1	Predatory Bacteria can Reduce Pseudomonas aeruginosa Induced Corneal
2	Perforation and Proliferation in a Rabbit Keratitis Model
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4	Eric G. Romanowski ¹ , Nicholas A. Stella ¹ , Brvn L. Brazile ² , Kira L. Lathrop ^{2,3} , Jonathan
5	M Eranke ³ Ian A Sigal ^{2,4} Tami Kim ⁵ Monnat Elsaved ⁵ Daniel E Kadouri ⁵ and Robert
J C	M.O. Charles 12.*
6	M.Q. Shanks
7	Short title. Predatory bacteria versus P. aeruginosa in the cornea
8	¹ The Charles T. Campbell Laboratory, Department of Ophthalmology, University of
9	Pittsburgh School of Medicine, Pittsburgh, PA
10	² Department of Ophthalmology, University of Pittsburgh School of Modicine, Pittsburgh
10	
11	PA
12	³ Center for Biological Imaging, University of Pittsburgh School of Medicine, Pittsburgh,
13	PA
14	[*] Department of Bioengineering, Swanson School of Medicine, University of Pittsburgh,
15	Pittsburgh PA
16	⁵ Department of Oral Biology, Rutgers School of Dental Medicine, Newark, NJ
17	* Corresponding author: Robert M. Q. Shanks, 203 Lothrop St. Pittsburgh, PA 15213,
18	Ph: 1-412-647-3537, Fax: Ph: 1-412-647-5880, email: shanksrm@upmc.edu
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20	Key Words: BALOs, Bdellovibrio bacteriovorus, corneal perforation, inflammation,
21	keratitis, predatory bacteria, Pseudomonas aeruginosa
22	

23 Abstract

24	Purpose: Pseudomonas aeruginosa keratitis is a severe ocular infection that can lead
25	to perforation of the cornea. In this study we evaluated the role of bacterial quorum
26	sensing in generating corneal perforation and bacterial proliferation and tested whether
27	co-injection of the predatory bacteria Bdellovibrio bacteriovorus could alter the clinical
28	outcome. P. aeruginosa with lasR mutations were observed among keratitis isolates
29	from a study collecting samples from India, so an isogenic lasR mutant strain of P.
30	aeruginosa was included.
31	Methods: Rabbit corneas were intracorneally infected with P. aeruginosa strain PA14 or
32	an isogenic $\Delta lasR$ mutant and co-injected with PBS or <i>B. bacteriovorus</i> . After 24 h, eyes
33	were evaluated for clinical signs of infection. Samples were analyzed by scanning
34	electron microscopy, optical coherence tomography, sectioned for histology, and
35	corneas were homogenized for CFU enumeration and for inflammatory cytokines.
36	Results: We observed that 54% of corneas infected by wild-type PA14 presented with a
37	corneal perforation (n=24), whereas only 4% of PA14 infected corneas that were co-
38	infected with B. bacteriovorus perforate (n=25). Wild-type P. aeruginosa proliferation was
39	reduced 7-fold in the predatory bacteria treated eyes. The $\Delta lasR$ mutant was less able to
40	proliferate compared to the wild-type, but was largely unaffected by <i>B. bacteriovorus</i> .
41	Conclusion: These studies indicate a role for bacterial quorum sensing in the ability
42	of <i>P. aeruginosa</i> to proliferate and cause perforation of the rabbit cornea. Additionally,
43	this study suggests that predatory bacteria can reduce the virulence of P. aeruginosa in

44 an ocular prophylaxis model.

