the identified AUC was
218 able to eradicate biofilms grown on both a tissue mimic such as agar (18, 19) and a material
219 commonly known to be important for bone infections (HA, (20)), indicating that the use of the
220 AUC metric for biofilm treatment is independent of the substrate rigidity and providing relevance

221 to other fields such as dentistry and orthopedics. The observation evaluating the clearance of
222 biofilms from two distinct substrates is important clinically as biofilms are able to grow in the
223 human body on soft tissues such as lung tissue or skin and on hard substrates such as bone or
224 implanted devices (20, 35, 36). Clearly, optimizing the AUC as a treatment effect metric needs
225 further work, but the values reported here provide a specific evaluation for tobramycin and *P.*
226 *aeruginosa*. Therefore, further research, including studies involving other antibiotic classes and
227 bacterial species, is needed for a more complete understanding of the ability to eradicate bacterial
228 biofilm infections.

229

230 **Methods and Methods**

231 *Bacterial strains and culture conditions*

232 The metabolically driven, bioluminescent strain *P. aeruginosa* Xen41 (Xenogen Corp., USA) was
233 used for the imaging portion of this study. Additionally, *P. aeruginosa* PAO1 (15) which is both
234 a standard lab strain and the parent strain for *P. aeruginosa* Xen41 was used in this study. Culture
235 plates were prepared by from glycerol stock cultures stored at -80°C using 100 mm petri plates
236 (Fisher Scientific, USA) containing 20 mL of Luria-Bertani (LB) agar. Streaked plates were
237 incubated at 37°C for 24 hours to allow for individual colonies to grow. Individual colonies were
238 examined for proper morphology and then isolated and transferred to 20 mL of LB broth. Broth
239 cultures were incubated overnight in a shaking incubator set to 37°C and 200 rpm.

240

241 *Generation of Biofilm Lawns*

242 Overnight broth cultures of *P. aeruginosa* Xen41 were diluted to an OD₆₀₀ of 0.1 in LB broth and
243 100 µL aliquots were spread onto plates containing 20 mL of sterile, LB agar. These plates were
244 incubated at 37°C with 5% CO₂ for 24 hours to allow for biofilm lawns to generate. Lawn
245 generation was confirmed both visually and by using *in vitro* imaging system (IVIS) imaging.

246

247 *Bioluminescence Imaging*

248 Bioluminescence imaging was completed using thirty second exposures in an IVIS system. A
249 pseudo-color heatmap was overlaid to aid in visualization of the metabolic activity of the biofilms.
250 The scale for the heatmap is as follows: red indicates high levels of metabolic activity, blue
251 indicates low levels of metabolic activity, and black indicates no metabolic activity.

252

253 *Exposure of Biofilm Lawns to Antibiotic*

254 After lawn generation, the biofilms were exposed to set masses of tobramycin (250 µg, 500 µg,
255 1000 µg, and 2000 µg). Briefly, a 100 mg/mL stock solution of tobramycin was created using
256 tobramycin powder (TCI America, USA) and sterile water. This solution was then used to create
257 dilutions for the other necessary masses of tobramycin. A sterile, filter paper disk was placed in
258 the center of each of the lawn biofilms and then 20 µL of the appropriate tobramycin solution was
259 placed on the disk to allow for exposure of the biofilm lawns to the desired masses of tobramycin.
260 After antibiotic placement, the plates continued to be incubated at 37°C with 5% CO₂ for five
261 additional days. IVIS images were taken and the zones of killing measured daily.

262

263 *Time-kill Curve Generation*

264 In order to measure the CFUs within the ZOK over time, time-kill curves were generated.
265 Biofilm lawns were generated as above and were either exposed to 1 mg of tobramycin on a
266 filter paper disk or not exposed for controls. At various time points, a 1 cm x 1 cm area of
267 biofilm lawn, located between 1 and 2 cm from the disk, was scraped using a sterile loop and
268 placed into 1 mL of sterile PBS. A dilution series was then made and plated. CFUs were counted
269 for each sample and curves were generated.

