



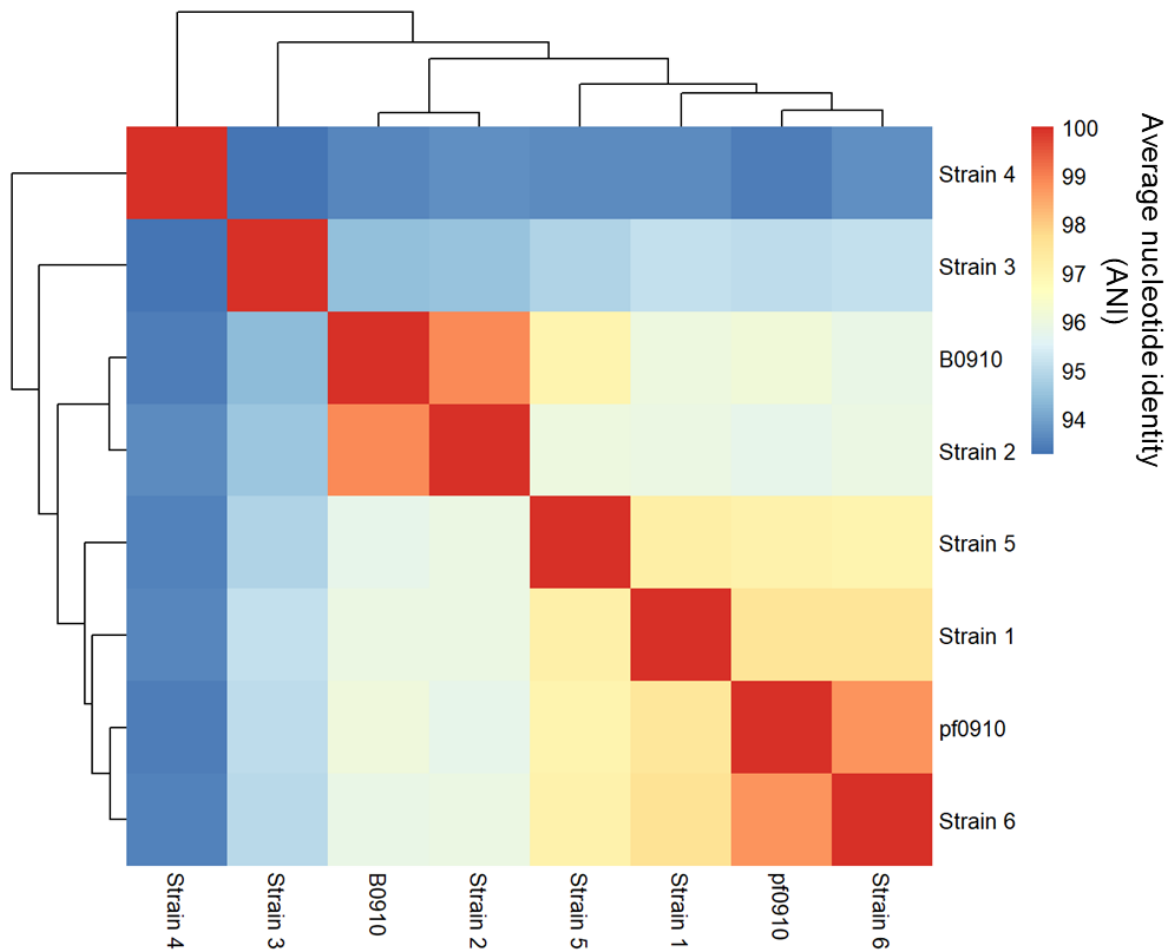
Supplement of

Effects of pH and light exposure on the survival of bacteria and their ability to biodegrade organic compounds in clouds: implications for microbial activity in acidic cloud water

Yushuo Liu et al.

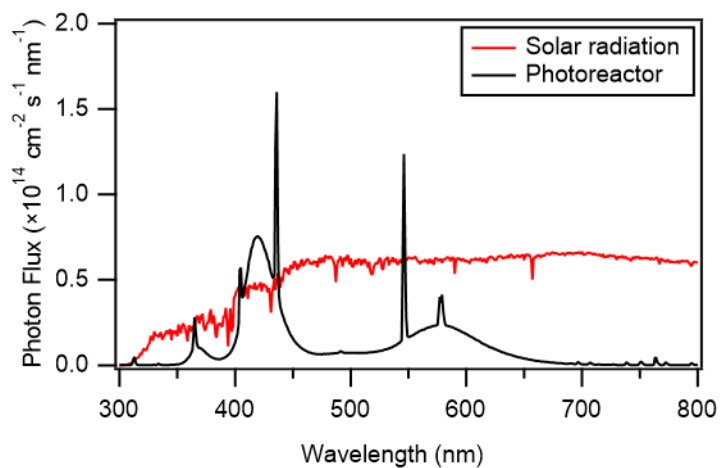
Correspondence to: Theodora Nah (theodora.nah@cityu.edu.hk)

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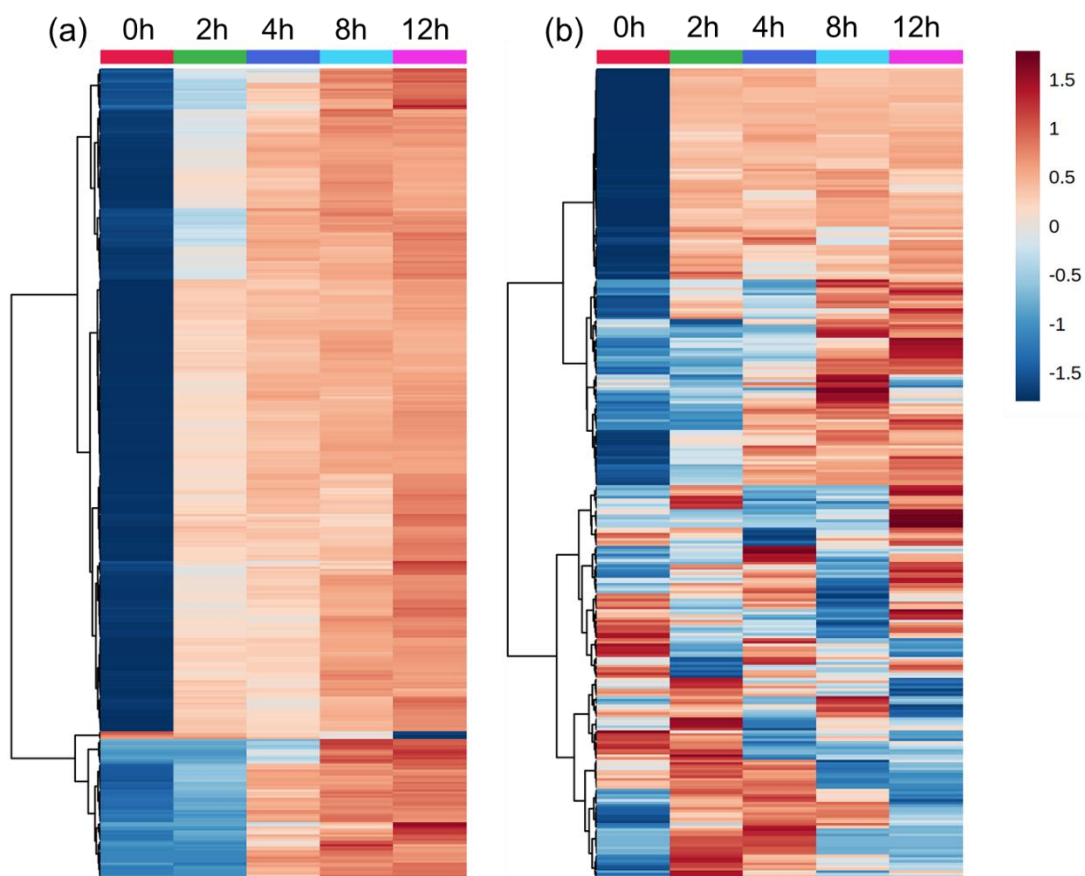
19 **Figure S1.** Average nucleotide identity (ANI) value of *Enterobacter* strains B0910, pf0910,
 20 and six others. **Strain 1:** *Enterobacter hormaechei* subsp. *oharae* DSM 16687; **Strain 2:**
 21 *Enterobacter hormaechei* subsp. *hoffmannii* DSM 14563; **Strain 3:** *Enterobacter hormaechei*
 22 ATCC 49162; **Strain 4:** *Enterobacter quasihormaechei*. GCF 004331385.1; **Strain 5:**
 23 *Enterobacter xiangfangensis* LMG27195; **Strain 6:** *Enterobacter hormaechei* subsp.
 24 *steigerwaltii* DSM 16691. Strains 1 to 6 are the closest identified neighbors with strains B0910
 25 and pf0910.



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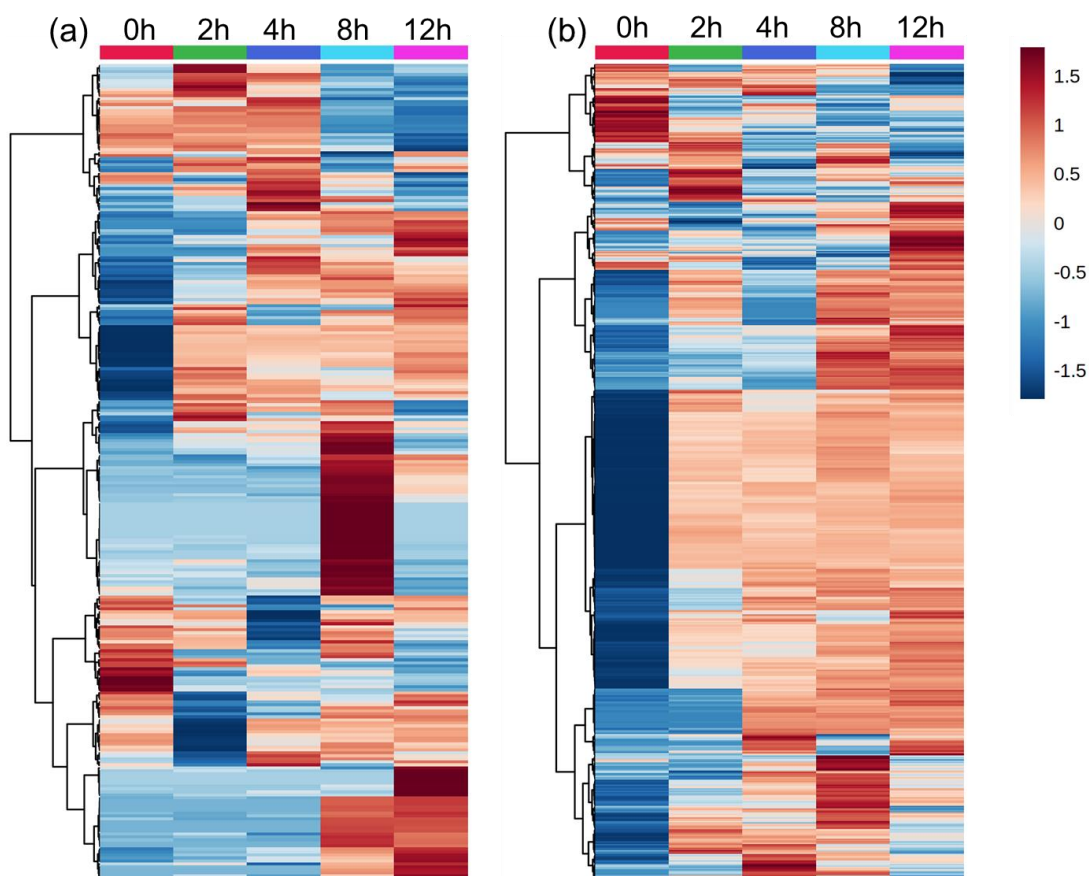
27 **Figure S2.** Photon flux inside of the photoreactor (black) and actinic flux for a fall day in Hong
28 Kong in the morning (red). One lamp with output centered at ~ 365 nm (RPR-3500A, Southern
29 New England Ultraviolet Company), four lamps with outputs centered at ~ 421 nm (RPR-
30 4190A, Southern New England Ultraviolet Company), and three lamps with outputs centered
31 at ~ 580 nm (RPR-5750A, Southern New England Ultraviolet Company) were used to
32 illuminate solutions in the photoreactor.

