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Photoinactivation of Eight Health-Relevant Bacterial Species: Determining the Importance of the Exogenous Indirect Mechanism

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1 Photoinactivation of Eight Health-Relevant Bacterial Species: Determining the Importance of the
2 Exogenous Indirect Mechanism

3

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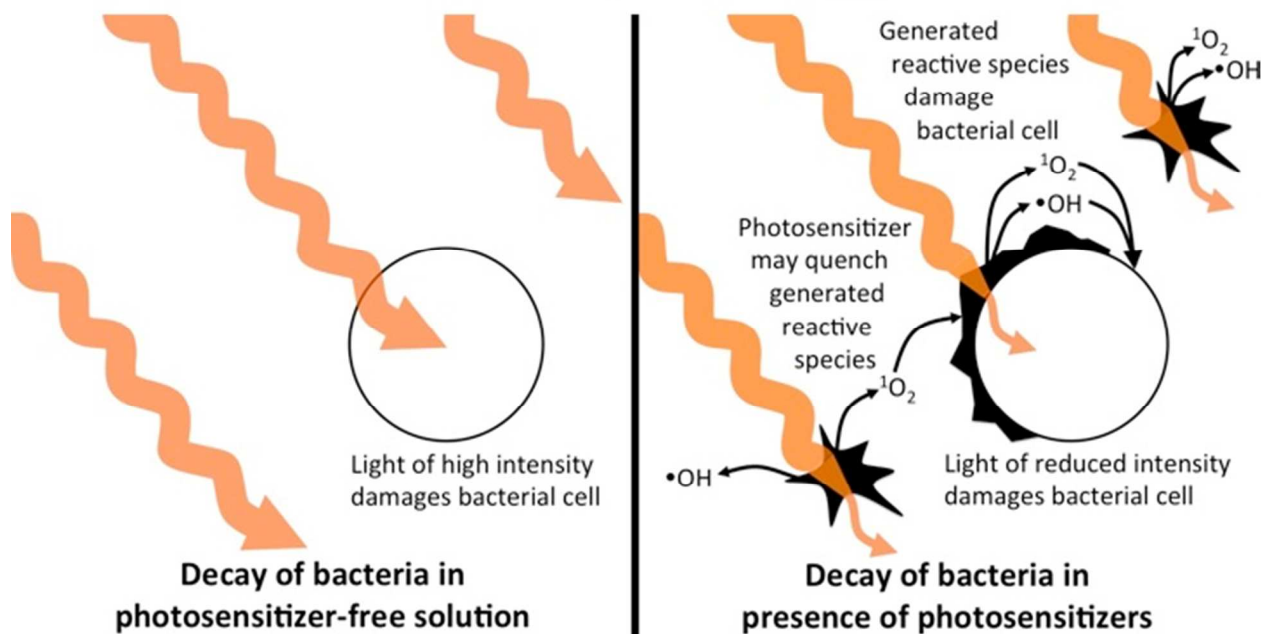
16 **Keywords:** bacteria, sunlight, disinfection, photoinactivation, exogenous, sensitizers, reactive
17 oxygen species, enterococci, E. coli

18 **ABSTRACT**

19 It is presently unknown to what extent the endogenous direct, endogenous indirect, and
20 exogenous indirect mechanisms contribute to bacterial photoinactivation in natural surface
21 waters. In this study, we investigated the importance of the exogenous indirect mechanism by
22 conducting photoinactivation experiments with eight health-relevant bacterial species
23 (*Bacteroides thetaiotaomicron*, *Campylobacter jejuni*, *Enterococcus faecalis*, *Escherichia coli*
24 K12, *E. coli* O157:H7, *Salmonella enterica* serovar Typhimurium LT2, *Staphylococcus aureus*,
25 and *Streptococcus bovis*). We used three synthetic photosensitizers (methylene blue, rose bengal,
26 and nitrite) and two model natural photosensitizers (Suwannee River natural organic matter and
27 dissolved organic matter isolated from a wastewater treatment wetland) that generated singlet
28 oxygen and hydroxyl radical. *B. thetaiotaomicron* had larger first order rate constants than all
29 other organisms under all conditions tested. The presence of the synthetic photosensitizers
30 generally enhanced photoinactivation of Gram-positive facultative anaerobes (*Ent. faecalis*,
31 *Staph. aureus*, *Strep. bovis*). Among Gram-negative bacteria, only methylene blue with *E. coli*
32 K12 and rose bengal with *C. jejuni* showed an enhancing effect. The presence of model natural
33 photosensitizers either reduced or did not affect photoinactivation rate constants. Our findings
34 highlight the importance of the cellular membrane and photosensitizer properties in modulating
35 the contribution of the exogenous indirect mechanism to the overall bacterial photoinactivation.
36
37

38 *TOC ART*

Do bacterial photoinactivation kinetics increase or decrease in the presence of photosensitizers?



39

40 *INTRODUCTION*

41 Forty-four percent of the world's population resides within 150 km of the coastline¹, and
42 eight of the world's ten largest urban centers are coastal. The shoreline and coastal waters are
43 valuable natural resources, providing society with food, recreation, revenue, and numerous
44 ecosystem services (such as nutrient cycling, runoff detoxification, and biodiversity). Despite the
45 importance of clean coastal waters to our economy and well-being, declining water quality along
46 the world's coastlines threatens ecosystem and human health^{2,3}.

47 It is estimated that globally, exposure to coastal waters polluted with pathogens from
48 wastewater causes an excess 120 million gastrointestinal (GI) and 50 million severe respiratory
49 illnesses per year⁴. In an effort to reduce recreational waterborne illnesses, US states are required
50 through provisions outlined in the BEACH Act to implement beach monitoring programs that
51 use densities of fecal indicator bacteria (FIB) to assess risk. Similar monitoring programs are in
52 place around the globe, guided by recommendations from the World Health Organization^{5,6}.

53 Although they are not usually etiologic agents of recreational waterborne illness, FIB are used to
54 evaluate beach water quality because their densities in coastal waters contaminated with
55 wastewater and urban runoff have been linked quantitatively to swimmer illness in epidemiology
56 studies⁷. When FIB densities exceed threshold values, beach advisories or closures are issued
57 warning swimmers that exposure may lead to illness.