45

46 **1. Introduction**

47 Predatory bacteria, such as *Bdellovibrio bacteriovorus*, have been suggested as an 48 alternative approach for treatment antibiotic resistant bacterial infections in general [1-6]. 49 B. bacteriovorus is a Gram-negative bacterium that is an obligate predator of other 50 Gram-negative bacteria. Given the preponderance of Gram-negative bacteria in causing 51 contact-lens related keratitis, predatory bacteria have been hypothesized as treatments 52 for eye infections [7-9], and are largely non-toxic and non-inflammatory when applied in 53 great numbers to a corneal epithelial cell line and intact and wounded ocular surfaces [7, 54 10]. These have been tested to prevent bacterial proliferation on the ocular surface with 55 some success and to prevent the development of keratitis in a mouse model using 56 Escherichia coli as a pathogen [11, 12].

57 Unlike E. coli, Pseudomonas aeruginosa is the leading cause of contact lens 58 associated keratitis, a blinding infection [13-15]. Antibiotic resistant P. aeruginosa 59 keratitis isolates are associated with worse clinical outcomes [16-19]. Previous work 60 showed that *B. bacteriovorus* predation is not influenced by antibiotic resistance status 61 of its prey [3, 5, 7] and is even effective at clearing antibiotic tolerant biofilms [20, 21]. A 62 recent study showed that around 20% of *P. aeruginosa* keratitis isolates isolated in India 63 from 2006-2010 had mutations in the *lasR* gene and that these strains were correlated 64 with worse visual outcomes in patients [22]. The LasR protein is a major regulator of 65 quorum sensing and important in regulation of numerous virulence-associated genes 66 [23].

In this study we evaluated whether *B. bacteriovorus* strain HD100 could act as a
"living antibiotic" in a rabbit keratitis infection model using a clade 2 (cytotoxic/ExoU⁺)
and highly virulent *P. aeruginosa* strain. This was a proof-of-concept prevention model
rather than a treatment model. The study found that the predatory bacteria caused mild-

71	moderate inflammation, but were able to reduce P. aeruginosa proliferation and most
72	importantly significantly reduced the frequency of corneal perforation events caused by
73	<i>P. aeruginosa</i> . We also observed that an isogenic <i>P. aeruginosa</i> $\Delta lasR$ mutant was
74	unable to replicate in the rabbit cornea and caused significantly fewer corneal
75	perforations, but was not significantly reduced by <i>B. bacteriovorus</i> in the cornea.
76	
77	2. Methods
78	2.1. Bacterial strains and culture
79	B. bacteriovorus HD100 (ATCC 15356) [24] was used for this study. Preparation of B.
80	bacteriovorus followed prior reports [10, 12] where B. bacteriovorus was incubated with
81	~10 ⁹ CFU/mI <i>E. coli</i> diaminopimelic acid auxotroph strain WM3064 for 24 h at 30°C. The
82	resulting mixture was passed multiple times through a 0.45- μm Millex $^{\! \mathrm{\tiny B}}\text{-}HV$ pore-size
83	filter (Millipore, Billerica, MA, USA) to remove remaining prey. Predators were washed
84	with phosphate buffered saline (PBS) and concentrated centrifugation. B. bacteriovorus
85	was suspended in PBS to 1.5×10^{10} PFU/ml <i>B. bacteriovorus</i> .
86	For a pathogen, the cytotoxic (ExoU+) / clade II strain UCBPP-PA14 (PA14) [25-
87	28] and an isogenic $\Delta lasR$ mutant were used [29]. These were inoculated from single
88	colonies into LB medium and grown with aeration at 30°C or 37°C. For inoculation, P.
89	aeruginosa were diluted in phosphate buffered saline (PBS) to achieve the inoculum
90	(~2-4x10 ³ CFU in 10 μl).
91	
92	2.2. Bacterial keratitis studies
93	The experiments in this study conformed to the ARVO Statement on the Use of Animals
94	in Ophthalmic and Vision Research and were approved by the University of Pittsburgh's
95	Institutional Animal Care and Use Committee (IACUC Protocols 15025331 and

96 18022194).