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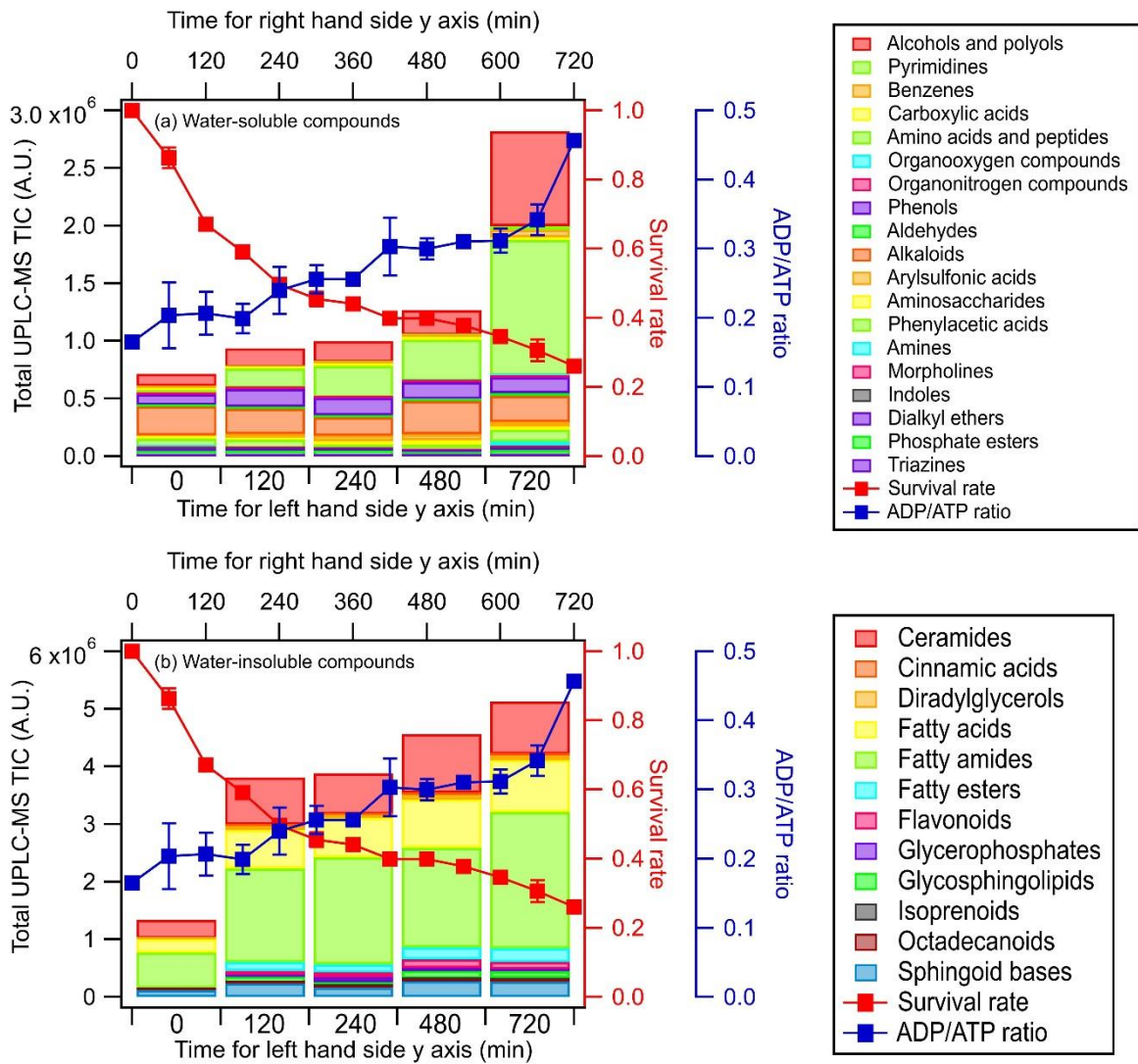
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35 **Figure S3.** Heat maps showing the time evolution of (a) water-soluble compounds and (b)
 36 water-insoluble compounds from *E. hormaechei* B0910 during exposure to simulated sunlight
 37 at pH 4.3. The heat maps were generated from non-targeted UPLC-MS analysis of samples
 38 with different light exposure times. 259 water-soluble compounds and 215 water-insoluble
 39 compounds were selected based on PLS-DA results (VIP > 1.0 criteria). The average UPLC-
 40 MS intensity of each compound at each light exposure time was obtained from the nine
 41 replicates. The average UPLC-MS intensities were subsequently log₁₀ transformed and auto
 42 scaled (i.e., mean-centered and divided by the standard deviation of each variable). The color
 43 scale ranges from red color for high abundance to blue for low abundance.



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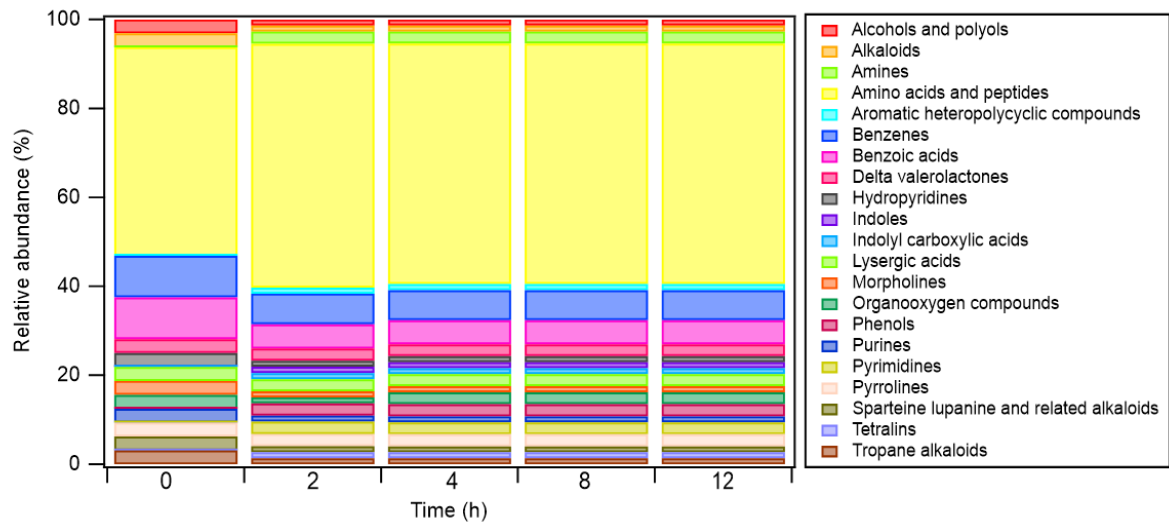
45 **Figure S4.** Heat maps showing the time evolution of (a) water-soluble compounds and (b)
 46 water-insoluble compounds from *E. hormaechei* pf0910 during exposure to simulated sunlight
 47 at pH 4.3. The heat maps were generated from non-targeted UPLC-MS analysis of samples
 48 with different light exposure times. 209 water-soluble compounds and 251 water-insoluble
 49 compounds were selected based on PLS-DA results (VIP > 1.0 criteria). The average UPLC-
 50 MS intensity of each compound at each light exposure time was obtained from the nine
 51 replicates. The average UPLC-MS intensities were subsequently log₁₀ transformed and auto
 52 scaled (i.e., mean-centered and divided by the standard deviation of each variable). The color
 53 scale ranges from red color for high abundance to blue for low abundance.



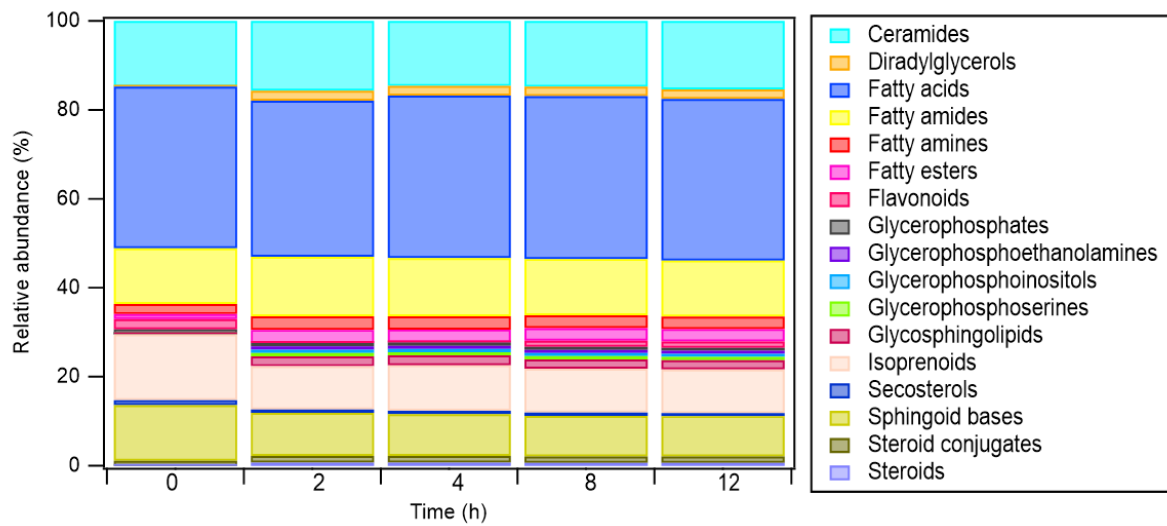
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55 **Figure S5.** Time evolution of the UPLC-MS total ion chromatograph (TIC) signals of (a)
 56 water-soluble compounds, and (b) water-insoluble compounds from *E. hormaechei* pf0910
 57 during exposure to simulated sunlight at pH 4.3 over time. These compounds are classified
 58 based on their chemical functionality. Also shown are the time evolution of the survival rate
 59 and ADP/ATP ratio of *E. hormaechei* pf0910.