58 The number of beach advisories and closures has grown in the US over the last 20 years,
59 in part due to the increasing number of beaches monitored by local and state agencies⁴.
60 Mitigation of microbial pollution in coastal waters has proven challenging because there are
61 many potential point and non-point sources (including runoff, sewage, bird and wild animal
62 feces, oceanic outfalls, decaying plants, and sediments)⁸, and there is imperfect knowledge about

63 the factors that modulate the abundance and distribution of different indicators and pathogens
64 once they are introduced into the environment^{9,10}. Science that yields insight into the physical,
65 chemical, and biological controls of the abundance and distribution of microbial pollution in
66 coastal waters is needed and of both national and international importance. Such knowledge
67 would guide remediation efforts, and assist in the identification of safe and dangerous conditions
68 for swimming.

69 Field observations in both marine and fresh surface waters suggest that sunlight is one of
70 the most important factors modulating FIB concentrations¹¹⁻¹³. Photoinactivation of bacteria, or
71 inactivation by means of sunlight or radiant energy, may occur via three different mechanisms:
72 endogenous direct, endogenous indirect, and exogenous indirect^{14,15}. Endogenous direct
73 photoinactivation results from the damage of cellular chromophores by photons, such as UVB
74 damage to DNA^{16,17}, while indirect photoinactivation involves energy or electron transfer to
75 form reactive species either within (endogenous) or external (exogenous) to the cell that can then
76 cause cell death. In practice, it is difficult to tease apart the endogenous mechanisms so they can
77 be considered together¹⁸. In coastal waters, colored dissolved organic matter^{19,20} can act as the
78 photosensitizers that absorb and transfer the light energy outside of the bacteria cell (exogenous
79 indirect), while chromophores within the cell can inadvertently do the same (endogenous
80 indirect).

81 It is presently unknown to what extent the three different mechanisms contribute to
82 bacterial inactivation in natural waters. The inactivation of *Escherichia coli* and to a lesser extent
83 enterococci has been measured in a wide range of sunlit waters and the observed inactivation rate
84 constants vary over several orders of magnitude^{15,21-26}. There are limited studies on sunlight
85 inactivation of waterborne bacterial pathogens such as *Campylobacter*, *Vibrio cholerae*, *Shigella*,

86 and *Salmonella*^{27–34}. These studies primarily have investigated how inactivation rate constants
87 vary between the pathogens and indicator organisms; the results are equivocal with some studies
88 reporting similar inactivation among organisms and others reporting widely differing inactivation
89 rate constants. Furthermore, a number of studies have attempted to elucidate the mechanisms of
90 bacterial inactivation in sunlit, natural waters by using FIB^{14,15,18,21,22,24,35–38}. The extent to which
91 each of the three photoinactivation mechanisms (endogenous direct, endogenous indirect, and
92 exogenous indirect) contributed to the overall photoinactivation varied between studies. Further,
93 it is unclear whether those results can be extended to bacterial pathogens.

94 The present study tests whether the exogenous indirect mechanism of bacterial
95 photoinactivation is important relative to the endogenous mechanisms in natural waters. We
96 conducted photoinactivation experiments in a solar simulator with eight health-relevant bacterial
97 species (*Bacteroides thetaiotaomicron*, *Campylobacter jejuni*, *Enterococcus faecalis*, *E. coli*
98 K12, *E. coli* O157:H7, *Salmonella enterica* serovar Typhimurium LT2, *Staphylococcus aureus*,
99 and *Streptococcus bovis*). Experiments were carried out in the presence and absence of model
100 synthetic and natural photosensitizers under full spectrum simulated sunlight. While studies have
101 previously investigated the exogenous indirect photoinactivation pathway for viruses^{19–21,39–43}, *E.*
102 *coli*^{14,15,18,21,22,36,37}, and enterococci^{14,15,18,21,37,44} this is the first study to do so for a wide range of
103 bacterial species and sensitizers. The results of this study will be used in future efforts to
104 accurately model and predict bacterial photoinactivation rate constants in natural waters as they
105 inform whether the exogenous indirect mechanism is critical to include.

106

107 ***EXPERIMENTAL MATERIALS AND METHODS***

108 **Bacterial cultivation.** The following bacteria were obtained from the American Type
109 Culture Collection (ATCC): *Bacteroides thetaiotaomicron* (ATCC 29741), *Campylobacter jejuni*
110 (ATCC 29428), *Enterococcus faecalis* (ATCC 19433), *Escherichia coli* K12 (ATCC 10798),
111 *Escherichia coli* O157:H7 (ATCC 43895), *Staphylococcus aureus* (ATCC 25923), and
112 *Streptococcus bovis* (ATCC 33317). The attenuated strain *Salmonella enterica* serovar
113 Typhimurium LT2 was obtained from the Falkow Lab of Stanford University. These organisms
114 were chosen to represent both Gram negative and positive bacteria. Each strain is a bacterial
115 pathogen or bacterial indicator organism. Individual properties and justifications for choice of
116 each bacterial species are provided in Table S1.

117 Tryptic soy broth (TSB, BD Bacto, Sparks, MD; ATCC medium 18) was used as the
118 growth media for the facultative anaerobic microorganisms (*Ent. faecalis*, *E. coli* K12, *E. coli*
119 O157:H7, *S. enterica*, *Staph. aureus*, and *Strep. bovis*) and made following manufacturer's
120 direction using deionized water. When TSB was used as the chemostat medium, it was diluted to
121 25% strength. Brucella broth (BD BBL, Sparks, MD; ATCC medium 1116) was used as the
122 growth media for the microaerophilic microorganism *Campylobacter jejuni*, and made following
123 manufacturer's direction using deionized water. The obligate anaerobe *Bacteroides*
124 *thetaitaomicron* was grown in modified TYG medium (see SI). All media were autoclaved at
125 121°C for 15 minutes prior to use. Facultative anaerobes were grown in an aerobic environment
126 (ambient air conditions) while *C. jejuni* and *B. thetaiotaomicron* were grown in a
127 microaerophilic and anaerobic environment, respectively (see SI).

128 A single stock of each bacterial species was generated for storage at -20°C. This single -
129 20°C stock was used to generate the biomass to seed the photoinactivation experiments in order
130 to minimize variance associated with starting bacterial population⁴⁵.

131 For facultative anaerobes, experiment seed was generated using chemostats (see SI) to
132 minimize the variability among replicates⁴⁵. Chemostats could not be used for microaerophiles
133 and obligate anaerobes due to the difficulty of controlling the atmosphere. *B. thetaiotaomicron*
134 and *C. jejuni* seed were generated from batch cultures (see SI). Cells were harvested by pelleting
135 and re-suspending the cells in 1 mM carbonate buffer saline (CBS, pH 7.64) three times.