97 Female New Zealand White rabbits (1.1-1.4 kg) were obtained from the 98 Oakwood Research Facility through Charles River Laboratories. The rabbits were 99 anesthetized with 40 mg/kg of ketamine and 4 mg/kg of xylazine administered 100 intramuscularly. The corneas of the right eyes only were anesthetized with topical 0.5% 101 proparacaine and injected intrastromally with the 10 µl of P. aeruginosa (~2000 CFU) in 102 PBS or PBS alone as a control. Actual inocula for each trial was determined using the 103 EddyJet 2 spiral plating system (Neutec Group Inc., Farmingdale, NY) on 5% trypticase 104 soy agar with 5% sheep's blood plates. Plates were incubated for ~18-20 h at 37°C and 105 the colonies were enumerated (Flash and Grow colony counting system, Neutec Group, 106 Inc). This was followed by a 25 μ I injection of *B. bacteriovorus* in PBS (~4x10⁸) or PBS. 107 At 24 h post-injection, the eyes were evaluated for ocular signs of inflammation 108 using a slit-lamp according to a modified McDonald-Shadduck grading system [30]. 109 Rabbits were systemically anesthetized with ketamine and xylazine as described above 110 and euthanized with Euthasol solution following the 2020 AVMA Euthanasia Guidelines. 111 Corneal buttons were harvested using a 10 mm trephine and placed into Lysing 112 Matrix A tubes (MP Biomedicals) containing 1 ml of PBS. The corneas were then 113 homogenized with an MP Fast Prep-24 homogenizer (MP Biomedicals), and the 114 numbers of corneal bacteria were enumerated as described above. The centrifuged and 115 filtered homogenate (Millipore 0.22 μ m PVDF filter) was frozen for ELISA analysis (IL-1 β 116 kit (Sigma-Aldrich RAB1108). MMP9 ELISA) following the manufacturer's guidelines. 117 Additional corneas were fixed with paraformaldehyde (4%) in PBS, embedded in 118 paraffin, and 5 µm sections were stained with hematoxylin and eosin. Sections were 119 viewed using an Olympus Provis AX-70 microscope and images captured with 120 MagnaFire 2.1 software.

121 In another set of animals, the whole eyes were removed following euthanasia
122 and stored individually in 25 mm of PBS and kept on ice. Within one hour, their corneas

123 were scanned by optical coherence tomography (OCT) using a Bioptigen Envisu R2210 124 system with InVivoVue software (Leica Microsystems) following the same general 125 approach described elsewhere [31]. Briefly, the system was modified with a broadband 126 superluminescent diode (Superlum, Dublin, Ireland $\lambda = 870$ nm, $\Delta\lambda = 200$ nm, 20,000 A-127 scans/second). OCT volume scans of the central cornea were acquired using a 10mm 128 telecentric lens for anterior segment imaging and a scan pattern: 6mm x 6mm x 2mm 129 (1000 x 100 x 1024 pixel sampling), with 1 Frame per B-scans, i.e. no repetitions. At 130 least three scans were obtained of each sample and the best quality scan chosen for 131 analysis. The most common issue was a high signal at the cornea apex that reduced 132 visibility directly underneath. Scans of a given eye were completed in less than six 133 minutes. The B-scans were loaded into Fiji for visualization as image volumes and to 134 export stills and movies [32].

135

136 2.3. Scanning Electron Microscopy

137 Electron microscopy was performed as previously described [12]. Rabbit corneal buttons

138 were fixed with glutaraldehyde (3%) at room temperature for 24h and washed with PBS.

139 Samples were pinned to prevent curling and post-fixed using osmium tetroxide (1%).

140 dehydrated with ethanol (30–100%), immersed in hexamethyldisilazane and air-dried.

141 Samples were mounted to aluminum stubs and sputter coated with gold/palladium (6

142 nm). Scanning electron microscopy was performed using a JEOL JSM-6335F scanning 143 electron microscope (3 kV).