(a) Water-soluble compounds



(b) Water-insoluble compounds



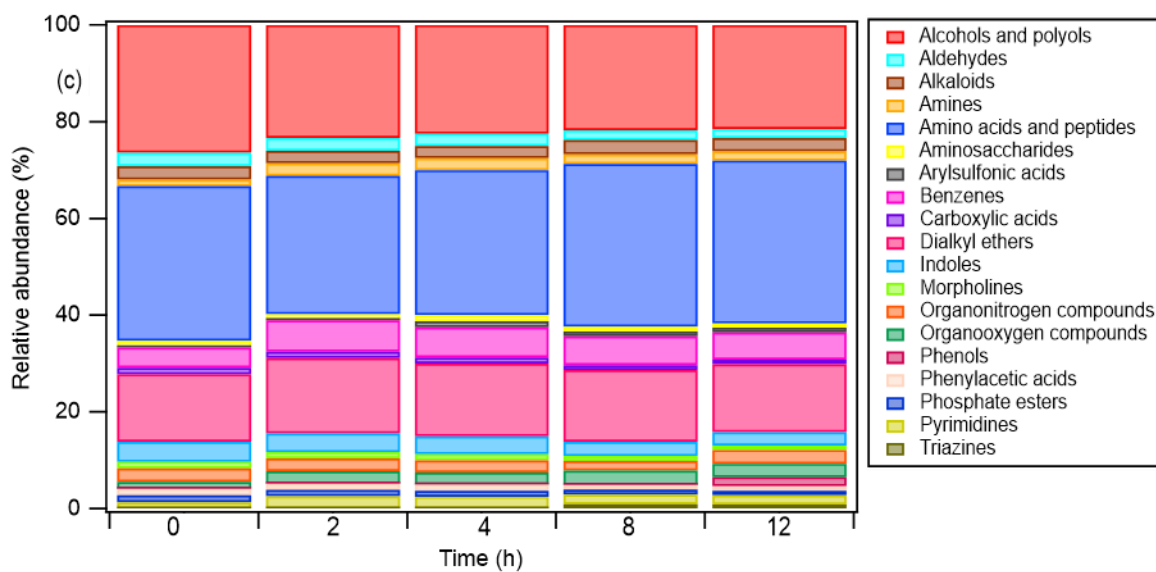
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61 **Figure S6.** Relative abundance of the different classes of (a) water-soluble compounds, and (b)

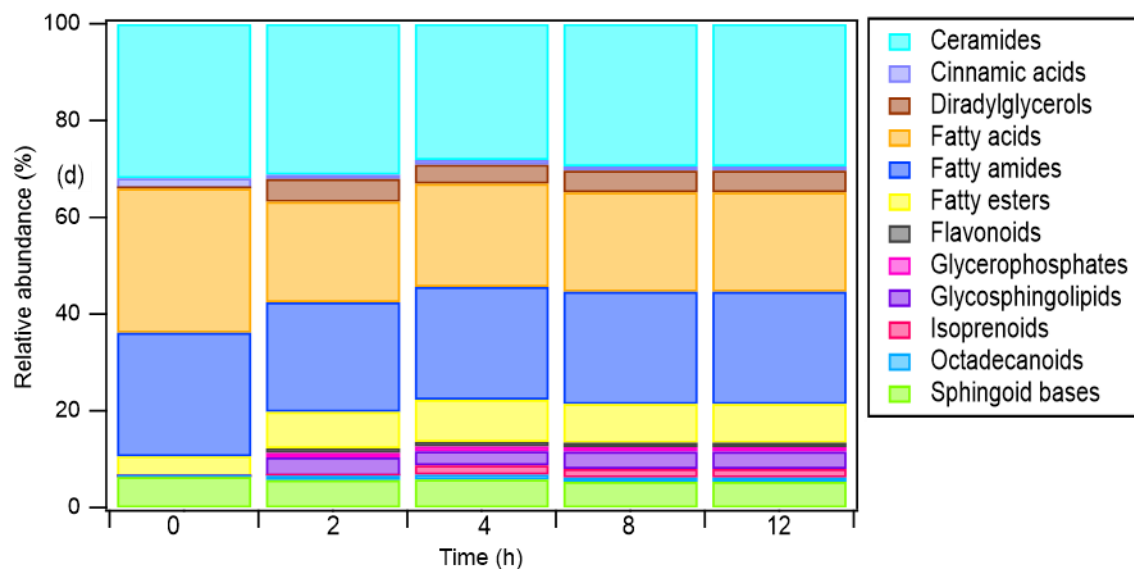
62 water-insoluble compounds from *E. hormaechei* B0910 during exposure to simulated sunlight

63 at pH 4.3.

(a) Water-soluble compounds

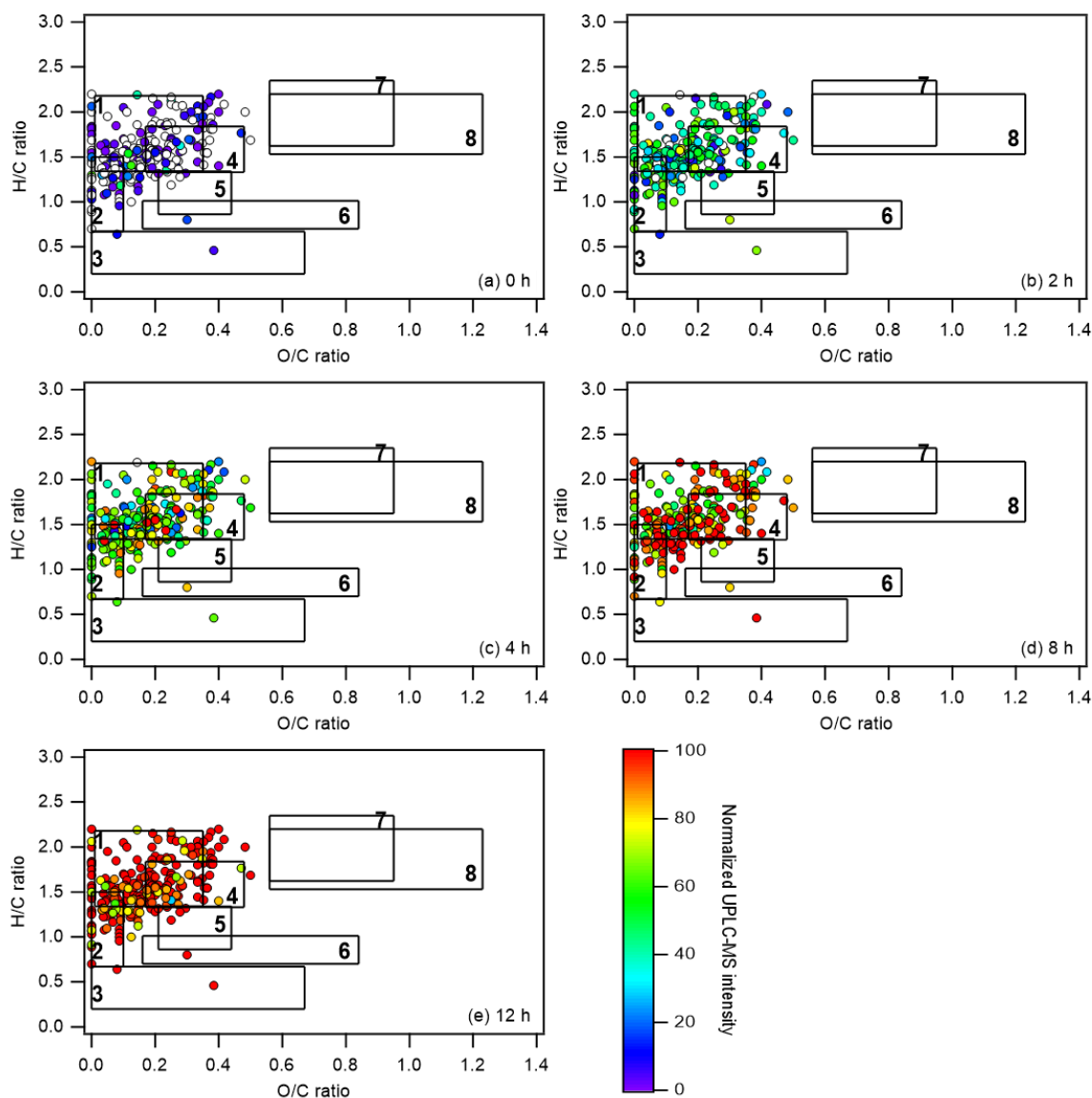


(b) Water-insoluble compounds



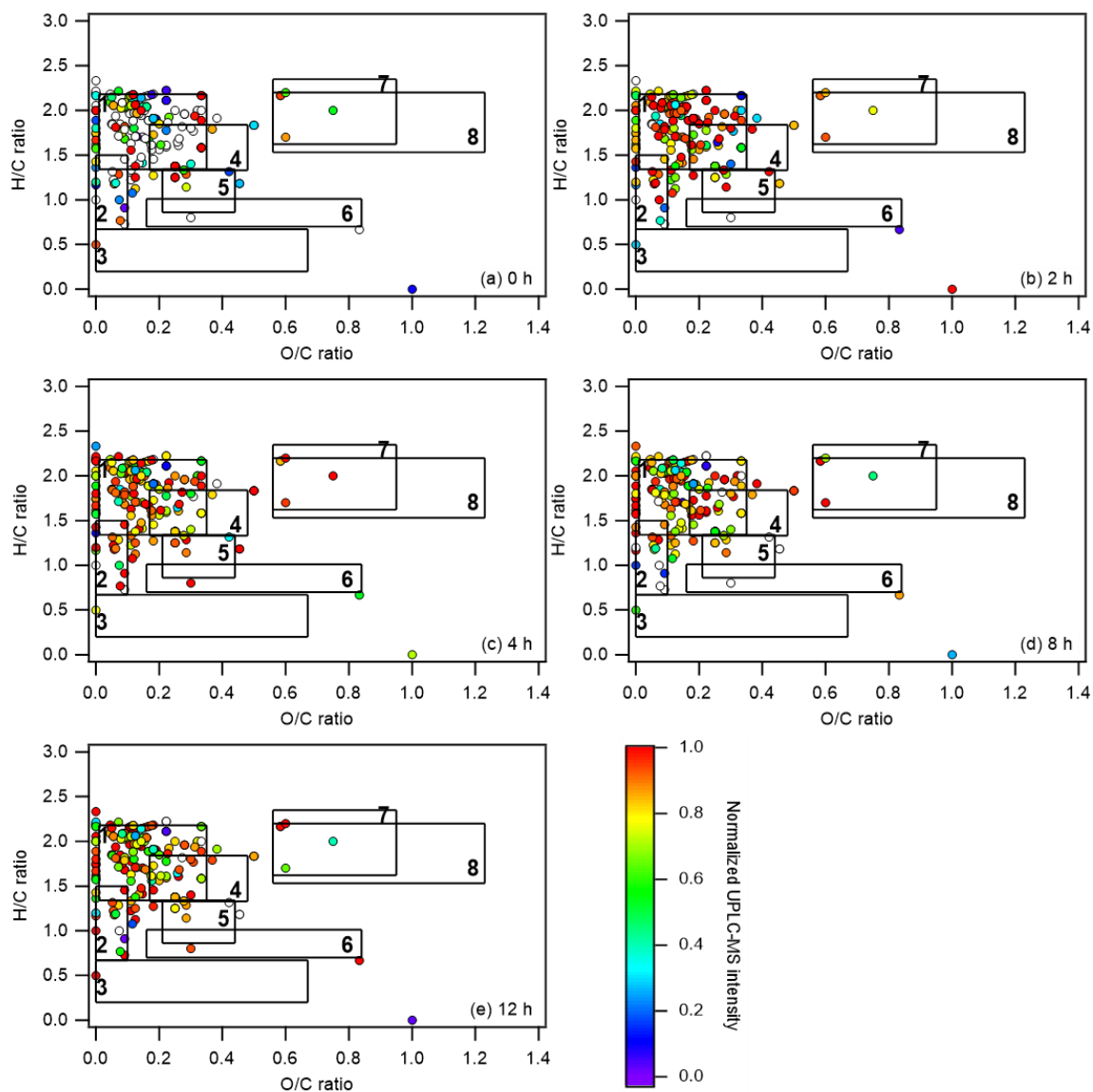
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65 **Figure S7.** Relative abundance of the different classes of (c) water-soluble compounds, and (d)
66 water-insoluble compounds from *E. hormaechei* pf0910 during exposure to simulated sunlight
67 at pH 4.3.