136 ***Experimental solutions and procedures.*** Stock solutions of Suwannee River natural
137 organic matter (SRNOM, International Humic Substance Society, St. Paul, MN), isolated DOM
138 (iDOM), methylene blue (MB), sodium nitrite, and rose bengal (RB) were filter sterilized before
139 use (for further details, including iDOM isolation method, see SI). The panel of photosensitizers
140 was chosen to include both natural (SRNOM and iDOM) and synthetic sensitizers (MB, RB, and
141 nitrite), a range of surface charges (anionic for SRNOM, iDOM, RB, and nitrite; cationic for
142 MB), and a range of reactive oxygen species generated (Table S2). Stock sensitizer solutions
143 were added to individual 100 mL experimental beakers to reach final concentrations of 0.5 μM
144 RB, 200 mg/L as NO_2^- , 0.5 μM MB, 20 mg C SRNOM/L, and 20 mg C iDOM/L each at a total
145 volume of 50 mL. All solutions were buffered using 1 mM CBS at pH 7.64. A single bacterial
146 species was then added to each reactor so that the final starting concentrations were $5.2 \times 10^3 -$
147 5.6×10^6 colony forming units (CFU)/ml for *B. thetaiotaomicron*, $9.2 \times 10^5 - 1.2 \times 10^7$ CFU/ml
148 for *C. jejuni*, $2.9 \times 10^6 - 3.0 \times 10^7$ CFU/ml for *E. coli* K12, $1.9 \times 10^6 - 1.6 \times 10^7$ CFU/ml for *E.*
149 *coli* O157:H7, $6 \times 10^5 - 1.1 \times 10^7$ CFU/ml for *Ent. faecalis*, $1.4 \times 10^7 - 2.9 \times 10^7$ CFU/ml for *S.*
150 *enterica*, $2.0 \times 10^7 - 4.2 \times 10^7$ CFU/ml for *Staph. aureus*, and $2.4 \times 10^4 - 1.9 \times 10^6$ CFU/ml for
151 *Strep. bovis* (Table S3).

152 Due to the time and cost of acquiring the iDOM, experiments with iDOM were limited to
153 representative Gram negative and Gram positive organisms: *E. coli* K12 and *Ent. faecalis*. We
154 also limited experiments with methylene blue to *E. coli* K12 and *Ent. faecalis*.

155 Beakers containing a single sensitizer and a single bacterial species were placed in a
156 recirculating water bath, set at a temperature of 15°C, in a solar simulator (Atlas Suntest CPS+;
157 Linsengericht-Altenhaßlau, Germany). Beakers were stirred continuously. An additional beaker
158 containing 1 mM CBS and the bacterial species was covered with foil and placed in the same
159 recirculating water bath to serve as dark control for each experiment; however, the dark control
160 was not continuously stirred and was only mixed upon sampling. The temperature of
161 experimental solutions was followed during initial experiments to confirm isothermal conditions
162 (15°C); thereafter temperature measurements were discontinued. The solar simulator was set at
163 an irradiance of 400 W/m² and was equipped with a coated quartz filter and a UV special glass
164 filter to block the transmission of wavelengths below 290 nm to simulate natural sunlight
165 (passing wavelength, 290 nm < λ < 800 nm)⁴⁶. The resulting light spectrum as measured using a
166 spectroradiometer (ILT950; International Light, Peabody, MA) compared favorably to the light
167 intensity at the ground level (17 m above sea level) in Palo Alto, California, USA (37.4292° N,
168 122.1381° W) at midday in the summer (Figure S1).

169 All experiments were performed in triplicate. A single chemostat (for facultative
170 anaerobes) or batch culture (for microaerophiles and obligate anaerobes) generated the bacterial
171 seed for an entire set of experiments (i.e., a CBS experiment, a SRNOM experiment, a rose
172 bengal experiment, and a nitrite experiment), but separate chemostats or batch cultures generated
173 the bacterial seeds for each experimental biological replicate (i.e, CBS replicate 1, CBS replicate
174 2, CBS replicate 3).

175 Upon exposure to the solar simulator light source, 0.5 ml samples were aseptically
176 withdrawn from the beakers over a time course (anywhere from every 2 to 60 minutes,
177 depending on how quickly the organism lost culturability during pilot experimental runs).
178 Samples were serially diluted with CBS and bacterial colonies were enumerated by spread
179 plating appropriate dilutions in duplicate on appropriate agars (see SI). Concentrations were
180 calculated using counts from all plates with between 10 and 400 colonies after accounting for the
181 dilution and volume applied to the agar.

182 **Chemical Probes.** The steady-state bulk concentrations of singlet oxygen ($^1\text{O}_2$) and
183 hydroxyl radical ($\cdot\text{OH}$) were indirectly measured in experimental solutions identical to those
184 used in the inactivation experiments by monitoring the decay of probe compounds furfuryl
185 alcohol (FFA) and phenol⁴⁷⁻⁴⁹ (see SI). The probe compounds did not decay due to direct
186 photolysis and only decayed in the presence of the photosensitizers (Figure S2).

187 **Data Analysis.** The inactivation data obtained from each experiment were fit using a
188 shoulder-log linear model⁵⁰:

$$\frac{C(t)}{C_0} = e^{-kt} \left(\frac{e^{kS}}{1 + (e^{kS} - 1)e^{-kt}} \right) \quad (1)$$

189 where t is time (min), $C(t)$ (CFU/ml) is the measured concentration at time t , C_0 is the measured
190 concentration at time 0, S (min) is the shoulder or lag time over which there is minimal
191 inactivation of the bacteria, and k (min^{-1}) is the rate constant for the log linear portion of the
192 inactivation curve after completion of the lag time. All fitted values are non-negative. If the 95%
193 confidence interval of the shoulder length, S , crossed zero, then the inactivation data were refit
194 using a simpler first order decay model:

$$\frac{C(t)}{C_0} = e^{-kt} \quad (2)$$

195 In some cases, biological replicates were fit with different models (either Equation 1 or Equation
196 2) if one replicate had a shoulder and the other did not. It should be noted that for the case when
197 the shoulder is not different from 0, then Equation 1 simplifies to Equation 2. Fit parameters, and
198 their standard deviations and 95% confidence intervals were obtained using IGOR PRO
199 (WaveMetrics Inc., Lake Oswego, OR). A χ^2 value was generated for each model (Eq. 1 and 2)
200 and the Pearson's χ^2 test for goodness of fit was used to determine the P value, with the assumed
201 null hypothesis that the model predicted the experimental data. Model parameters were corrected
202 for light screening of UVB photons as described in the SI, and the corrected parameters are
203 hereon referred to as \hat{k} and \hat{S} and reported with units m^2/MJ and MJ/m^2 , respectively.