144

145 2.4. In vitro predation assay

146 P. aeruginosa were grown at 37°C in LB broth on a tissue culture rotor, centrifuged and

147 suspended in HEPES buffer (25 mM HEPES with 3 mM MgCl₂ and 2 mM CaCl₂).

Aliquots of P. aeruginosa (0.5 ml, ~1x10⁹ CFU) and B. bacteriovorus (0.5 ml, ~2x10⁸ 148

- 149 PFU) and 4 ml of HEPES buffer were added to 15 ml conical polypropylene tubes and
- 150 incubated at 30°C. At 24 h bacteria were serial diluted and plated on LB agar plates to
- 151 determine surviving *P. aeruginosa*. Log reductions were calculated as compared to a
- 152 predator free control survival at the same time point.
- 153

154 2.5. Statistical analysis

- 155 Kruskal-Wallis with Dunn's post-test was used for non-parametric analysis and ANOVA
- 156 with Tukey's post-test or Student's T-tests for parametric analysis. Contingency analysis
- 157 was performed with Chi-Square test followed by Fisher's Exact test for pair-wise
- 158 comparisons. All analyses were done with GraphPad Prism software.
- 159

160 3. Results

- 161 3.1. Evaluation of the antimicrobial efficacy of *B. bacteriovorus* in a *P. aeruginosa* 162 keratitis model.
- 163 The ability of the predatory bacteria *B. bacteriovorus* strain HD100 to prevent *P.*
- 164 aeruginosa proliferation in a bacterial keratitis model was evaluated. P. aeruginosa was
- 165 used because it is a major cause of corneal infection. In this study either 10 µl of P.
- 166 aeruginosa (~2000 CFU) or the PBS vehicle was injected into the corneal stroma (Figure
- 167 1A), followed by a second injection of 25 µl of either *B. bacteriovorus* strain HD100
- 168 (4x10⁸ PFU) or the PBS vehicle into the same needle track.

169 The virulent P. aeruginosa strain UCBPP-PA14 (PA14) [25-28] was injected into 170 the cornea at an average of 1,777±353 CFU/cornea. This strain is a clade 2 strain, also 171 known as a cytotoxic strain, that is ExoU positive and ExoS negative [28]. This strain has 172 not been previously reported in a rabbit keratitis model to our knowledge. PA14 was able 173 to replicate 40-fold to 7.25x10⁴ CFU/cornea by 24 hours (Figure 1B). The eyes were 174

evaluated with a slit lamp and fluorescein staining using a MacDonald-Shadduck-based

scoring system [30] and there was a clear increase in corneal inflammation score by 24
h (Figure 1C-D, 2). This led to corneal perforation in just over half of the infected corneas
(54%) (Figure 1E, 2).

By contrast, *B. bacteriovorus* caused intermediate inflammation compared to PBS and PA14 groups (Figure 1C-E, 2). Moreover, *B. bacteriovorus* injection did not cause corneal perforations (Figure 1D, 2).

B. bacteriovorus injected following *P. aeruginosa* correlated with a significant reduction in *P. aeruginosa* proliferation $(1.03 \times 10^4 \text{ CFU} \text{ at } 24\text{h}, \text{p} < 0.01)$ but was still higher than the inoculum. This reduction did not significantly influence the overall corneal score, but was associated with a remarkable reduction in the frequency of corneal perforations (4%, p<0.001) (Figure 1D).

186 Fluorescein staining and slit lamp examination 24 h post-inoculation revealed 187 minor pathology of the corneas infected with B. bacteriovorus that consisted of redness 188 and a small infiltrate that did not cause a fluorescein stained ulcer (Fig 2). By contrast, 189 the *P. aeruginosa* injected eyes showed major purulent discharge from the large central 190 corneal ulcer and anterior chamber effects (Figure 2). Eyes injected with both had an 191 intermediate phenotype similar to those injected with both the $\Delta lasR P$. aeruginosa and 192 predatory bacteria. The eyes injected with just the $\Delta lasR$ mutant bacteria had minimal 193 inflammation, though had a corneal ulcer of moderate size (Figure 2).