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69 **Figure S8.** Van Krevelen diagrams of water-soluble compounds from *E. hormaechei* B0910
70 during exposure to simulated sunlight at pH 4.3 taken at different time points of the experiment:
71 (a) 0 h, (b) 2 h, (c) 4 h, (d) 8 h, and (e) 12 h. The color of each symbol denotes its UPLC-MS
72 intensity at that specific time point normalized to its maximum UPLC-MS intensity obtained
73 during the entire experiment. Symbols that are colored white indicates that these compounds
74 were not detected at that specific time point. The Van Krevelen diagrams are divided into eight
75 chemical classes based on their O/C and H/C ratios: (1) lipids, (2) unsaturated hydrocarbons,
76 (3) condensed aromatic structures, (4) peptides, (5) lignin, (6) tannin, (7) amino sugars, and (8)
77 carbohydrates.

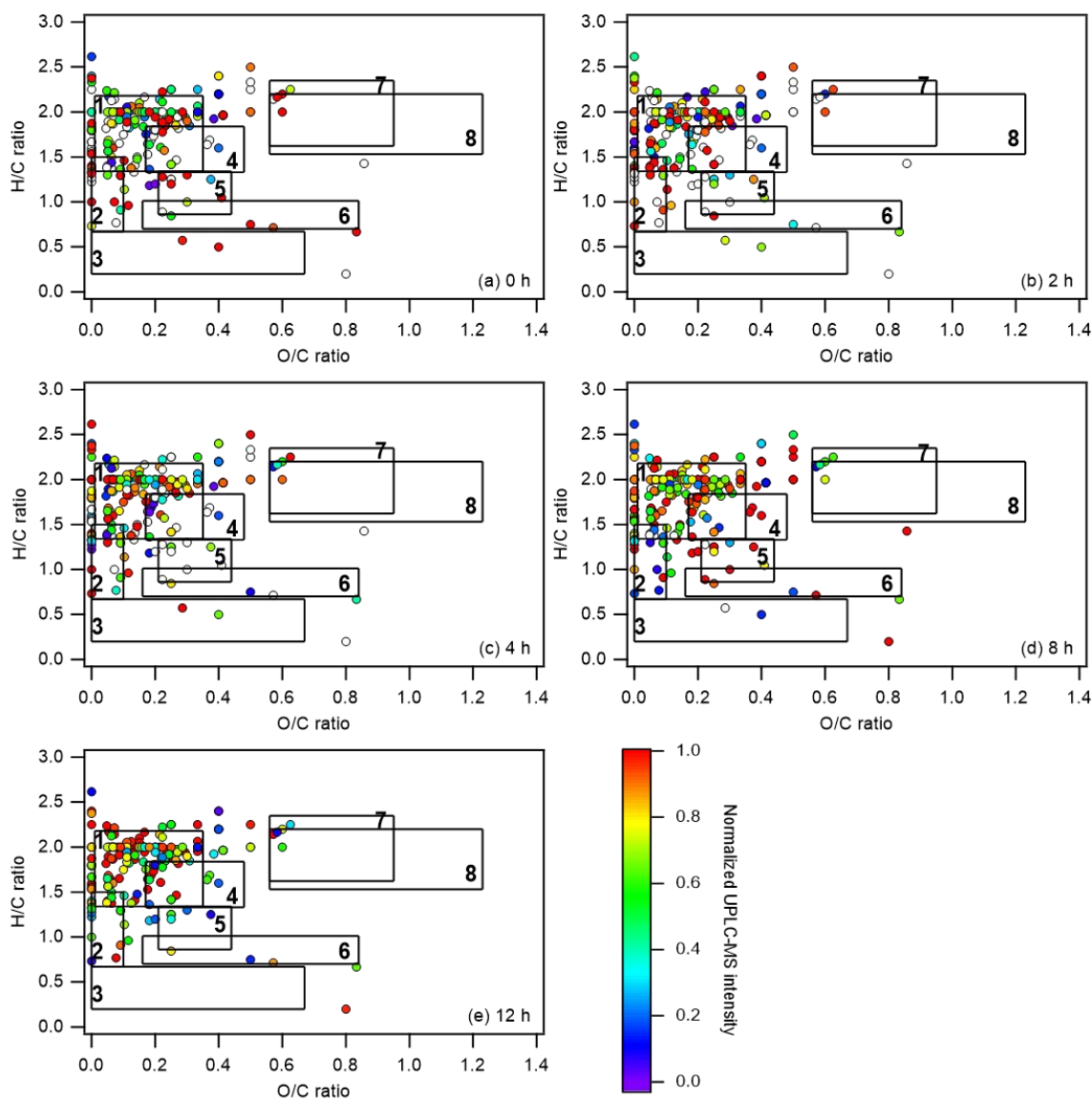


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79 **Figure S9.** Van Krevelen diagrams of water-insoluble compounds from *E. hormaechei* B0910
 80 during exposure to simulated sunlight at pH 4.3: (a) 0 h, (b) 2 h, (c) 4 h, (d) 8 h, and (e) 12 h.

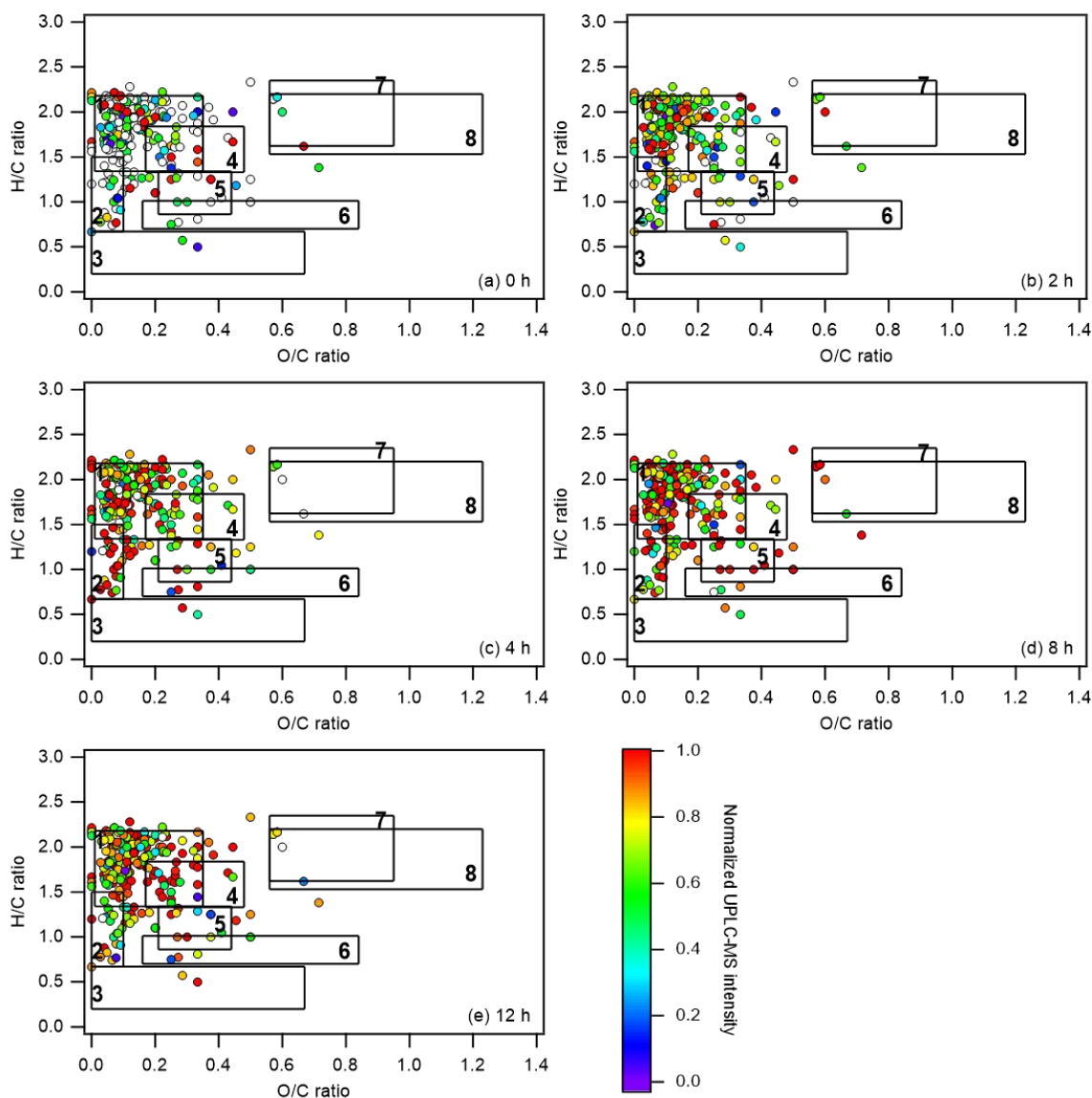
81 The color of each symbol denotes its UPLC-MS intensity at that specific time point normalized
 82 to its maximum UPLC-MS intensity obtained during the entire experiment. Symbols that are
 83 colored white indicates that these compounds were not detected at that specific time point. The

84 Van Krevelen diagrams are divided into eight chemical classes based on their O/C and H/C
 85 ratios: (1) lipids, (2) unsaturated hydrocarbons, (3) condensed aromatic structures, (4) peptides,
 86 (5) lignin, (6) tannin, (7) amino sugars, and (8) carbohydrates.