204 \hat{k} values were compared between bacterial species and experimental solutions using
205 analysis of variance (ANOVA) and Tukey post hoc tests. \hat{k} values were \log_{10} transformed for
206 the analysis to achieve normality. The presence or absence of a shoulder was compared between
207 experiments using Pearson's chi-square test. Statistics were completed using IBM SPSS statistics
208 (version 22).

209 \hat{k} values from treatment and controls conducted with organisms from the same
210 chemostat or batch culture (Table S4) were compared pairwise, with the null hypothesis that the
211 \hat{k} values were the same¹⁸. This approach was taken rather than pooling biological replicates as
212 variability among replicate experiments conducted with different bacterial seed has been well
213 documented⁴⁵. If the paired t-test indicated the control and treatment were significantly different
214 ($p < 0.05$) in two or more of the three biological replicate experiments and the effect indicated the
215 same trend (i.e. enhanced or suppressed inactivation in treatment relative to the control), then the
216 treatment was deemed to have a potential effect on photoinactivation, and the \hat{k} values in

217 treatment and controls were considered different. We chose this approach to be inclusive in
218 considering treatments with potential effects.

219 The inactivation rate constants \hat{k} (corrected for light screening) were pooled and
220 averaged across experimental treatments and CBS controls for which the \hat{k} values were different
221 and higher in the treatments than in the controls. $\hat{k}_{endogenous}$, defined as the inactivation rate
222 constant in the absence of exogenous sensitizers, was estimated as the average \hat{k} from the CBS
223 controls for each particular bacterial species. $\hat{k}_{exogenous}$, defined as the increase in inactivation rate
224 constant due to the presence of exogenous sensitizers, was estimated as the difference between
225 the average \hat{k} from the experiments with a particular photosensitizer and the average \hat{k} of the
226 CBS controls ($\hat{k}_{endogenous}$). $k_{exogenous}$ was then expressed in units of per time by multiplying by the
227 depth-average UVB intensity (averaged across control and treatment replicates) (Table S5).
228 Pearson's r assessed whether $k_{exogenous}$ was correlated with steady state singlet oxygen or
229 hydroxyl radical concentrations.

230

231 **RESULTS**

232 **Curve fits.** Inactivation data (Figure 1) were fit using either a shoulder-log linear model
233 or first-order decay model. Model fit parameters and their 95% confidence intervals can be found
234 in Table S3. For 114 (99%) of the data series of the 115 individual experiments, the Pearson's χ^2
235 test values for fitted curves were above 0.05, indicating that the applied model well fitted the
236 data (Table S4).

237 A shoulder was observed for 79 of the 115 inactivation curves ranging from 0.000055 to
238 0.0051 MJ/m² (Table S4). The remaining 36 data series did not require a shoulder and were fit

239 with the simplified model. Pearson χ^2 tests showed that the proportion of experiments with a
240 shoulder varied between bacterial species and sensitizer ($p < 0.05$ for both). For example,
241 experiments conducted with iDOM ($n=6$) all had a shoulder and nearly 90% of experiments
242 conducted in CBS ($n=31$) exhibited a shoulder; the remaining sensitizer experiments showed
243 similar proportions of models with and without a shoulder. All experiments conducted with *Ent.*
244 *faecalis* ($n=21$) and *Strep. bovis* ($n=12$) had shoulders and 85% of the experiments conducted
245 with *Staph. aureus* ($n=13$) had a shoulder, while 17% of experiments done with *S. enterica* had a
246 shoulder. For experiments with the other organisms, close to 50% of models included a shoulder.
247 A coarser analysis indicates that of the 47 experiments conducted with Gram-positive bacteria,
248 96% had a shoulder while 50% of the experiments conducted with Gram-negative bacteria
249 ($n=68$) had a shoulder.

250 Inactivation data of the dark controls were fit with a first-order decay model; however,
251 the slopes did not differ from zero ($p < 0.05$, data not shown).

252 ***Photoinactivation in the absence of photosensitizers.*** In CBS, \hat{k} values ranged from
253 1045 to 16907 m^2/MJ with a median of 1925 m^2/MJ (Table S4). \hat{k} was significantly different
254 between organisms (ANOVA, $p < 0.05$). Tukey post hoc testing indicated *B. thetaiotaomicron* \hat{k}
255 was greater than \hat{k} of all other organisms (average $\hat{k} = 12678 \text{ m}^2/\text{MJ}$). *C. jejuni*, *Strep. bovis*,
256 and *Staph. aureus* decayed similarly ($\hat{k} = 2100\text{-}3950 \text{ m}^2/\text{MJ}$) with lower \hat{k} than *B.*
257 *thetaiotaomicron*, but higher \hat{k} than the remaining organisms (1470-1944 m^2/MJ).

258 ***Effect of photosensitizers on photoinactivation.*** The measured bulk phase steady state
259 singlet oxygen and hydroxyl radical concentrations for all photosensitizers at the chosen
260 concentration are shown in Table S2. For RB the singlet oxygen concentration was $1.0 \times 10^{-12} \text{ M}$
261 (Table S2). MB, SRNOM, and iDOM all generated similar steady state concentrations of singlet

262 oxygen, ranging from 8.6×10^{-14} to 1.2×10^{-13} M (Table S2). The singlet oxygen concentration
263 for the nitrate solution is not reported due to potential side reactions and contribution of hydroxyl
264 radical to FFA decay^{51–53}. MB, RB, and nitrite generated similar steady state concentrations of
265 hydroxyl radicals, from 4.5×10^{-15} to 8.3×10^{-15} M, while SRNOM and iDOM generated steady
266 state concentrations approximately 10 times lower ($\sim 1.4 \times 10^{-16}$ to 2.1×10^{-16} M (Table S2)).
267 Note that phenol is a rather unspecific probe compound for hydroxyl radical as it may
268 considerably react with other types of reactive species⁴⁷. Therefore, the presented values
269 represent upper limits for bulk phase steady state hydroxyl radical concentrations. The
270 contribution of carbonate radical to phenol depletion under the chosen experimental conditions
271 was calculated and can be assumed negligible (see SI).

272 \hat{k} measured in the presence of photosensitizers ranged from 367 to 20510 m²/MJ with a
273 median of 2424 m²/MJ (Table S4). \hat{k} varied between organisms, photosensitizer present in
274 solution, and their interaction (ANOVA, $p < 0.05$ for all three terms). Post hoc testing indicated *B.*
275 *thetaitaomicron* generally had the highest \hat{k} , followed by *C. jejuni*, then the Gram-positive
276 facultative anaerobes followed by the Gram-negative facultative anaerobes. Among the
277 experimental treatments containing sensitizers, treatments with MB had among the highest \hat{k} .
278 Treatments containing RB and nitrite had smaller \hat{k} than those containing MB but higher \hat{k} than
279 those containing iDOM and SRNOM which tended to have the lowest \hat{k} .