270

271 *Computational Modeling*

272 In order to calculate the antibiotic concentration versus time necessary for development of the
273 ZOK, an analytical solution was used to estimate the tobramycin concentration, $c(r, t)$ for a 120
274 hour treatment time. The solution for the diffusion of antibiotic bead or disk in an agar plate is
275 given by:

276
$$c(r, t) = \frac{m}{h_0} \frac{1}{4\pi D(t+t_0)} \exp\left(-\frac{r^2}{4D(t+t_0)}\right),$$

277 where the diffusion coefficient of tobramycin in agar is D and was experimentally determined to
278 be $3.84 \times 10^{-10} \text{ m}^2/\text{s}$, m is the mass of antibiotic added to the source, h_0 is the height of the agar,
279 and r is the position coordinate relative to the center of a paper disk (19, 21, 22, 37). The model
280 assumes that antibiotic source is finite with a radius, $r_o = 3 \text{ mm}$. The model assumed that the inner
281 radius of the petri dish, 70 mm (inner diameter is 140 mm), was >10 times the radius of the paper
282 disks (3 mm) used, whereas the depth of agar layer was ~3.6 mm. After model generation, an area
283 under the curve (AUC) value was calculated in MATLAB for each curve using the trapezoidal rule
284 function, and the mean of the AUCs was also calculated.

285

286 *Testing of AUC on Hydroxyapatite*

287 In order to assess whether the AUC required for *P. aeruginosa* biofilm eradication could be
288 translated to biofilms grown on alternative surfaces, biofilms of *P. aeruginosa* PAO1 were grown
289 on 0.5 inch hydroxyapatite (HA) coupons (Biosurface Technologies, USA). Overnight LB broth
290 cultures of *P. aeruginosa* PAO1 were prepared as above. Overnight cultures were diluted to an
291 OD₆₀₀ of 0.1 in LB broth. HA coupons were added to the wells of a 12 well plate before 3 mL of
292 the diluted overnight culture were also added to the wells. The well plate was incubated at 37°C
293 with 5% CO₂ for 24 hours to allow biofilms to form on the HA coupons. Three coupons were then
294 exposed to tobramycin, and three coupons were exposed to PBS only as a control. Briefly, the
295 coupons were removed from the well plate and rinsed in PBS to remove any planktonic bacteria.
296 The coupons were then placed in a fresh 12 well plate for exposure to tobramycin or PBS only.
297 Three mL of a tobramycin solution was used for antibiotic exposure of the coupons for 24 hours
298 at a concentration of 182 µg/mL in order to equal the mean AUC value of 4,368 µg*hr/mL. During
299 the tobramycin exposure, the coupons continued to be incubated at 37°C with 5% CO₂. After 24
300 hours, coupons were removed from the well plate to obtain CFU counts. The coupons were rinsed
301 in PBS before being placed into wells of a fresh 12 well plate containing 3 mL of PBS. This plate
302 was then placed into a water bath sonicator and sonicated for 30 minutes. After sonication, the
303 PBS was removed from each well and centrifuged in order to pellet the bacteria. The pellet was
304 then resuspended in 200 µL of LB broth and a dilution series was created. The dilution series was
305 plated on 20 mL of LB agar and CFU counts were obtained.

306

307 *Testing of AUC on Plastic and Agar*

308 In addition to AUC validation on HA coupons, the AUC was also validated on plastic pegs of an
309 MBEC assay biofilm inoculator plate (Innovotech, Canada) as well as MBEC assay biofilm
310 inoculator plate pegs coated in LB agar. Briefly, LB agar coated pegs were prepared as follows.
311 200 μ L of molten LB agar was placed into the wells of a 96 well plate. The MBEC plate, which
312 had previously been cooled to -80°C , was then placed onto the 96 well plate to allow the pegs to
313 dip into the molten LB agar. After 3 seconds, the plate was removed and the agar was allowed to
314 finish cooling in a coating on the pegs. After LB agar coating, both LB agar coated pegs and
315 uncoated pegs were exposed to *P. aeruginosa* PAO1 to allow biofilms to form. An overnight
316 culture of PAO1 was diluted in LB broth at a ratio of 1:1000. 150 μ L of the diluted culture was
317 added to wells of 96 well plates. MBEC plates both coated in LB agar and uncoated were placed
318 onto the 96 well plates with diluted culture. These plates were then incubated at 37°C with 110
319 rpm shaking for 24 hours to allow biofilms to develop. After incubation, the pegs were rinsed in
320 200 μ L of PBS for 10 seconds. After rinsing, the MBEC plates were placed on 96 well plates
321 containing either 200 μ L of 182 $\mu\text{g}/\text{mL}$ tobramycin or 200 μ L of PBS as a control. These plates
322 were then incubated for an additional 24 hours at 37°C with 110 rpm shaking. After incubation,
323 the MBEC plates were rinsed in wells containing 200 μ L PBS and then placed on a fresh 96 well
324 plate with wells containing 200 μ L PBS. These plates were then sonicated in a water bath sonicator
325 for 30 minutes. After sonication, the bacteria containing PBS was removed and plated in a dilution
326 series. CFU counts were obtained for each sample.

327

328 *Statistical analysis*

329 All experiments were replicated in triplicate. ANOVA analysis was completed by GraphPad Prism
330 (v8.2.1) with a p-value of 0.05 being considered significant.