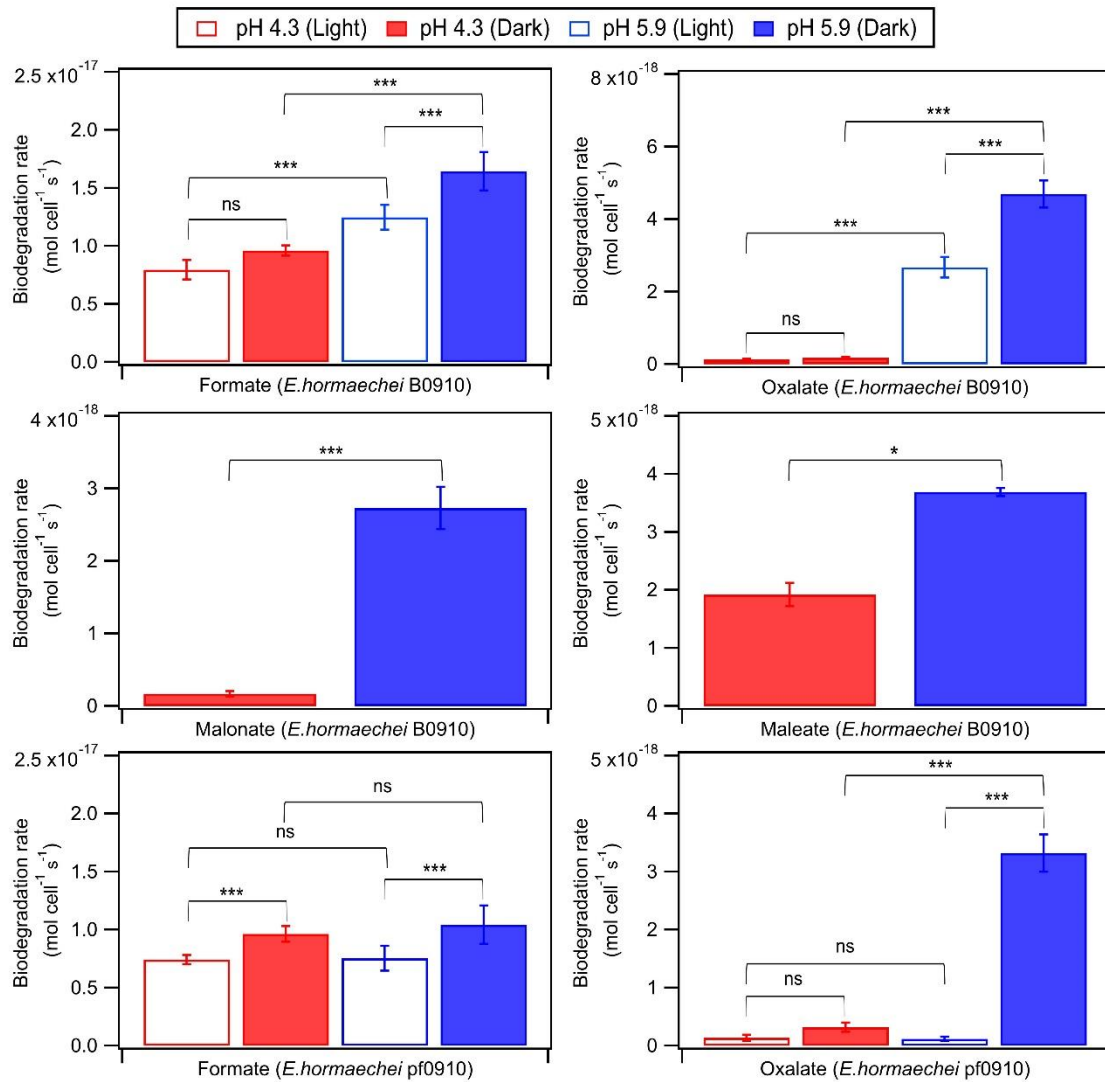
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88 **Figure S10.** Van Krevelen diagrams of water-soluble compounds from *E. hormaechei* pf0910
 89 during exposure to simulated sunlight at pH 4.3: (a) 0 h, (b) 2 h, (c) 4 h, (d) 8 h, and (e) 12 h.
 90 The color of each symbol denotes its UPLC-MS intensity at that specific time point normalized
 91 to its maximum UPLC-MS intensity obtained during the entire experiment. Symbols that are
 92 colored white indicates that these compounds were not detected at that specific time point. The
 93 Van Krevelen diagrams are divided into eight chemical classes based on their O/C and H/C
 94 ratios: (1) lipids, (2) unsaturated hydrocarbons, (3) condensed aromatic structures, (4) peptides,
 95 (5) lignin, (6) tannin, (7) amino sugars, and (8) carbohydrates.



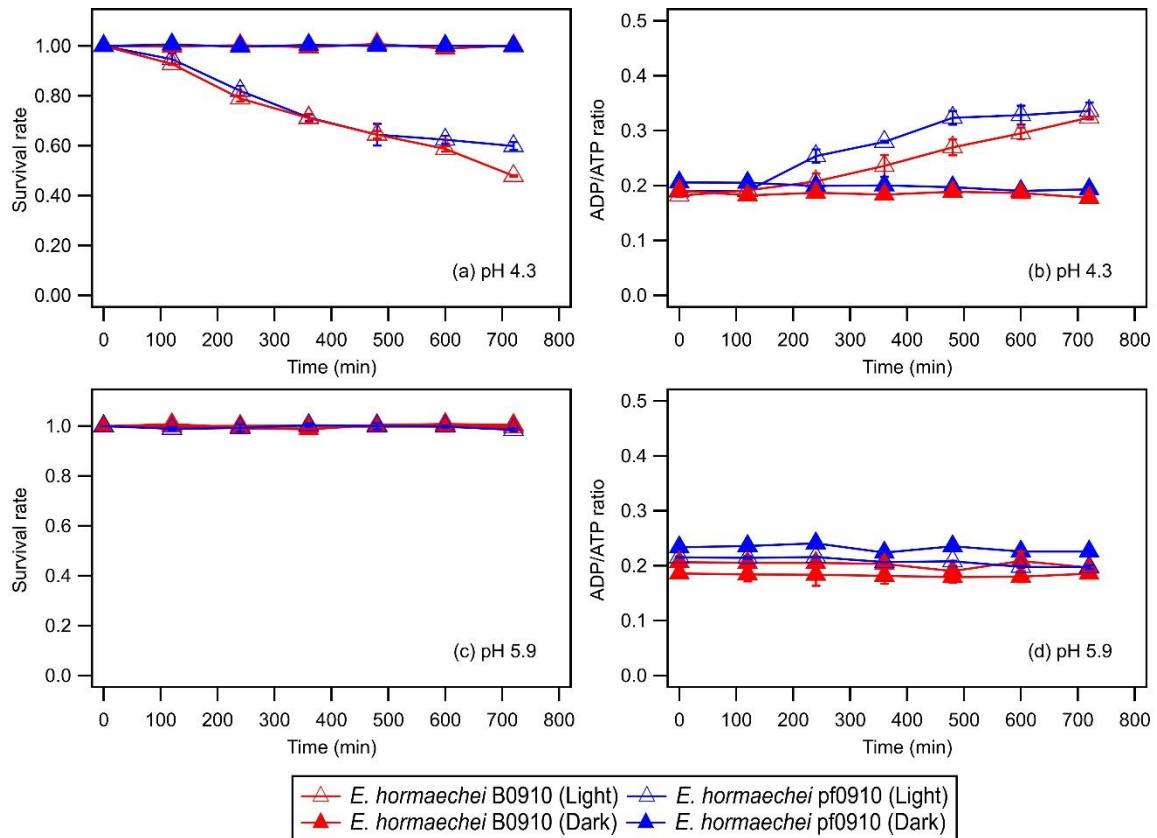
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97 **Figure S11.** Van Krevelen diagrams of water-insoluble compounds from *E. hormaechei*
 98 pf0910 during exposure to simulated sunlight at pH 4.3: (a) 0 h, (b) 2 h, (c) 4 h, (d) 8 h, and (e)
 99 12 h. The color of each symbol denotes its UPLC-MS intensity at that specific time point
 100 normalized to its maximum UPLC-MS intensity obtained during the entire experiment.
 101 Symbols that are colored white indicates that these compounds were not detected at that
 102 specific time point. The Van Krevelen diagrams are divided into eight chemical classes based
 103 on their O/C and H/C ratios: (1) lipids, (2) unsaturated hydrocarbons, (3) condensed aromatic
 104 structures, (4) peptides, (5) lignin, (6) tannin, (7) amino sugars, and (8) carbohydrates.



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106 **Figure S12.** Biodegradation rates of oxalate, maleate, and malonate by (a) *E. hormaechei*
 107 B0910 and (b) *E. hormaechei* pf0910 under light and dark conditions at pH 4.3 and pH 5.9.
 108 Error bars represent one standard deviation from the mean of biological triplicates. Statistical
 109 analysis was performed using the Student's t test (ns: not significant, *: p value < 0.05, **: p
 110 value < 0.01, ***: p value < 0.001).



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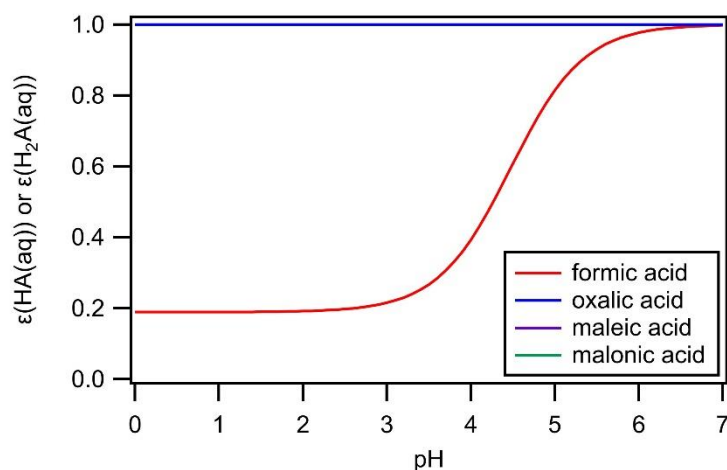
112 **Figure S13.** Survival and ADP/ATP ratios of *E. hormaechei* B0910 and *E. hormaechei* pf0910
 113 under illuminated and dark conditions at pH 4.3 and pH 5.9 in the solutions containing the
 114 seven organic acids. Error bars represent one standard deviation from the mean of biological
 115 triplicates.

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121 **Figure S14.** Calculated pH-dependent molar fraction of formic acid in the aqueous phase
 122 ($\epsilon(HA(aq))$) and pH-dependent molar fractions of oxalic acid, malonic acid, and maleic acid
 123 in the aqueous phase ($\epsilon(H_2A(aq))$) under cloud water conditions (Section S6 and Table S8).
 124 A liquid water concentration of $10^6 \mu\text{g m}^{-3}$ (Ervens et al., 2011) was assumed in these
 125 calculations. A significant fraction of formic acid will be in the gas phase at pH 4 and 5 under
 126 cloud water conditions, whereas all of the oxalic acid, malonic acid, and maleic acid will be in
 127 the aqueous phase at pH 4 and 5 under cloud water conditions (note that their values overlap
 128 one another at $\epsilon(H_2A(aq)) = 1$). These differences were due primarily to the large differences
 129 in their water solubility (i.e., Henry's law constants) (Table S8).

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141 **Table S1.** Chemical composition of the artificial cloud water used to prepare bacterial cells
 142 and perform experiments that investigated the effects of cloud water pH and light exposure on
 143 the survival and energetic metabolism of bacteria. In the experiments, the pH of the artificial
 144 cloud water was adjusted while keeping the final organic and inorganic ion composition the
 145 same.

Organic ion	μM	Inorganic ion	μM
Formate	17.1	Na^+	93
Acetate	10.2	NH_4^+	235
Pyruvate	2.7	K^+	8
Oxalate	10.3	Mg^{2+}	23
		Ca^{2+}	49
		Cl^-	138
		SO_4^{2-}	305

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149 **Table S2.** Chemical composition of the artificial cloud water used for organic acid
 150 biodegradation experiments.

Organic ion	μM	Inorganic ion	μM
Formate	50	Na^+	930
Acetate	50	NH_4^+	2350
Pyruvate	50	K^+	80
Oxalate	50	Mg^{2+}	230
Succinate	50	Ca^{2+}	490
Maleate	50	Cl^-	1380
Malonate	50	SO_4^{2-}	3050
Glutarate	50		
MSA	50		

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Table S3. Genes involved in the pH homeostasis in the two *E. hormaechei* strains.