280 Based on the criteria described in the methods section, we identified which sensitizers
281 had a potential effect on the photoinactivation of each bacterial species. For *Ent. faecalis*, the
282 presence of MB, RB, and nitrite increased \hat{k} compared to the CBS control while iDOM
283 decreased \hat{k} . For *Staph. aureus*, RB and nitrite increased \hat{k} and SRNOM decreased \hat{k} . For

284 *Strep. bovis*, RB increased \hat{k} . For *E. coli* K12, SRNOM and RB decreased \hat{k} while MB
285 increased \hat{k} relative to CBS. For *E. coli* O157:H7, RB, nitrite, and SRNOM decreased \hat{k} . For *S.*
286 *enterica*, SRNOM decreased \hat{k} . SRNOM and nitrite decreased \hat{k} of *B. thetaiotaomicron*. RB
287 increased \hat{k} of *C. jejuni*. No other differences, identified based on our criteria, were observed.
288 The average \hat{k} from pooled biological replicate experiments for the CBS control and
289 photosensitizer treatments are shown in Figure 2.

290 $\hat{k}_{exogenous}$ was between 900 and 11681 m^2/MJ for bacteria-photosensitizer treatments
291 where the sensitizer increased \hat{k} relative to the control (Table 1). These values correspond to
292 $k_{exogenous}$ between 0.07 and 1.07 min^{-1} (Table 1). $\hat{k}_{exogenous}$ and $k_{exogenous}$ were not positively
293 significantly correlated with the steady state concentrations of singlet oxygen or hydroxyl radical
294 in the experimental solutions (\square ranged from -0.5 to 0.4, $p > 0.05$).

295 The starting concentrations of bacteria varied among biological replicates. Even though
296 we aimed to have the same starting concentration, the differences were unavoidable. Multiple
297 linear regression modeling of k or \hat{k} of each bacterial species as a function of photosensitizer
298 and \log_{10} -transformed starting concentration indicated that the starting concentration no
299 significant impact on the rate constants ($p > 0.05$).

300

301 **DISCUSSION**

302 In sensitizer-free solution, the obligate anaerobe *B. thetaiotaomicron* has a rate constant
303 approximately three times greater than other bacteria. *B. thetaiotaomicron* is able to generate
304 proteins to alleviate oxidative stress, but only if the stress is applied while the bacteria have
305 access to the nutrients necessary to sustain repair mechanisms^{54,55}. *B. thetaiotaomicron* was

306 grown in anaerobic (<1% O₂) environments, where oxygen is scarce. As such, it may not have
307 developed the cellular machinery to deal with oxidative stress. However, simply being exposed
308 to an oxic environment did not lead to inactivation, as the dark controls for *B. thetaiotaomicron*
309 decayed minimally over the length of the experiment (data not shown).

310 A comparison between photoinactivation rate constants reported herein with those
311 reported in previous studies is difficult because prior studies generally do not report the spectrum
312 of the light source used in their experiments, the geometry of their reactors, and/or the
313 absorbance of their experimental solutions. This makes it difficult to correct their reported rate
314 constants (e.g., in units of per time) for UVB light screening for comparison with our results.
315 Photoinactivation rate constants (uncorrected for light screening) reported in previous studies
316 span several orders of magnitude (Table S6): *C. jejuni* ($k = 0.12 - 0.61 \text{ min}^{-1}$)²⁸, *E. faecalis* ($k =$
317 $0.003 - 0.64 \text{ min}^{-1}$)^{21,45,56,57}, *E. coli* ($k = 0.002 - 0.12 \text{ min}^{-1}$)^{22,32,58,59}, *Salmonella enterica* ($k =$
318 $0.001 - 0.80 \text{ min}^{-1}$)^{30,32-34,60,61}, *Staphylococcus spp.* ($k = 0.19 \text{ min}^{-1}$)⁶², *Bacteroides ovatus*
319 (measured by QPCR, $k = 0.06 - 0.15 \text{ min}^{-1}$)⁶³.

320 The rate constants reported herein for bacteria in sensitizer-free solutions are attributed
321 solely to endogenous mechanisms and may be representative of those in clear natural waters (like
322 open ocean waters). This assumes that carbonate radicals were not present at significant
323 concentrations in the CBS solution. Indeed, the phenol probe, which reacts with both hydroxyl
324 and carbonate radicals⁶⁴, did not decay in 1 mM CBS (Figure S2), indicating minimal steady
325 state concentrations of both radicals. It should be noted that direct application of the rate
326 constants to clear natural waters may be hindered due to our use of laboratory-grown bacteria
327 instead of those sourced, uncultured, from the natural environment. A recent study suggests the
328 latter have differential susceptibility to photoinactivation⁵⁶. Using uncultivated organisms from

329 the natural environment (including sewage) for decay experiments represents a great
330 methodological challenge; the vast majority, if not all, pathogen inactivation experiments have
331 been conducted using laboratory-cultured organisms. Future work will need to overcome this
332 challenge to determine if results from laboratory-cultivated organisms can be extended to natural
333 systems.

334 The addition of the photosensitizers enhanced photoinactivation in some instances, and
335 diminished it in others. The panel of photosensitizers were chosen to include both natural and
336 synthetic sensitizers, a range of surface charges, and varying proportions of reactive oxygen
337 species generated, and were added at concentrations that would generate those reactive oxygen
338 species at the high end or just above what is found in the environment. In sunlit surface waters,
339 such as rivers and lakes, including municipal wastewater effluent impacted water bodies, the
340 steady-state concentration of singlet oxygen has been reported to be $5.9 \times 10^{-14} - 15 \times 10^{-14}$
341 $M^{48,65}$, while hydroxyl radical steady state concentrations are several orders of magnitude lower
342 ranging from 10^{-19} to $2.7 \times 10^{-16} M^{66,67}$, although the hydroxyl radical steady state concentration
343 may be as high as $10^{-14} M$ in the presence of nitrate⁶⁸. The anionic dye rose bengal was the only
344 sensitizer to generate bulk phase steady state singlet oxygen concentrations above that reported
345 in the natural environment; however, rose bengal, the cationic dye methylene blue, and
346 photoactive nitrite all generated bulk-phase hydroxyl radicals concentration approximately ten
347 times higher than would appear in the natural environment. Suwannee River NOM and isolated
348 DOM each generated environmentally relevant bulk phase concentrations of reactive oxygen
349 species.