Optical coherence tomography (OCT) was done on a subset of eyes following sacrifice of the rabbits at 24h post-inject (Figure 3). This reveals the shape of the cornea and unlike the PBS only injected eye, the *B. bacteriovorus* (HD100) injected eyes had clear inflammation including an infiltrate and overall edema. The loss of the corneal integrity was demonstrated in the *P. aeruginosa* PA14 wild-type injected eyes where the massive infiltrate prevented unobstructed corneal imaging and resulted in shadows. By contrast, the *P. aeruginosa* wild-type and HD100 coinjected eyes had clear damage,

201 ulceration, and infiltrate, but maintained overall shape. The $\Delta lasR$ mutant with or without 202 HD100 had intermediate phenotypes. Movies depicting OCT volumes of corneal layers 203 are provided in the supplemental information and clearly demonstrate the perforation of 204 the *P. aeruginosa* strain PA14 infected eye (Supplemental movies S1-3). 205 To evaluate whether infiltrates were made by neutrophils, H&E staining of 206 corneas was performed. The histology provided consistent results with the fluorescein 207 staining with respect to maintenance or loss of corneal epithelium (Figure 4). Neutrophils 208 and severe damage of the anterior stroma were observed in corneas infected by PA14 209 and to a lesser extent by *B. bacteriovorus* (Figure 4 and Figure S1). 210 Low magnification scanning electron microscopy showed the perforation in the 211 PA14 infected cornea that was absent in the other corneas (Figure 5). Similar to 212 fluorescein results (Figure 2), a clear ulcer was evident in corneas infected with PA14 213 (PA14 + PBS) and *B. bacteriovorus* (PBS + HD100) (Figure 5). At higher power, red and 214 white blood cells were observed on the ocular surface of most eyes, especially those 215 infected with PA14 (Figure 5). 216 Consistent with the corneal inflammation MMP9 (92-kDa neutrophil gelatinase or 217 gelatinase B), an enzyme that digests gelatin and collagen types IV and V [33], and the 218 pro-inflammatory cytokine interleukin-1 β (IL-1 β) were highly induced in corneas infected 219 with PA14, and to a lesser extend by *B. bacteriovorus* (Figure 6). The co-infection of 220 PA14 and *B. bacteriovorus* did not significantly alter these pro-inflammatory markers. 221 However, *B. bacteriovorus* by itself was less inflammatory than PA14 for both MMP9 and 222 IL-1β (p<0.05). 223

225 A recent study showed natural *lasR* mutants at high frequency among bacterial keratitis

3.2. LasR promotes corneal pathogenesis by a cytotoxic *P. aeruginosa* strain.

224

isolates [22]. Therefore, the above experiments were also carried out with an isogenic

227 variant of PA14 with a deletion of the *lasR* guorum sensing regulator. This strain was 228 previously validated by restoring the wild-type allele and is defective in expression of a 229 number of potential virulence associated phenotypes such as protease production [22]. 230 Though slightly more $\Delta lasR$ mutant bacteria were inoculated compared to the WT 231 $(3,805 \pm 1703 \text{ CFU/cornea}, p=0.03)$, the CFU per cornea at 24 h was significantly lower 232 at 5.9×10^3 CFU/cornea which is 12-fold lower than wild-type PA14, p<0.01 (Figure 1B). 233 Corneal inflammation scores were also lower, but did not reach significance (p>0.05), 234 (Figure 1C). Nevertheless, the $\Delta lasR$ mutant was defective in causing corneal 235 perforation compared to the WT (Figure 1D, p=0.002). Histology and SEM analysis 236 demonstrate reduced corneal damage (Figure 2-4), yet the MMP9 and IL-1ß levels were 237 similar to the WT (Figure 6). Both the wild-type PA14 and $\Delta lasR$ mutant was susceptible 238 to *B. bacteriovorus in vitro*. The mean Log₁₀ reduction *in vitro* was 4.9±0.05 CFU/ml for 239 PA14 and 4.7±0.15 CFU/ml for *AlasR*, p>0.05, n=3. However, the predatory bacteria did 240 not significantly lower the corneal burden. While corneal scores were high, the frequency 241 of perforation was low (0%). However, MMP9 and IL-1 β were levels in $\Delta lasR$ and B. 242 bacteriovorus inoculate corneas was indistinguishable from PA14 wild-type infected 243 eyes. 244 245 4. Discussion 246 This study demonstrated the severe pathogenesis caused by strain PA14 in a rabbit 247 corneal infection model. 54% of corneas perforated by 24 h limiting the length of the