Transporters	Protein subunits	<i>E. hormaechei</i> B0910	<i>E. hormaechei</i> pf0910
		CDS	CDS
F1F0-type ATP synthase	Subunit a, AtpB	MOG78_16595	MMW20_13045
	Subunit c, AtpE	MOG78_16590	MMW20_13050
	Subunit b, AtpF	MOG78_16585	MMW20_13055
	Subunit delta, AtpH	MOG78_16580	MMW20_13060
	Subunit alpha, AtpA	MOG78_16575	MMW20_13065
	Subunit gamma, AtpG	MOG78_16570	MMW20_13070
	Subunit beta, AtpD	MOG78_16565	MMW20_13075
	Subunit epsilon, AtpC	MOG78_16560	MMW20_13080
Kdp-type high-affinity potassium transporter	Potassium-binding ATPase subunit KdpA	MOG78_10080	MMW20_19865
	Potassium-binding ATPase subunit KdpB	MOG78_10085	MMW20_19860
	Potassium-binding ATPase subunit KdpC	MOG78_10090	MMW20_19855
	Potassium-binding ATPase subunit KdpF	MOG78_10075	Gene sequence found but CDS is not annotated. (Chromosome genome nucleotide position: 3800683-3800772)
Kup-type low-affinity potassium transporter	Kup	MOG78_16640	MMW20_13000

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167 **Table S4.** Stoichiometric ranges of the eight chemical classes in VK diagrams (Bauer et al.,
 168 2002; Jaenicke, 2005).

Chemical class	H/C	O/C
Amino sugar (Burrows et al., 2009)	$1.62 \leq \text{H/C} \leq 2.35$	$0.56 \leq \text{O/C} \leq 0.95$
Carbohydrate (Jaenicke, 2005)	$1.53 \leq \text{H/C} \leq 2.20$	$0.56 \leq \text{O/C} \leq 1.23$
Lignin (Möhler et al., 2007)	$0.86 \leq \text{H/C} \leq 1.34$	$0.21 \leq \text{O/C} \leq 0.44$
Lipid (Burrows et al., 2009)	$1.34 \leq \text{H/C} \leq 2.18$	$0.01 \leq \text{O/C} \leq 0.35$
Peptide (Attard et al., 2012)	$1.33 \leq \text{H/C} \leq 1.84$	$0.17 \leq \text{O/C} \leq 0.48$
Tannin (Hu et al., 2018)	$0.70 \leq \text{H/C} \leq 1.01$	$0.16 \leq \text{O/C} \leq 0.84$
Unsaturated hydrocarbons (Amato et al., 2005)	$0.67 \leq \text{H/C} \leq 1.5$	$0 \leq \text{O/C} \leq 0.10$
Condensed aromatic structures (Delort et al., 2010)	$0.20 \leq \text{H/C} \leq 0.67$	$0 \leq \text{O/C} \leq 0.67$

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170 **Table S5.** Genes involved in the biodegradation of organic acids in the two *E. hormaechei*
 171 strains.

Organic acid	Genes	<i>E. hormaechei</i> B0910		<i>E. hormaechei</i> pf0910	
		Biodegradation	CDS	Biodegradation	CDS
		Yes/No	Absent/Present	Yes/No	Absent/Present
Formic acid	Formate dehydrogenase	Yes	MOG78_16880; MOG78_16875; MOG78_16870; MOG78_06810	Yes	MMW20_12765; MMW20_12770; MMW20_12775; MMW20_22665
	Oxalate decarboxylase	Yes	Absent	Yes	Absent
	Oxalate oxidase	Yes	Absent	Yes	Absent
Oxalic acid	Formyl-CoA:oxalate CoA-transferase	Yes	Absent	Yes	Absent
	Succinyl-CoA:oxalate CoA-transferase	Yes	Absent	Yes	Absent
	Hypothetical protein (Cupin 2 protein) ^a	Yes	MOG78_20825	Yes	MMW20_08875
Malonic acid	Malonate decarboxylase	Yes	MOG78_18565; MOG78_18550; MOG78_18545; MOG78_18540; MOG78_18530	No	MMW20_11060; MMW20_11075; MMW20_11080; MMW20_11085; MMW20_11095
	Malonate CoA-transferase	Yes	Absent	No	Absent
	Malonate-semialdehyde dehydrogenase	Yes	Absent	No	Absent
	Malonyl-CoA/methylmalonyl-CoA synthetase	Yes	Absent	No	Absent
Maleic acid	Maleate isomerase	Yes	Absent	No	Absent
	Maleate hydratase	Yes	Absent	No	Absent
	3-isopropylmalate dehydratase ^a	Yes	MOG78_13080; MOG78_13075	No	MMW20_17000; MMW20_17005
Acetic acid	Acetyl-CoA synthetase	No	MOG78_14765	No	MMW20_14980
	Acetate kinase	No	MOG78_01250	No	MMW20_05785
	Aldehyde dehydrogenase	No	MOG78_17415	No	MMW20_12230
	ActP ^b	No	MOG78_14775	No	MMW20_14970

	SatP ^b	No	MOG78_13285	No	MMW20_16750
Methane sulfonic acid	Alkanesulfonate monooxygenase	No	MOG78_08820; MOG78_08810	No	MMW20_21175; MMW20_21185
Glutaric acid	Succinate-semialdehyde dehydrogenase / glutarate-semialdehyde dehydrogenase ^c	No	MOG78_19060; MOG78_13695	No	MMW20_10560; MMW20_16345
	Glutaryl-CoA synthetase	No	Absent	No	Absent
	Glutarate dioxygenase	No	Absent	No	Absent

^a Genes are not canonical but may involve in the biodegradation of organic acids.

^b Transporter proteins involved in uptake of acetic acid for biodegradation

^c No reverse catalysis in the direction from glutarate to glutarate-semialdehyde has been reported in the literature.

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178 **Table S6.** Concentration of radicals and cells used to estimate the loss rates by biodegradation
179 and chemical reactions in Table S6.

Radical concentration/ Cell concentration	Area	Concentration	Reference
·OH (M)	Remote	2.2×10^{-14}	(Vaitilingom et al., 2010)
	Marine	2.0×10^{-12}	(Vaitilingom et al., 2013)
	Urban	3.5×10^{-15}	(Morris et al., 2014)
NO ₃ · (M)	Remote	5.1×10^{-15}	(Morris et al., 2017)
	Marine	6.9×10^{-15}	(Hu et al., 2018)
	Urban	1.4×10^{-13}	(Huang et al., 2021)
Cell (cell L ⁻¹)		8.0×10^7	(Zhang et al., 2021)

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181 **Table S7.** Estimations of the loss rates of formate, oxalate, and malonate by biodegradation and chemical reactions (i.e., $\cdot\text{OH}$ oxidation (daytime) and $\text{NO}_3\cdot$ (nighttime)). These loss rates were calculated based on
 182 concentrations and pH measured at the different sites. Equations used in these calculations can be found in Section S6. References used to obtain the pH of cloud/rainwater and organic acids are indicated in superscripts.
 183 The biodegradation and chemical reaction loss rates calculated here were used to generate Figure 5.

184 **Daytime**

Location (remote)	Category	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})		
				bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (remote)		bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (remote)
Mount Lu(Möhler et al., 2007)	Cloud	3.81 (Vaitilingom et al., 2012)	10.83 (Wei et al., 2017)	1.34×10^{-10}		5.72×10^{-10}	4.95 (Zhu et al., 2018)	1.06×10^{-12}		1.74×10^{-11}
Mount Lu(Wei et al., 2017)	Rain	4.44 (Peng et al., 2019)	10.21 (Amato et al., 2005)	1.26×10^{-10}		5.39×10^{-10}	2.54 (Burrows et al., 2009)	5.44×10^{-13}		8.92×10^{-12}
Mount Heng(Amato et al., 2017)	Cloud	3.8 (Amato et al., 2005)	19.65 (Amato et al., 2017)	2.43×10^{-10}		1.04×10^{-9}	5.11 (Wei et al., 2017)	1.10×10^{-12}		1.80×10^{-11}
Mount Heng(Delort et al., 2010)	Rain	4.35 (Vaitilingom et al., 2010)	14.30 (Vaitilingom et al., 2013)	1.77×10^{-10}		7.55×10^{-10}	1.66 (Ervens and Amato, 2020)	3.55×10^{-13}		5.83×10^{-12}
Mangdang Mountain(Ariya et al., 2002)	Rain	4.81 (Husárová et al., 2011)	7.90 (Vaitilingom et al., 2011)	9.78×10^{-11}		4.17×10^{-10}	1.80 (Jaber et al., 2020)	3.86×10^{-13}		6.34×10^{-12}
Taiwan(Jaber et al., 2021)	Cloud	3.91 (Joly et al., 2015)	5.74 (Davey and O'toole, 2000)	7.11×10^{-11}		3.03×10^{-10}	6.60 (Delort et al., 2010)	1.42×10^{-12}		2.32×10^{-11}
Kleiner Feldberg, Germany(Flemming and Wingender, 2010)	Cloud	3.9-4.6 (Vaitilingom et al., 2012)	3.26 (Matulova et al., 2014)	4.03×10^{-11}		1.72×10^{-10}	ND			
Whiteface Mountain, USA(Amato et al., 2005)	Cloud	3.1-4.4 (Peng et al., 2019)	25.20 (Amato et al., 2005)	3.12×10^{-10}		1.33×10^{-9}	9.66 (Pye et al., 2020)	2.07×10^{-12}		3.40×10^{-11}
Rax, Austria(Shah et al., 2020)	Cloud	3.84 (Li et al., 2020)	13.25 (Pye et al., 2020)	1.64×10^{-10}		7.00×10^{-10}	5.11 (Shah et al., 2020)	1.10×10^{-12}		1.80×10^{-11}
Sonnblick, Austria(Qu and Han, 2021)	Cloud	5.0-6.5 (Anglada et al., 2015)	6.30 (Joly et al., 2015)		9.79×10^{-11}	3.33×10^{-10}	1.89 (Vaitilingom et al., 2011)		3.61×10^{-12}	6.65×10^{-12}
Mount Tai, China(Joly et al., 2015)	Cloud	4.6 (Jaber et al., 2021)	31.80 (Jaber et al., 2020)	3.94×10^{-10}		1.68×10^{-9}	11.10 (Li et al., 2020)	2.38×10^{-12}		3.91×10^{-11}
Shangzhong(Qu and Han, 2021)	Rain	ND	4.95 (Peng et al., 2019)	6.13×10^{-11}		2.61×10^{-10}	1.16 (Chen et al., 2012)	2.48×10^{-13}		4.07×10^{-12}
São Paulo State, Brazil(Després et al., 2012)	Rain	4.96 (Ding et al., 2015)	7.80 (Zhou et al., 2018)		1.21×10^{-10}	4.12×10^{-10}	1.20 (Prokof'eva et al., 2021)		2.29×10^{-12}	4.22×10^{-12}

Location (Marine)	Category	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})		
				bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (marine)		bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (marine)
Puerto Rico	Cloud	5.5 (Romano et al., 2019)	1.00 (Ruiz-Gil et al., 2020)		1.55×10^{-11}	4.80×10^{-9}	0.50 (Romano et al., 2021)		9.55×10^{-13}	1.60×10^{-10}
Puerto Rico(Tsai and Kuo, 2013)	Rain	5.3 (Löflund et al., 2002)	0.20 (Sun et al., 2016)		3.11×10^{-12}	9.60×10^{-10}	0.00 (Li et al., 2020)			