350 The addition of the synthetic photosensitizers (rose bengal, nitrite, methylene blue)
351 tended to have an enhancing effect on the photoinactivation of Gram-positive facultative

352 anaerobes (*Ent. faecalis*, *Staph. aureus*, *Strep. bovis*). Among the Gram-negative bacteria, only
353 methylene blue with *E. coli* K12 and rose bengal with *C. jejuni* had an enhancing effect, with the
354 remaining synthetic photosensitizers having either a negligible or diminishing effect on
355 photoinactivation rate constants. The Gram-staining procedure characterizes the cell wall
356 structure, designating the microorganisms with thinner peptidoglycan layers and an additional
357 outer membrane containing two lipid bilayers as Gram-negative, and those with thicker
358 peptidoglycan layers and without the additional outer membrane as Gram-positive⁶⁹. The Gram-
359 negative property of an additional outer membrane may confer some protection against the
360 exogenous indirect mechanism of photoinactivation^{44,70} that the Gram-positive bacteria do not
361 have, possibly explaining the trends found in this and past studies^{71,72}. Methylene blue, a
362 cationic photosensitizer, enhanced photoinactivation of both the Gram-positive *Ent. faecalis* and
363 the Gram-negative *E. coli* K12. The cationic photosensitizers have a greater affinity for the
364 negatively charged phospholipids of Gram-negative bacteria and enter more easily via self-
365 promoted uptake in comparison with the anionic photosensitizers, such as rose bengal, and
366 neutral photosensitizers^{73,74}, and are therefore potentially better able to adsorb to the cellular
367 membrane and generate reactive oxygen species close to vital cellular components within the
368 bacteria³⁷.

369 The natural photosensitizers either diminished or did not affect photoinactivation rate
370 constants of all bacteria tested. Considering that most natural photosensitizers such as Suwannee
371 River natural organic matter are anionic at circumneutral pH⁷⁵, the charge repulsion and
372 decreased associated sorption properties may prevent the natural sensitizers from adsorbing to
373 the cellular membrane which is also negatively charged, to damage it with reactive oxygen
374 species generated at locally high concentrations^{20,76}. The photosensitizers may also interact with

375 the bacteria to shield the susceptible targets of photoinactivation⁷⁷, thereby reducing
376 photoinactivation to a greater degree than that for which can be corrected by light screening.
377 Lastly, photosensitizers may react with transient reactive oxygen species⁷⁸⁻⁸², thereby acting as a
378 sink for reactive oxygen species generated within or outside the cell. We used relatively high, yet
379 still environmentally relevant, concentrations of NOM in our experiments; future work could
380 explore if lower or higher NOM concentrations give rise to different results regarding exogenous
381 indirect photoinactivation. The natural photosensitizers we used in this study may differ from the
382 multitude of photosensitizers found in the natural environment in terms of sizes, charges, ROS-
383 generating potentials, and potential for associating with cellular membranes. It is possible that
384 other NOM could induce exogenous indirect photoinactivation.

385 When the presence of a photosensitizer did enhance photoinactivation, the resulting
386 exogenous indirect photoinactivation rate constant did not correlate with bulk phase steady state
387 concentrations of singlet oxygen or hydroxyl radical. The lack of a correlation suggests that the
388 exogenous indirect mechanism depends on more than the bulk phase reactive oxygen species
389 concentrations alone. A strong possibility is that the exogenous indirect mechanism is mediated
390 by interactions between the cell and the sensitizer^{20,76}. When the sensitizer is associated with the
391 bacterial surface, localized concentrations of reactive species may be much higher than in the
392 bulk phase^{19,20,76}. There are other reactive species that we did not consider which may have
393 contributed to and correlate with exogenous indirect photoinactivation rate constants. For
394 example, it cannot be excluded that, in the 1 mM CBS buffer solution used as matrix in the
395 experiments, no secondary carbonate radicals formed in the presence of photosensitizers.
396 Carbonate radicals are formed when carbonate or bicarbonate scavenge hydroxyl-radical or
397 through scavenging reactions with excited triplet state NOM^{64,83}. If carbonate radicals did form,

398 they could potentially enhance bacterial photoinactivation. Additionally, triplet state NOM may
399 also contribute to photoinactivation. However so far, no study has provided evidence that under
400 environmentally relevant conditions carbonate radical or triplet state NOM contributes
401 significantly to bacterial inactivation. One study concluded that carbonate radicals do not
402 contribute to the inactivation of rotavirus³⁹, while another predicted that carbonate radicals
403 contributed ~1% and ~0.2% to the overall photoinactivations of the model viruses MS2 and
404 phiX174 in pond water, respectively⁸⁴. For viruses, the triplet state has been associated with the
405 exogenous indirect photoinactivation mechanism for rotavirus^{39,43}. Future work could examine
406 how sensitive different bacterial species are to different reactive species following what was
407 done by Mattle et al. for three model viruses⁸⁴. While this approach would allow us to determine
408 second order rate constants for each reactive species relative to its bulk phase concentration, it
409 may be problematic since we suspect that adsorption of the photosensitizer to the cell and
410 localized high concentrations of reactive species near the photosensitizer play an important role
411 in mediating the exogenous indirect mechanism.

412 The amount of UVB light relative to other wavelengths likely affects the contribution of
413 the exogenous indirect mechanism to the overall photoinactivation of bacteria. Removal of UVB
414 entirely from the described experiments enhances the relative contribution of exogenous
415 photoinactivation to total photoinactivation significantly⁸⁵. Thus, differences in the amount of
416 UVB present in experimental systems may affect conclusions about the importance of the
417 exogenous indirect mechanism (see supporting information of Fisher et al.⁵⁶).

418 Our photoinactivation rate constants were correct for UVB light screening (280-320 nm)
419 solely while others have corrected rate constants using UVA+UVB light (280-400 nm).
420 Including the fluence from UVA light in the calculations of photoinactivation rate constants does

421 not change the results reported herein (see Tables S7 and S8 and Figure S3). Future work to
422 uncover the precise wavelengths of light responsible for endogenous bacterial photoinactivation
423 (through determining organism-specific action spectra^{41,86,87}) will aid in the choice of
424 wavelengths to use for light screening calculations.