study for ethical reasons. While *B. bacteriovorus* on their own caused some

inflammation, they prevented *P. aeruginosa* proliferation and corneal perforation.

In this study, MMP9 was strongly induced by *P. aeruginosa*. MMP9 is a

251 metalloproteinase secreted by neutrophils, macrophages, and corneal epithelial cells

[33]. Transcription of the MMP9 gene was strongly induced when corneal cells are

253 exposed to bacteria [34, 35]. MMP9 is involved in remodeling of the corneal basement 254 membrane and regulates corneal healing, but has also been linked with corneal 255 perforation [36, 37]. Tetracyclines inhibit MMP9 activity, and that is one reason that 256 tetracyclines have been given systemically to patients with corneal ulcers or perforations 257 [38]. In animal models tetracyclines promoted wound healing and prevented corneal 258 perforation [39-41]. Our results suggest that MMP9 may contribute to, but is not fully 259 responsible for corneal perforation caused by P. aeruginosa as MMP9 levels remained 260 in infected corneas that did not perforate. Other factors such as neutrophil elastase. 261 MMP8, or bacterial proteases such as elastase B, which is positively regulated by LasR 262 may be key contributors to corneal perforation in our model [42, 43]. 263 IL-1 is a major proinflammatory cytokine during bacterial keratitis [44] and 264 promotes neutrophil mediated corneal damage in mouse models of P. aeruginosa 265 keratitis [45, 46]. Although a role for IL-1 in preventing corneal perforation in young 266 Swiss mice has also been described [45]. An in vitro study showed that keratocytes 267 produced collagenolytic compounds in an IL-1 dependent manner and that collagen 268 degradation could be prevented with an IL-1 inhibitor [47]. IL-1 release from damaged 269 corneal epithelial cells acts as an alarmone and induces increased expression of IL-6 270 and -8 and reduced barrier function in corneal epithelial cells [48]. Similarly, anti-IL-1 β 271 antibody pretreatment reduced MMP9 induction in a mouse (C57Black/6) P. aeruginosa 272 keratitis model leading to the overall model that *P. aeruginosa* induced IL-1 induces the 273 expression of MMPs resulting in the destruction of corneal tissue [44].

In our study, predatory bacteria failed to induce IL-1β production, while this is an
unusual for a Gram-negative bacteria in the corneal stroma, *B. bacteriovorus* have
unusual physiology that may reduce its ability to activate the immune system. Namely, it
has a modified lipopolysaccharide structure that does not activate TLR4 and a
membrane sheathed flagellum, so is unlikely to activate TLR5-mediated inflammation