Location (Urban)	Category	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})			
				bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (urban)		bio (pH ~4)	bio (pH ~5)	$\cdot\text{OH}$ (urban)	
Puy de dome(Vaitilingom et al., 2010)	Cloud	6.1 (Vaitilingom et al., 2011)	4.90 (George et al., 2015)			7.61×10^{-11}	2.35×10^{-8}	1.00 (Huang et al., 2018)		1.91×10^{-12}	3.20×10^{-10}
Shenzhen, South China(Misovich et al., 2021)	Rain	4.56 (Li et al., 2020)	2.26 (Tsai and Kuo, 2013)	2.80×10^{-11}			0.58 (Löfflund et al., 2002)	1.23×10^{-13}		3.22×10^{-13}	
Anshun(Sun et al., 2016)	Rain	4.67 (Li et al., 2020)	8.77 (Vaitilingom et al., 2010)	1.09×10^{-10}			2.84 (Jaber et al., 2020)	6.09×10^{-13}		1.59×10^{-12}	
Newark US East Coast(Jaber et al., 2021)	Rain	4.6 (Vaitilingom et al., 2011)	4.44 (Jaber et al., 2020)	5.50×10^{-11}			0.68 (Jaber et al., 2021)	1.46×10^{-13}		3.81×10^{-13}	
Hong Kong SAR(Bearson et al., 1997)	Cloud	3.87 (Lund et al., 2014)	17.10 (Bearson et al., 1997)	2.12×10^{-10}			10.30 (Davey and O'toole, 2000)	2.21×10^{-12}		5.77×10^{-12}	
Puy de dome(Delort et al., 2010)	Cloud	3.9 (Flemming and Wingender, 2010)	33.20 (Vaitilingom et al., 2012)	4.11×10^{-10}			9.30 (Matulova et al., 2014)	1.99×10^{-12}		5.21×10^{-12}	

185 ND: No data

186 **Nighttime**

Location (remote)	Category	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})			Malonate (μM)	Malonate loss rate (M s^{-1})		
				bio (pH ~4)	bio (pH ~5)	$\text{NO}_3\cdot$ (remote)		bio (pH ~4)	bio (pH ~5)	$\text{NO}_3\cdot$ (remote)		bio (pH ~4)	bio (pH ~5)	$\text{NO}_3\cdot$ (remote)
Mount Lu(Guan and Liu, 2020)	Cloud	3.81 (Davey and O'toole, 2000)	10.83 (Delort et al., 2010)	1.69×10^{-10}		2.32×10^{-12}	4.95 (Flemming and Wingender, 2010)	2.07×10^{-12}		1.11×10^{-12}	ND			
Mount Lu(Vaitilingom et al., 2012)	Rain	4.44 (Matulova et al., 2014)	10.21 (Bianco et al., 2018)	1.59×10^{-10}		2.19×10^{-12}	2.54 (Laszakovits and Mackay, 2022)	1.06×10^{-12}		5.69×10^{-13}	ND			
Mount Heng(Watson et al., 2007)	Cloud	3.8 (Rivas-Ubach et al., 2018)	19.65 (Matulova et al., 2014)	3.06×10^{-10}		4.21×10^{-12}	5.11 (Bianco et al., 2016)	2.14×10^{-12}		1.15×10^{-12}	ND			
Mount Heng(Tyagi et al., 2015)	Rain	4.35 (Jaber et al., 2021)	14.30 (Bianco et al., 2019)	2.23×10^{-10}		3.06×10^{-12}	1.66 (Vaitilingom et al., 2010)	6.94×10^{-13}		3.71×10^{-13}	ND			
Mangdang Mountain(Vaitilingom et al., 2011)	Rain	4.81 (Fankhauser et al., 2019)	7.90 (Makuc et al., 2001)	1.23×10^{-10}		1.69×10^{-12}	1.80 (Tilgner et al., 2021)	7.55×10^{-13}		4.04×10^{-13}	1.40 (Koutny et al., 2006)		5.16×10^{-12}	4.00×10^{-14}
Taiwan(Guan and Liu, 2020)	Cloud	3.91 (Vaitilingom et al., 2011)	5.74 (Jaber et al., 2020)	8.95×10^{-11}		1.23×10^{-12}	6.60 (Jaber et al., 2021)	2.77×10^{-12}		1.48×10^{-12}	0.16 (Herrmann et al., 2010)	3.65×10^{-14}		4.57×10^{-15}

Kleiner Feldberg, Germany(Vaitilingom et al., 2011)	Cloud	3.9-4.6 (Jaber et al., 2020)	3.26 (Jaber et al., 2021)	5.08×10^{-11}		6.98×10^{-13}	ND				ND		
Whiteface Mountain, USA(Herrmann et al., 2010)	Cloud	3.1-4.4 (Ervens et al., 2003)	25.20 (Herrmann et al., 2010)	3.93×10^{-10}		5.40×10^{-12}	9.66 (Vaitilingom et al., 2010)	4.05×10^{-12}	2.17×10^{-12}		7.69 (Vaitilingom et al., 2011)	1.75×10^{-12}	2.20×10^{-13}
Rax, Austria(Fankhauser et al., 2019)	Cloud	3.84 (Pye et al., 2020)	13.25 (Li et al., 2020)	2.07×10^{-10}		2.84×10^{-12}	5.11 (Pye et al., 2020)	2.14×10^{-12}	1.15×10^{-12}		1.92 (Shah et al., 2020)	4.38×10^{-13}	5.49×10^{-14}
Sonnblick, Austria(Qu and Han, 2021)	Cloud	5.0-6.5 (Vaitilingom et al., 2012)	6.30 (Zhu et al., 2018)		1.32×10^{-10}	1.35×10^{-12}	1.89 (Peng et al., 2019)		1.19×10^{-11}	4.24×10^{-13}	0.38 (Ariya et al., 2002)	1.42×10^{-12}	1.10×10^{-14}
Mount Tai, China(Vaitilingom et al., 2010)	Cloud	4.6 (Husárová et al., 2011)	31.80	4.96×10^{-10}		6.81×10^{-12}	11.10 (Vaitilingom et al., 2011)	4.65×10^{-12}	2.49×10^{-12}		ND		
Shangzhong(Xu et al., 2009)	Rain		4.95	7.71×10^{-11}		1.06×10^{-12}	1.16	4.84×10^{-13}	2.59×10^{-13}		ND		
São Paulo State, Brazil(Coelho et al., 2011)	Rain	4.96 (Coelho et al., 2011)	7.80 (Coelho et al., 2011)		1.63×10^{-10}	1.67×10^{-12}	1.20 (Coelho et al., 2011)		7.57×10^{-12}	2.69×10^{-13}	ND		

Location (marine)	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})			Malonate (μM)	Malonate loss rate (M s^{-1})		
			bio (pH ~4)	bio (pH ~5)	NO_3^- (marine)		bio (pH ~4)	bio (pH ~5)	NO_3^- (marine)		bio (pH ~4)	bio (pH ~5)	NO_3^- (marine)
Puerto Rico(Gioda et al., 2011)	Cloud	5.5 (Gioda et al., 2011)	1.00 (Gioda et al., 2011)			0.50 (Gioda et al., 2011)				ND			
Puerto Rico(Gioda et al., 2011)	Rain	5.3 (Gioda et al., 2011)	0.20 (Gioda et al., 2011)			ND				ND			
Puy de dôme(Vaitilingom et al., 2013)	Cloud	6.1 (Vaitilingom et al., 2013)	4.90 (Vaitilingom et al., 2013)			1.00 (Vaitilingom et al., 2013)				0.40 (Vaitilingom et al., 2012)		1.47×10^{-12}	1.55×10^{-14}

Location (urban)	pH	Formate (μM)	Formate loss rate (M s^{-1})			Oxalate (μM)	Oxalate loss rate (M s^{-1})			Malonate (μM)	Malonate loss rate (M s^{-1})		
			bio (pH ~4)	bio (pH ~5)	NO_3^- (urban)		bio (pH ~4)	bio (pH ~5)	NO_3^- (urban)		bio (pH ~4)	bio (pH ~5)	NO_3^- (urban)
Shenzhen, South China(Huang et al., 2010)	Rain	4.56 (Huang et al., 2010)	2.26 (Huang et al., 2010)			0.58 (Huang et al., 2010)				ND			
Anshun(Zhang et al., 2011)	Rain	4.67 (Zhang et al., 2011)	8.77 (Zhang et al., 2011)			2.84 (Zhang et al., 2011)				ND			
Newark US East Coast(Song and Gao, 2009)	Rain	4.6 (Song and Gao, 2009)	4.44 (Song and Gao, 2009)			0.68 (Song and Gao, 2009)				0.29(Song and Gao, 2009)		6.61×10^{-14}	2.27×10^{-13}
Hong Kong SAR(Li et al., 2020)	Cloud	3.87 (Li et al., 2020)	17.10 (Li et al., 2020)			10.30 (Li et al., 2020)				1.36 (Zhao et al., 2019)		3.10×10^{-13}	1.07×10^{-12}
Puy de dome(Vaitilingom et al., 2013)	Cloud	3.9 (Vaitilingom et al., 2013)	33.20 (Vaitilingom et al., 2013)			9.30 (Vaitilingom et al., 2013)				3.50 (Vaitilingom et al., 2013)		7.97×10^{-13}	2.74×10^{-12}

187 ND: No data

188 **Table S8.** Acid dissociation constants and Henry's law coefficients at 25 °C used to generate
 189 $\varepsilon(HA(aq))$ and $\varepsilon(H_2A(aq))$ S curves in Figure S14

Organic acid	First acid dissociation constant (K_{a1}) (mol L ⁻¹)	Second acid dissociation constant (K_{a2}) (mol L ⁻¹)	Henry's law constant (H_{HA} or H_{H_2A}) (mol L ⁻¹ atm ⁻¹)
Formic acid	1.78×10^{-4} (Haynes, 2014)	Not applicable	9.53×10^3
Oxalic acid	5.62×10^{-2} (Haynes, 2014)	1.55×10^{-4} (Haynes, 2014)	6.11×10^8 (Nah et al., 2018) ^a
Malonic acid	1.48×10^{-3} (Williams, 2022)	2.04×10^{-6} (Williams, 2022)	3.85×10^{10} (Compernelle and Müller, 2014)
Maleic acid	1.26×10^{-2} (Weast and Astle, 1981)	8.51×10^{-7} (Weast and Astle, 1981)	1.42×10^{10} (Lide and Frederikse, 1995)

190 ^aWhile we used the Henry's law coefficient provided by Nah et al. (2018), it should be noted
 191 that the authors obtained this value by taking the average of $H_{C_2H_2O_4}$ values provided by Clegg
 192 et al. (1996), Compernelle and Muller (2014) and Saxena and Hildemann (1996), and
 193 accounted for the effect of temperature using the equations provided by Sander (2015).

194

195 Section S1. Genome assembly, annotation, and taxonomic analysis

196 Genome assembly of the sequencing reads was performed using the NECAT pipeline
 197 (v0.0.1_update20200803) (Chen et al., 2021) with the default parameters. The reads were first
 198 corrected (PREP_OUTPUT_COVERAGE = 40, CNS_OUTPUT_COVERAGE = 30,
 199 MIN_READ_LENGTH = 3000) and then the corrected reads were assembled
 200 (OVLFP_FAST_OPTIONS = -n 500 -z 20 -b 2000 -e 0.5 -j 0 -u 1 -a 1000,
 201 OVLFP_SENSITIVE_OPTIONS = -n 500 -z 10 -e 0.5 -j 0 -u 1 -a 1000). Both the correction
 202 and assembly steps were progressive with multiple processing steps to improve the accuracy
 203 and completeness. The quality of the assembled genomes was evaluated using the
 204 Benchmarking Universal Single-copy Orthologs (BUSCO v5.3.1) tool based on the database
 205 of enterobacterales_odb10 (Manni et al., 2021). For both strains B00910 and pf0910, complete
 206 circular chromosomes and plasmids were obtained.