425 Some decay profiles exhibited a shoulder, most predominantly among the Gram-positive
426 bacteria. A shoulder may arise due to a threshold effect, for which a cell can withstand a certain
427 level of stress before death, or because multiple cellular targets independently require
428 inactivation to reach cell death⁸⁸⁻⁹¹. Field modeling efforts typically have not incorporated a
429 shoulder component¹⁸ and it is unclear if the shoulder is an artifact of the experimental set up or
430 is experienced by organisms in natural surface waters. Future work will need to address this.

431 ***Environmental relevance.*** Colored dissolved organic matter and other natural
432 photosensitizers in water have the potential to both enhance photoinactivation via the exogenous
433 indirect mechanism and diminish photoinactivation by blocking the light that would contribute to
434 the direct and indirect endogenous mechanisms. This research suggests that the bacterial
435 membrane properties (Gram-positive versus Gram-negative), and interactions between bacteria
436 and photosensitizer, determine whether or not the exogenous indirect pathway is important
437 relative to other inactivation pathways for bacteria. Furthermore, this research demonstrates the
438 inability of Suwannee River NOM and analogous NOM to significantly accelerate bacterial
439 photoinactivation under the conditions tested.

440

441 ***SUPPORTING INFORMATION***

442 Includes additional experimental details, calculations for light screening, calculations of
443 carbonate radical contribution, table of ROS concentrations in solution, tables of raw data from

444 photoinactivation experiments, results of model fittings and statistical analysis, and the natural
445 and simulated light spectra. This material is available free of charge.

446

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- 723

Bacteria:	$\hat{k}_{exogenous}$	$k_{exogenous}$
Photosensitizer	[m ² MJ ⁻¹]	[min ⁻¹]
<i>Ent. faecalis</i> :		
0.5 μM Methylene Blue	11681 ± 1958	1.07 ± 0.18
0.5 μM Rose Bengal	2788 ± 367	0.24 ± 0.03
200 mg/L Nitrite	900 ± 262	0.07 ± 0.02
<i>Staph. aureus</i> :		
0.5 μM Rose Bengal	1992 ± 904	0.16 ± 0.07
200 mg/L Nitrite	905 ± 327	0.07 ± 0.03
<i>Strep. bovis</i> :		
0.5 μM Rose Bengal	1709 ± 898	0.16 ± 0.08
<i>E. coli</i> K12:		
0.5 μM Methylene Blue	1160 ± 548	0.09 ± 0.04
<i>C. jejuni</i> :		
0.5 μM Rose Bengal	8972 ± 4786	0.79 ± 0.42

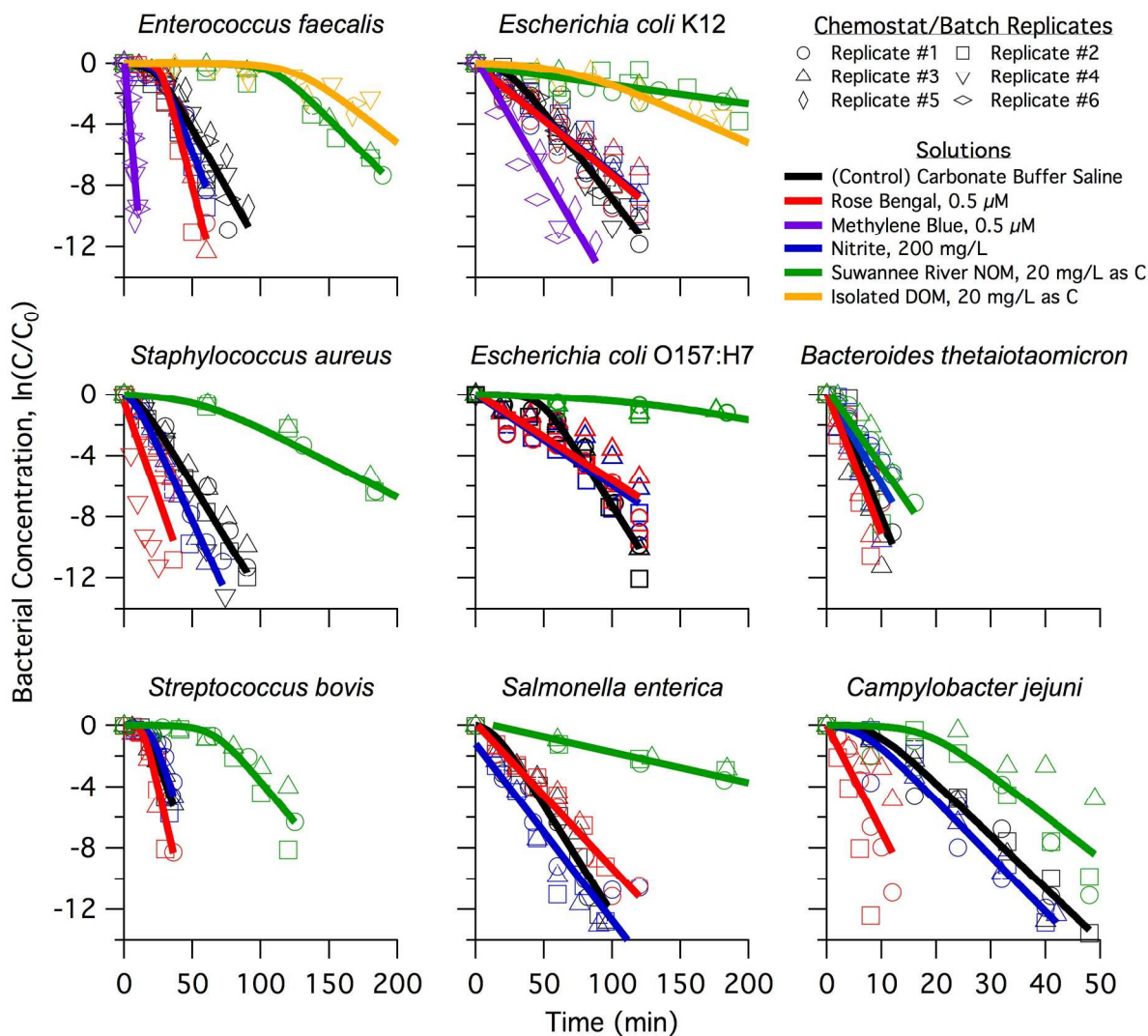
724

725 **Table 1.** Exogenous photoinactivation rate constants for bacteria in a photosensitizer treatment726 where the observed \hat{k} was higher than that in the control solution for at least two of the three727 replicates as determined by pairwise comparison. $\hat{k}_{exogenous} \pm$ standard error was estimated as the728 difference between the average \hat{k} from the pooled experiments with a photosensitizer and the729 average \hat{k} of the pooled control experiments, then the $k_{exogenous} \pm$ standard error was calculated730 from the depth-average UVB intensity (averaged across control and treatment replicates). k