279 [49, 50]. By contrast, all other infection groups strongly induced IL-1^β that correlated with 280 high levels of MMP9. The intermediate levels of MMP9 induced by *B. bacteriovorus* 281 suggest that there are IL-1-independent mechanisms to induce MMP9. However, only 282 one time point was evaluated, so IL-1 may have been induced at early time point by B. 283 bacteriovorus. That said, IL-1 induction by bacterial keratitis was reported to be 284 maintained over the course of several days in a mouse model weakening the likelihood 285 that IL-1 levels were higher at earlier time points [45]. Although the levels of MMP9 286 measured in the predator group was significantly lower than that measured in the P. 287 aeruginosa infected arm, one should consider that the concentration of the predator 288 injected was 5 logs higher (\sim 4x10⁸ vs 2x10³ respectively). The low inflammatory 289 stimulating effect of B. bacteriovorus seen in this study is in agreement with both in vitro 290 [7, 51, 52] and *in vivo* studies that show that pro inflammatory cytokines, including IL-1, 291 are not significantly elevated after exposure to the predator [10, 53-56]. 292 The role of the LasR quorum sensing marker in *P. aeruginosa* keratitis has 293 previously been characterized in mouse models using a clade l/invasive/ExoS+ strain 294 PA01 and similar, but not isogenic lasR mutant [57]. Prior work has evaluated the 295 importance of LasR in *P. aeruginosa* keratitis. These studies both used PAO1, an ExoS+ 296 / invasive / clade I strain and a randomly mutated variant of PAO1 that was selected for 297 streptomycin resistance, PAO1s with the *lasR* gene replaced by a tetracycline resistance 298 cassette. The first study demonstrated that in a murine scratch model, 33-fewer lasR 299 mutant bacteria were necessary to establish infections in C3H/HeN mice [58]. The 300 second study used the same bacterial strains with a murine scratch keratitis model using 301 BALB/c mice and did not find a difference in the ability of the two bacteria to cause 302 keratitis, however the lasR mutant was not as successful in proliferation with ~10-fold 303 lower CFU/cornea being measured [59]. Our study, by contrast, used a ExoU+ / 304 cytotoxic / clade II strain in a rabbit intrastromal injection model. While both the scratch

305 and intrastromal injection models have important limitations, the intrastromal injection 306 model ensures highly similar inoculation numbers, which are important for a treatment 307 study, especially when the lasR mutant and wild-type express adhesins at different 308 levels [22]. The results of this study suggest that once past the epithelial barrier the lasR 309 mutant is less able to proliferate and this may account for the highly reduced frequency 310 of perforations, additionally, the lower expression of collagen digesting proteases such 311 as elastase B by the lasR mutant may also contribute to the highly reduced frequency of 312 corneal perforations [57, 59]. More work is needed to understand the complex role of 313 LasR regulated quorum sensing in *P. aeruginosa* keratitis, nevertheless, this study 314 supports the model that in at least one cytotoxic strain, LasR is required for full levels of 315 pathogenesis. 316

318 proliferation and prevent corneal perforation in a keratitis model, however, further study 319 to test efficacy against established infections by multiple types of *P. aeruginosa* clinical 320 isolates are necessary to move the concept of using predatory bacteria as a therapeutic 321 forward.

This proof-of-concept study supports that predatory bacteria can prevent bacterial

322

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338 6. Disclosure/Conflict of Interest Statement

- 339 The authors have no commercial interests regarding this study.
- 340

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508 8. Figure Legends

509 Figure 1. Predatory bacteria and *lasR* mutation reduce *P. aeruginosa* pathogenesis in

- 510 **a rabbit keratitis model. A.** Diagram of model in which the rabbit corneal stroma (arrow) is
- 511 injected with *P. aeruginosa* strain PA14, isogenic Δ*lasR* mutant, or PBS followed by PBS or
- 512 *B. bacteriovorus* strain HD100. **B-D**. Each point represents a rabbit. **a** indicates significant
- 513 difference from control group, p<0.05. n.s. indicates not significantly different than the control
- group. **, p<0.01; ***, p<0.001.**B**. CFU from corneas at 24 h post-injection. Means and SDs