207 Genome annotation was performed using Prokka (v1.14.6) (Seemann, 2014) with the
 208 default parameters. Whole genome-based taxonomic analysis was conducted using the Type

209 (Strain) Genome Server (TYGS) (Meier-Kolthoff and Göker, 2019). Average Nucleotide
210 Identity (ANI) was calculated by fastANI (v1.33) (Jain et al., 2018). Metabolic pathways were
211 analyzed using the KEGG Mapper (Kanehisa et al., 2022) and the RAST server (Aziz et al.,
212 2008). The sequences of the two genomes have been deposited in NCBI under the BioProject
213 number PRJNA812965.

214 **Section S2. Extraction of water-insoluble and water-soluble biological material and** 215 **organic compounds for UPLC-MS analysis**

216 A modified Bligh & Dyer (BD) protocol was performed to extract water-insoluble
217 organic compounds (Sündermann et al., 2016). Briefly, 3 mL of methanol (Duskan, LC-MS
218 grade)/chloroform (RCI, HPLC grade) (1:2, v/v) was added to a filtered 5 mL sample solution
219 and vortexed for 5 min, after which the samples were centrifuged at 3000 rpm for 10 min at 10
220 °C. The bottom layer was collected into a clean 2 mL centrifuge tube and dried in a concentrator
221 using nitrogen gas. The dried extracts were redissolved in 500 µL of acetonitrile (Duskan, LC-
222 MS grade) and stored at -20 °C prior to UPLC-MS analysis. Solid-phase extraction (SPE) was
223 performed to remove the inorganic salts and extract the water-soluble organic compounds using
224 hydrophobic lipophilic balanced (HLB) cartridges (Oasis HLB, 6cc 500 mg). The HLB
225 cartridges were first preconditioned with 1 mL methanol and 2 mL Milli-Q water. A 10 mL
226 filtered sample solution was then loaded into the SPE cartridge and washed with 20 mL Milli-
227 Q water under vacuum at a flow rate of 5 mL/min. The elution was performed by adding 1.5
228 mL methanol (Duskan, LC-MS grade). The eluent was evaporated to dryness under nitrogen
229 gas and reconstituted in 500 µL acetonitrile (Duskan, LC-MS grade).

230 **Section S3. UPLC-MS operation, data processing, and statistical analysis**

231 Chromatographic separation was performed on a Kinetex HILIC LC column (100 × 2.1
232 mm, 2.6 µm, 100 Å, Phenomenex). The flow rate was fixed at 0.3 mL/min with ultra-pure
233 water containing 5 mM ammonium acetate (Fisher, LC-MS grade) as mobile phase A and
234 acetonitrile (Duskan, LC-MS grade) for mobile phase B. The following gradient program was
235 used: 0 to 2 min 95% A; 2 to 4 min linear gradient to 80% B; 4 to 11 min linear gradient to

236 65% B; 11 to 12.5 min 65% B; 12.5 to 13 min linear gradient to 95% B; 13 to 15 min
237 equilibration wash with 95% B. Injection volume was set at 10 μ L. The information dependent
238 analysis (IDA) acquisition was acquired with MS scan (100 to 1200 m/z) followed by MS/MS
239 scan (50 to 1200 m/z) in positive ion mode. The following MS conditions were used: 30 PSI
240 curtain gas, 60 PSI ion source gas, 3000 V ESI ion spray voltage, 320 $^{\circ}$ C source temperature,
241 10 V collision energy for MS, and 80 V declustering potential. MS/MS was acquired with a
242 collision energy was 20 V with 5 V spread. The raw MS data was processed for peak detection,
243 retention time correction, alignment, and integration using the XCMS software built into the
244 web-based Galaxy platform (<https://umsa.cerit-sc.cz/>) (Gowda et al., 2014). The processed data
245 was then uploaded to MetaboAnalyst 5.0 (<https://www.metaboanalyst.ca/>) (Pang et al., 2021)
246 to identify cellular compounds that had prominent ion intensities.

247 The raw UPLC-MS data first underwent preprocessing, normalization, and quality
248 control steps using the XCMS software built into the web-based Galaxy platform (available at:
249 <https://umsa.cerit-sc.cz/>). The raw data was processed for peak detection, alignment, and
250 framing. This generated a table that displayed the retention time, mass-to-charge ratio (m/z),
251 and the intensity/peak area for each peak. The quality control step was performed to assess the
252 stability of the intensities of peaks (“features”) between samples. This was performed using
253 quality control samples, which were mixtures of equal amounts of experimental samples taken
254 at each time point of the experiment. The relative standard deviation (RSD) of each feature in
255 the quality control sample was compared to those in the experimental samples. Features with
256 higher RSD in the quality control sample than in the experimental samples were excluded,
257 while features with $RSD < 30\%$ were retained for further analysis. Multivariable statistical
258 analysis was performed on the retained features using principal component analysis (PCA) with
259 95% confidence ellipse and partial least squares discrimination analysis (PLS-DA) to identify
260 potential discriminations between the experimental samples. Heatmaps were generated to
261 determine how the retained features changed at different time points during the experiment. A
262 selection of discriminant ions and buckets was done based on the variable importance in
263 projection (VIP) values. Features with VIP values greater than 1.0 were used for the
264 identification step. MS/MS analysis was performed for the structural identification of

265 compounds. The structure of each compound was deduced based on its adducts, isotopes, and
266 MS/MS fragments using the SCIEX OS-Q software (AB Sciex). Information about
267 compounds' chemical structures, m/z, and retention times were subsequently uploaded to
268 MetaboAnalyst 5.0 (<https://www.metaboanalyst.ca/>), which used this information to identify
269 the compounds.

270 **Section S4. IC operation**

271 Organic acid concentrations were measured using a Dionex ICS-1100 (ThermoFisher
272 Scientific) system. Separation was achieved using a Dionex IonPac AS18 (4 × 250 mm) anion
273 exchange column (Thermo Scientific) equipped with a Dionex IonPac AG18 (4 × 50 mm)
274 guard column (Thermo Scientific). 16 mM potassium hydroxide (Fisher, ≥85%) was used as
275 the mobile phase at a flow rate of 1.0 mL/min for a 30 min run time. Each aliquot of solution
276 was passed through a syringe filter before IC analysis.

277 **Section S5. Estimation of biodegradation and chemical reaction rates (M s⁻¹) in cloud 278 water**

279 **S5.1. Biodegradation**

280 The decay in the concentration of a specific organic acid as a function of time (0 to 12
281 hours) during a biodegradation experiment can be described by the following equation:

$$282 \frac{d[Acid]}{dt} = k'_{cell} \times [Acid] = k_{cell,acid} \times [cell]_{experiment} \times [Acid]_{experiment}$$

283 where k'_{cell} (s⁻¹) is the pseudo first order rate constant obtained from fitting the decay of the
284 organic acid, and $[Acid]_{experiment}$ (mol L⁻¹) is the initial concentration of the organic acid
285 used in the biodegradation experiment. k'_{cell} is the product of the concentration of bacteria cells
286 used in the experiment ($[cell]_{experiment}$, cell L⁻¹) and the biodegradation rate constant
287 ($k_{cell,acid}$, L cell⁻¹s⁻¹).

288 The loss rate of the organic acid in cloud water resulting from biodegradation is:

$$289 \frac{d[Acid]_{cloud}}{dt} = k_{cell,acid} \times [cell]_{cloud} \times [Acid]_{cloud}$$

290 where $[cell]_{cloud}$ ($cell L^{-1}$) is the concentration of bacteria cells present in cloud water, and
 291 $[Acid]_{cloud}$ ($mol L^{-1}$) is the concentration of the organic in cloud water.

292 **S5.2. Chemical reactions**

293 The loss rates of the organic acid in cloud water resulting from reactions with $\cdot OH$ and
 294 $NO_3\cdot$ are:

$$295 \frac{d[Acid]_{cloud}}{dt} = k_{OH,acid} \times [\cdot OH]_{cloud} \times [Acid]_{cloud}$$

$$296 \frac{d[Acid]_{cloud}}{dt} = k_{NO_3,acid} \times [NO_3\cdot]_{cloud} \times [Acid]_{cloud}$$

297 where $k_{OH,acid}$ ($L mol^{-1}s^{-1}$) and $k_{NO_3,acid}$ ($L mol^{-1}s^{-1}$) are the rate constants for the
 298 reactions of the organic acid with $\cdot OH$ and $NO_3\cdot$, respectively, and $[\cdot OH]_{cloud}$ ($mol L^{-1}$) and
 299 $[NO_3\cdot]_{cloud}$ ($mol L^{-1}$) are the concentrations of $\cdot OH$ and $NO_3\cdot$ in cloud water, respectively.

300 **Section S6. Gas-aqueous phase partitioning of monocarboxylic and dicarboxylic acids**

301 Meskhidze et al. (2003) and Guo et al. (2016) previously introduced the concept of “S
 302 curves”, which describe how the pH of the aqueous phase affects the gas-aqueous partitioning
 303 of acidic and basic species. It is assumed that the equilibrium between gas and aqueous phases
 304 involves the dissolution of the acidic/basic species into the aqueous phase, followed by the
 305 dissociation of the dissolved species. Assuming unity activity coefficients, for monocarboxylic
 306 acids (HA, e.g., formic acid), the pH-dependence of the molar fraction of HA in the aqueous
 307 phase ($\varepsilon(HA(aq))$) is described by the following equation (Nah et al., 2018):

$$308 \varepsilon(HA(aq)) = \frac{H_{HA}WRT(10^{-pH} + K_{a1}) \times 0.987 \times 10^{-14}}{10^{-pH} + H_{HA}WRT(10^{-pH} + K_{a1}) \times 0.987 \times 10^{-14}}$$

309 where W is liquid water concentration ($\mu g m^{-3}$), H_{HA} ($mole L^{-1} atm^{-1}$) is the Henry’s law
 310 constants for monocarboxylic acid, K_{a1} ($mole L^{-1}$) is the first acid dissociation constant, R is
 311 the gas constant ($8.314 m^3 Pa K^{-1} mol^{-1}$), and T is temperature (K). The complete derivation for
 312 $\varepsilon(HA(aq))$ can be found in the SI of Guo et al. (2015).

313 Assuming unity activity coefficients, for dicarboxylic acids (H_2A , e.g., oxalic acid,

314 malonic acid, and maleic acid), the pH-dependence of the molar fraction of H₂A in the aqueous
315 phase ($\varepsilon(H_2A(aq))$) can eventually be simplified to the following equation (Nah et al., 2018):

$$316 \quad \varepsilon(H_2A(aq)) \cong \frac{H_{H_2A} W R T (10^{-pH} + K_{a1}) \times 0.987 \times 10^{-14}}{10^{-pH} + H_{H_2A} W R T (10^{-pH} + K_{a1}) \times 0.987 \times 10^{-14}}$$

317 where W is liquid water concentration ($\mu\text{g m}^{-3}$), H_{H_2A} (mole L⁻¹ atm⁻¹) is the Henry's law
318 constants for monocarboxylic acid, K_{a1} (mole L⁻¹) is the first acid dissociation constant, R is
319 the gas constant (8.314 m³ Pa K⁻¹ mol⁻¹), and T is temperature (K). The complete derivation for
320 $\varepsilon(H_2A(aq))$ can be found in the SI of Nah et al. (2018), which also includes discussions of the
321 assumptions made during the derivation process which will lead to the disappearance of the
322 second acid dissociation constant (K_{a2}) term during the process of simplifying the equation.

323

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