731 values, their standard deviations, and their 95% confidence intervals for individual experiments

732 may be found in Table S3, while \hat{k} values, their standard deviations, and their 95% confidence

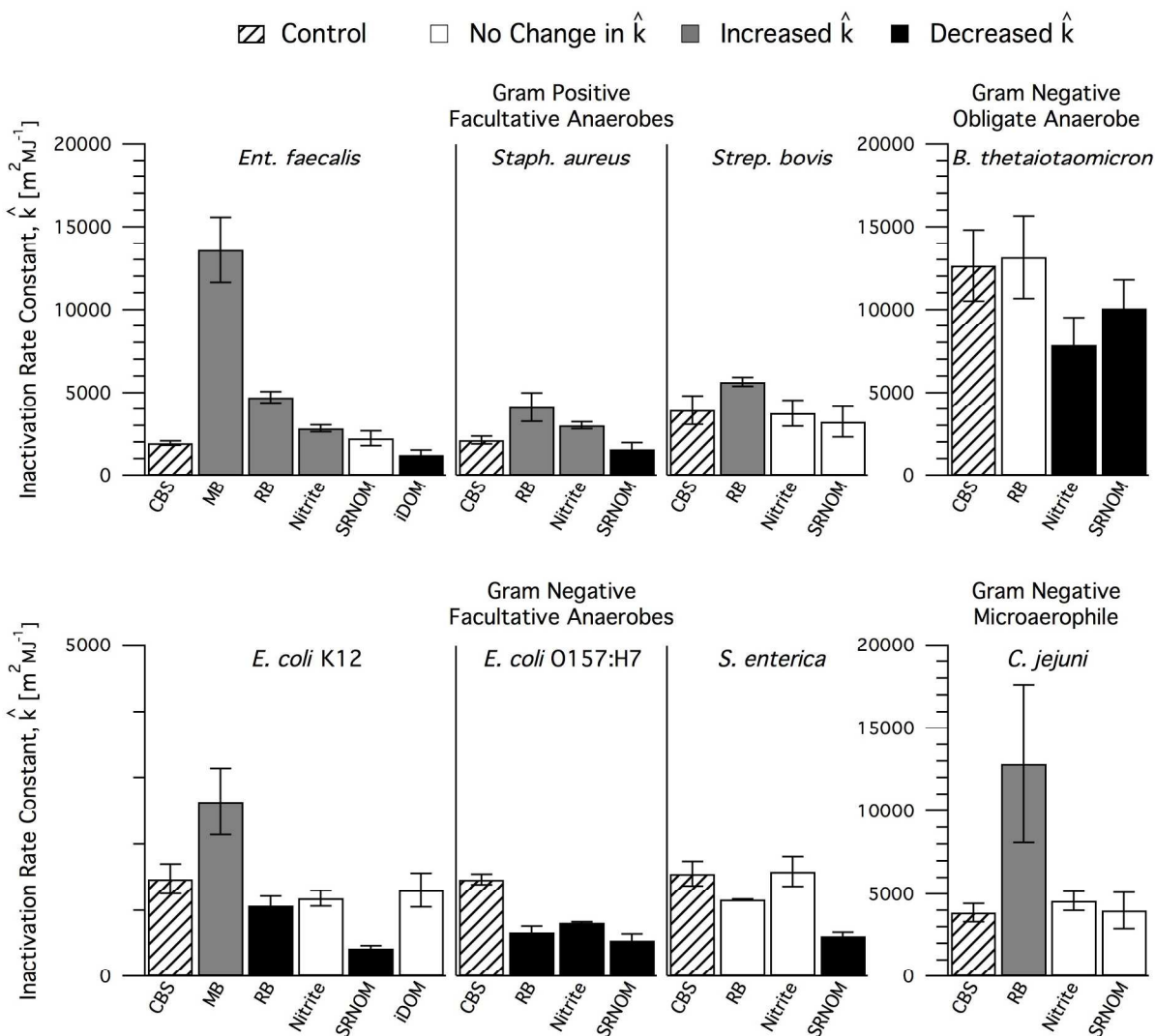
733 intervals for individual experiments may be found in Table S4.



734
 735 **Figure 1.** Photoinactivation curves of Gram-positive facultative anaerobes (*Ent. faecalis*, *Staph.*
 736 *aureus*, *Strep. bovis*), Gram-negative facultative anaerobes (*E. coli* K12, *E. coli* O157:H7, *S.*
 737 *enterica*), an obligate anaerobe (*B. thetaiotaomicron*), and a microaerophile (*C. jejuni*) in 1 mM
 738 carbonate buffer saline with various photosensitizers. The data from the replicate experiments
 739 (shown as different symbols, minimum three for each bacteria and solution) were aggregated
 740 together and fitted with either shoulder-log linear or first order kinetics model (solid lines on
 741 graphs) for ease of viewing; however the decay curves obtained from the pooled replicates were
 742 not used for hypothesis testing. As described in the methods, three biological paired replicates

743 were completed for each treatment and control. The number of replicates as indicated in the
744 legend is 6 because in some cases, not all treatments could be completed at the same time with a
745 single control, so the treatments and their paired controls were split between 2 separate
746 experiments (see Table S3).

747



748

749 **Figure 2.** Inactivation rate constants of Gram-positive facultative anaerobes, Gram-negative
 750 facultative anaerobes, an obligate anaerobe, and a microaerophile in 1 mM carbonate buffer
 751 saline (CBS) with various photosensitizers (CBS = no photosensitizer, MB = 0.5 μM methylene
 752 blue, RB = 0.5 μM rose bengal, Nitrite = 200 mg/L NO_2^- , SRNOM = 20 mg/L as C Suwannee
 753 River natural organic matter, iDOM = 20 mg/L as C dissolved organic matter isolated from a
 754 treatment wetland) when irradiated with simulated sunlight. Values are averages and standard
 755 errors from pooled biological replicates; note that pooled values were not used in hypothesis
 756 testing and are shown here for visualization. The stripped bars are the control solutions (CBS),

757 the grey bars indicate the photosensitizer increased \hat{k} , the black bars indicate the photosensitizer
758 decreased \hat{k} , the white bars indicate there was no change in \hat{k} compared to the control, as
759 determined by paired t-tests between each biological replicates matched treatment and control
760 (see methods). The \hat{k} values, their standard deviations, and their 95% confidence intervals for
761 individual experiments may be found in Table S4.