515	are shown. C. Corneal inflammation score based on MacDonnald-Shadduck scoring
516	system. Medians and interquartile ranges are shown and Kruskal-Wallis analysis with
517	Dunn's post-test was performed. D. Corneal perforation frequency. Indicated groups were
518	compared by Fisher's Exact test.
519	
520	Figure 2. Bright-field microscopy and fluorescein staining of infected eyes.
521	Representative images are shown. Fluorescein staining indicates corneal ulceration and
522	loss of corneal epithelium.
523	
524	Figure 3. Optical coherence tomography of infected eyes. Representative images are
525	shown. Shadows are due to infiltrates that prevent imaging.
526	
527	Figure 4. Histological analysis demonstrates neutrophil infiltrates. Representative
528	images are shown. Magnification was 10X. Polymorphonuclear neutrophils are evident from
529	inside the stroma and in at the anterior chamber interface.
530	
531	Figure 5. Scanning micrographs of rabbit corneal surfaces. Representative images are
532	shown. Magnification was 25x (top) and $\epsilon 500x$ (bottom). The corneal perforation is clearly
533	shown in the PA14 + PBS injected cornea, and a corneal ulcer was evident in the PA14 with
534	HD100 injected cornea.
535	
536	Figure 6. <i>P. aeruginosa</i> strain PA14 stimulation of gelatinase B (MMP9) and pro-
537	inflammatory cytokine IL-1® from rabbit corneas. Medians and interquartile ranges of
538	pro-inflammatory cytokines are shown. Each point represents one rabbit. Kruskal-Wallis with

- 539 Dunn's post-test was performed, **a** indicates significant difference from control group,
- 540 p<0.05. n.s. indicates not significantly different than the control group. *, p<0.05; **, p<0.01.



Figure 1. Predatory bacteria and *lasR* mutation reduce *P. aeruginosa* pathogenesis in a rabbit keratitis model. A. Diagram of model in which the rabbit corneal stroma (arrow) is injected with *P. aeruginosa* strain PA14, isogenic $\Delta lasR$ mutant, or PBS followed by PBS or *B. bacteriovorus* strain HD100. B-D. Each point represents a rabbit. a indicates significant difference from control group, p<0.05. n.s. indicates not significantly different than the control group. **, p<0.01; ***, p<0.001.B. CFU from corneas at 24 h post-injection. Means and SDs are shown. C. Corneal inflammation score based on MacDonnald-Shadduck scoring system. Medians and interquartile ranges are shown and Kruskal-Wallis analysis with Dunn's post-test was performed. D. Corneal perforation frequency. Indicated groups were compared by Fisher's Exact test.



Figure 2. Bright-field microscopy and fluorescein staining of infected eyes. Representative images are shown. Fluorescein staining indicates corneal ulceration and loss of corneal epithelium.





PA14 + PBS



ΔlasR + PBS





PA14 + HD100



ΔlasR + HD100





Figure 3. Optical coherence tomography of infected eyes. Representative images are shown. Shadows are due to infiltrates that prevent imaging.

PBS + PBS

PBS + HD100





PA14 + PBS

PA14 + HD100



Figure 4. Histological analysis demonstrates neutrophil infiltrates. Representative images are shown. Magnification was 10X. Polymorphonuclear neutrophils are evident from inside the stroma and in at the anterior chamber interface.



Figure 5. Scanning micrographs of rabbit corneal surfaces. Representative images are shown. Magnification was 25x (top) and ≥500x (bottom). The corneal perforation is clearly shown in the PA14 + PBS injected cornea, and a corneal ulcer was evident in the PA14 with HD100 injected cornea.

≥500X



Figure 6. *P. aeruginosa* strain PA14 stimulation of gelatinase B (MMP9) and proinflammatory cytokine IL-1 β from rabbit corneas. Medians and interquartile ranges of pro-inflammatory cytokines are shown. Each point represents one rabbit. Kruskal-Wallis with Dunn's post-test was performed, **a** indicates significant difference from control group, p<0.05. n.s. indicates not significantly different than the control group. *, p<0.05; **, p<0.01.