331

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335 Blood Institute (R01HL141941; SP).

336 Conflict of interest - None declared.

337

338 Figure Legends

339

340 Figure 1 – Emergence of ZOC and ZOK. IVIS images of a representative *P. aeruginosa* Xen41
341 plate showing ZOC emergence after exposure to tobramycin. A biofilm lawn was grown for 24
342 hours before being exposed to tobramycin on Day 0. Over time, a ZOC (yellow dotted circle) can
343 be seen as the antibiotic clears the background lawn. On Day 4, variant colonies begin to grow
344 within the ZOC and a ZOK (red dotted circle) becomes apparent. The ZOK is the region in which
345 there are no variant colonies or other discernable bacterial activity. In the images, red indicates
346 high levels of metabolic activity, blue indicates low levels of metabolic activity, and black
347 indicates no metabolic activity.

348

349 Figure 2 – IVIS Imaging of *P. aeruginosa* Biofilms Exposed to Tobramycin. Biofilm lawns of *P.*
350 *aeruginosa* Xen41 were generated on agar before being exposed to various amounts of tobramycin
351 for a total of five days. Zones of clearance (ZOC) can be seen growing over time for each mass of
352 tobramycin. Although variant colonies emerge from the ZOC, there is a distinct region within the
353 ZOC, marked explicitly in Figure 1, known as the zone of killing (ZOK). In the images, red
354 indicates high levels of metabolic activity, blue indicates low levels of metabolic activity, and
355 black indicates no metabolic activity.

356

357 Figure 3 – ZOK Expansion Curves for Tobramycin. The ZOK of the plates from Figure 2 were
358 measured over time. The zones follow linearly until the peak (indicated by black arrows), which
359 occurs as variant colonies begin to emerge within the ZOC, and the diameter of the ZOK becomes
360 evident. As these variant colonies continue to emerge and begin to grow, the ZOK reduces. After
361 the peak, the ZOK continues to become more apparent as the area lacking emergence of variant

362 colonies. The ZOK decrease continues as variant colonies emerge until the plot begins to plateau,
363 at which point the ZOK was stable. Data are reported as mean \pm SD (n=3). *p<0.05, **p<0.01.

364

365 Figure 4 – Time-kill Curve for the ZOK. CFUs within the ZOK of 1 mg of tobramycin were
366 measured over time. Controls CFUs were measured for plates without tobramycin exposure. At 48
367 hours post tobramycin exposure, a decrease in CFUs begins to be seen, and by 72 hours post
368 tobramycin exposure, no CFUs were able to be recovered from within the ZOK. Data are reported
369 as mean \pm SD. *p<0.05, ***p<0.001, ****p<0.0001, n=3.

370

371 Figure 5 – Computational Modeling of Antibiotic Diffusion. The diffusion model shows the spread
372 of tobramycin over time. The radii used to determine the zones of killing for each antibiotic
373 quantity was experimentally measured (Figure 2) and used as an input to the model for the
374 computation of the concentration. All plots show an increase in tobramycin concentration followed
375 by a decrease following the Fickian diffusion model.

376

377 Table 1 – AUC Values Associated with Complete Biofilm Eradication in Our Agar Plate Biofilm
378 Model. Values for the AUC ($\mu\text{g}\cdot\text{hr}/\text{mL}$) for each mass of tobramycin were calculated from the
379 plots of Figure 4. The mean of each of these values was also calculated and likely represents the
380 minimum AUC necessary for complete eradication of biofilms, including antibiotic resistant and
381 tolerant variants.

	250 μg	500 μg	1000 μg	2000 μg	Average
AUC	3560 $\mu\text{g}\cdot\text{hr}/\text{mL}$	4730 $\mu\text{g}\cdot\text{hr}/\text{mL}$	4330 $\mu\text{g}\cdot\text{hr}/\text{mL}$	4870 $\mu\text{g}\cdot\text{hr}/\text{mL}$	4372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$

382

383

384 Figure 6 – Eradication of *P. aeruginosa* Biofilms Grown on Plastic, LB Agar, and Hydroxyapatite.
385 *P. aeruginosa* PAO1 biofilms grown on plastic pegs, LB agar coated pegs, and hydroxyapatite
386 coupons were exposed to 182 $\mu\text{g}/\text{mL}$ of tobramycin for 24 hours, equaling an AUC value of
387 approximately 4372.5 $\mu\text{g}\cdot\text{hr}/\text{mL}$. Complete biofilm eradication was seen in all of the tobramycin
388 treated samples. **** $p < 0.0001$, $n = 3$.

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