Inactivation of antibiotic resistant bacteria and inhibition of horizontal resistance gene transfer is more effective by 222 than 254 nm UV

Yijing Liu¹, Natalie M. Hull^{1,2*}

¹ College of Engineering, Department of Civil, Environmental and Geodetic Engineering, The Ohio State University, Columbus, OH 43210, USA

²The Sustainability Institute, The Ohio State University, Columbus, OH, 43210, USA

*Corresponding author: <u>hull.305@osu.edu</u>, current email: <u>nataliehull@boisestate.edu</u>

Keywords:

UV Disinfection, Antibiotic Resistant Bacteria, Antibiotic Resistance Gene, Far-UVC Disinfection, Antimicrobial Resistance, Water Treatment, Waterborne Pathogens

Supplemental Material

Table S1 Summarized log logistic parameters fitting to UV dose curves for ARB inactivation, extracted and intracellular gene damage and HGT inhibition with kinetic information

Target	UV Source	Condition	b(cm²/mJ per log reduction)	d (log reduction)	e (mJ/cm ²)	$ \begin{array}{c} \mathbf{D}_1 \\ (\mathbf{mJ/cm}^2) \end{array} $	D ₂ (mJ/cm ²)
Cells	222 nm	T	-0.67±0.04	10.66±0.97	119.29±32.54	4.11	28.61
Cells	254 nm	Inactivation	-0.64±0.14	21.57±0.17	998.89±12.55	8.99	13.54
1017 bp		Extracellular	-0.44±0.10	8.73±5.00	394.21±919.16	3.85	33.05
870 bp	222 nm	Extracellular	-0.65 ± 0.07	5.30±0.80	88.99±42.91	9.50	40.82
832 bp		Extracellular	-0.54±0.11	9.36±7.53	721.54±1743.74	14.24	72.87
266 bp		Extracellular	-0.53±0.10	9.09±6.69	1054.59±2225.60	20.67	126.03
1017 bp		Intracellular	-0.47±0.16	12.09±13.82	702.48±2882.74	4.03	25.26
870 bp		Intracellular	-1.0670.23	3.66±0.36	23.10±5.49	9.25	41.25
832 bp		Intracellular	-0.71±0.09	4.56±0.76	72.12±36.44	12.21	64.91
266 bp		Intracellular	-0.66±0.12	8.40±6.38	657.94±1156.39	32.40	97.51
1017 bp	254 nm	Extracellular	-0.52±0.20	14.17±23.31	1070.48±5262.33	7.47	42.94
870 bp		Extracellular	-0.90 ± 0.07	4.73±0.37	57.61±11.58	13.33	57.51
832 bp		Extracellular	-0.56±0.03	6.98±1.39	372.15±234.23	15.19	91.99
266 bp		Extracellular	-0.49±0.30	6.34±11.78	620.72±3898.87	19.80	150.79
1017 bp		Intracellular	-0.57±0.07	14.60±8.92	1054.19±1681.70	3.85	21.82
870 bp		Intracellular	-0.56±0.10	11.61±9.49	956.89±2171.48	9.50	27.48
832 bp		Intracellular	-0.73±0.12	9.58±7.32	574.23±943.64	14.24	51.15
266 bp		Intracellular	-1.12±0.26	3.54±1.35	119.53±83.69	20.67	114.33
HGT Inhibition	254 nm	Extra a allocita a	-0.88±0.53	3.04±3.08	112.59±264.52	8.57	236.81
	222 nm	Extracellular	-0.48±0.19	5.51±4.13	191.98±568.91	50.23	60.14
	254 nm	Intracellular	-1.33±0.19	1.52±0.17	56.74±12.55	20.14	NA
	222 nm		-0.62±0.10	13.14±9.55	1094.24±1819.24	92.90	70.07



Figure S1 Growth curves (OD vs. time) for *B.subtilis* 1A189 (green) and *B.subtilis* 1A1 (red). Error bars represent SEM of three averaged biological replicates, with three technical replicates each, at each collecting time. Modeled parameters are shown in Table S2.



Figure S2 Growth curves (Log₁₀ CFU/mL vs. time) for *B.subtilis* 1A189 (green) and *B.subtilis* 1A1 (red). Error bars represent SEM of three averaged biological replicates, with three technical replicates each, at each collecting time. Modeled parameters are shown in Table S3.

Table S2. Summary of four parameter estimates with their corresponding standard deviation of log logistic model depicting the growth curve of OD at 600 nm over time curve for *B.subtilis* 1A189 and *B.subtilis* 1A1

B.subtilis	b (1/hour)	d (OD ₆₀₀)	e (hour)
1A189	-4.29±2.06	0.52±0.09	26.02±3.02
1A1	-8.01±1.96	0.39±0.02	29.09±1.01

Table S3. Summary of slope and intercept of linear model with their corresponding standard deviation depicting the growth curve of Log10 CFU/mL over time for *B.subtilis* 1A189 and *B.subtilis* 1A1

B.subtilis	Slope (log ₁₀ CFU/mL/hour)	Intercept (log ₁₀ CFU/mL)
1A189	0.07±0.001	5.55±0.12
1A1	0.08±0.001	5.09±0.12



Figure S3 (a) D_2 for 1-log₁₀ reduction and (b) Electrical Energy per Order at D_2 of: cell inactivation, extracted DNA damage at various amplicon lengths, intracellular DNA damage at various amplicon lengths, and HGT inhibition of intracellular and extracted DNA with 254 nm (blue) and 222 nm (red) UV.