

1 **Applying Raman micro-spectroscopy to evaluate the effects of nutrient cations**  
2 **on alkane bioavailability to *Acinetobacter baylyi* ADP1**

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## 20 **Abstract**

21 Contamination with petroleum hydrocarbons, such as crude and mineral oils, can cause extensive  
22 damage to ecological systems. On oil contaminated sites, alkanes are major components; many  
23 indigenous bacteria exhibit the ability to access and/or degrade alkanes. However, their ability to  
24 do so is affected by external properties of the soil, including nutrient cations which can enhance  
25 or reduce their performance. This study therefore used Raman micro-spectroscopy techniques to  
26 study how nutrient cations affect alkanes' bioavailability to *Acinetobacter baylyi* ADP1 (a known  
27 degrader). Following treatment with Na, K, Mg and Ca at 10 mM, *A. baylyi* ADP1 was exposed to  
28 7 n-alkanes (decane, dodecane, tetradecane, hexadecane, nonadecane, eicosane and tetracosane)  
29 and 1 alkane mixture (mineral oil). Raman spectral alterations identified from multivariate analysis  
30 indicated that bacterial availability of alkanes varied with carbon chain lengths and additional  
31 cations altered the bacterial response to n-alkane molecules. Sodium significantly increased the  
32 bacterial affinity towards short-chain alkanes (decane and dodecane), and K and Mg enhanced the  
33 bioavailability of medium-chain alkanes (tetradecane and hexadecane). In contrast, the bacterial  
34 response was inhibited by Ca for all alkanes. Similar results were observed in the treatment of  
35 mineral oil. Our study employed a novel Raman spectral assay to offer a deep insight into how  
36 nutrient cations affect bioavailability of alkanes, suggesting nutrient cations present in natural  
37 environments can play a key role to influence the harmful effects of such contaminants and could  
38 be optimised to enhance a bioremediation strategy.

39

## 40 **1. Introduction**

41 Rapid industrial development has caused the increasing use of petroleum hydrocarbons and their  
42 derivatives, such as crude oil and mineral oil. From the 1960's, more than 40 large oil spill  
43 incidents have been reported, *e.g.*, the Deepwater Horizon oil spill in Gulf of Mexico, the Xingang  
44 oil spill in Dalian, and the Exxon Valdez oil spill in Prince William Sound.<sup>1-4</sup> The release of  
45 hydrocarbons into the environment can result in extensive damage to natural ecological systems  
46 and serious threats to human health.<sup>5-8</sup> Oil spill accidents and relevant contaminated sites have  
47 therefore received considerable attention in recent decades. Biosensor and bioremediation methods  
48 are deemed as an environmentally friendly strategy to monitor and remove petroleum  
49 hydrocarbons from the environment.<sup>9-13</sup> Since alkanes are the main components of hydrocarbon  
50 contamination, whole-cell bioreporters have been used to evaluate bioavailability and  
51 bioaccessibility. Some have even been applied practically in the field, but low reproducibility has  
52 limited their use for alkane monitoring.<sup>4,13</sup> Numerous bacterial species show high hydrocarbon  
53 affinity in natural environments and can have sensitive responses to the bioavailable fractions of  
54 relevant alkanes.<sup>14,15</sup> However, environmental physicochemical conditions and nutrient  
55 deficiencies can limit bacterial activities, which consequently affect their behaviour towards  
56 alkanes.<sup>16,17</sup> Because bacterial selection and accumulation is significantly related to the  
57 bioavailability of alkanes,<sup>18</sup> it is necessary to understand how nutrients influence bacterial  
58 behaviours.

59 At hydrocarbon contaminated sites, bacterial cells have to collect and/or accumulate their  
60 surrounding alkane molecules.<sup>18</sup> The bioavailable alkane affects the sensory systems of bacterial  
61 cells, including selection, accumulation and/or utilization of alkanes.<sup>19</sup> Briefly, bacteria's sensory  
62 system consists of methyl-accepting chemotaxis protein (MCP), histine kinase (CheA) and signal

63 transduction protein (CheW).<sup>20</sup> Alkane sensitive bacteria possess various MCP-encoding genes to  
64 interact with extracellular alkane molecules. In *Pseudomonas putida* GPo1, *alkN* gene has been  
65 found to encode an MCP for alkane collection.<sup>21</sup> The *tlpS* gene, located downstream of *alkB1*, is  
66 believed to encode the MCP in *Pseudomonas aeruginosa* PAO1 for hexadecane accumulation.<sup>22</sup>  
67 *Acinetobacter baylyi* ADP1 exhibits strong selection and accumulation towards alkanes.<sup>23-25</sup> The  
68 fimbriae on the membrane surface of *Acinetobacter* cells drive them to access alkane droplets to  
69 activate alkane hydroxylase AlkM for further degradation.<sup>23-25</sup> The emulsan-coding *wee* gene  
70 cluster in *Acinetobacter* contributes to the bacterial emulsification of alkane molecules when  
71 attaching on alkane droplets.<sup>26</sup> Uneven distributions of cations usually associate with alkane  
72 contamination, resulting in extra cationic strengths on the bioavailability of alkanes in bacterial  
73 cells. Therefore, this needs to be considered for its influence on bacterial accumulation of alkanes.  
74 Conventional methods that are used to study bacterial accumulation behaviours include swarm  
75 plates,<sup>27</sup> capillary assays,<sup>28</sup> temporal stimulation of tethered cells<sup>29</sup> and automated tracking of  
76 swimming cells.<sup>30</sup> They are useful methods to study bacterial movements towards specific  
77 chemicals, but require longstanding and labour-intensive pre-treatment work. Raman micro-  
78 spectroscopy is widely used in biological studies.<sup>31</sup> As a fast, reproducible and non-destructive  
79 approach providing vibrational information of biomolecules, it has been applied to investigate  
80 bacterial responses to their surrounding environments.<sup>32,33</sup> Spectral alterations in Raman peaks  
81 allow the measurement of cationic effects on bacterial selection and accumulation for alkane  
82 molecules.

83 To our knowledge, this is the first study to use Raman spectral assays to gather information on  
84 how nutrients affect the bioavailability of alkanes and alkane mixtures. We used Raman micro-  
85 spectroscopy to identify the influences of four cations (Na, K, Mg, Ca) on the selection and

86 accumulation of alkanes around or inside *A. baylyi* cells. We demonstrated that the bioavailable  
87 fraction of alkanes is highly specific to external cations on oil contaminated sites, and the stability  
88 of bacterial accumulation towards alkanes is measurable by Raman spectral assay. This method  
89 contributes to the deep understanding of changes in bacterial behaviours with different  
90 environmental conditions, and has the potential as a tool for the risk assessment of environmental  
91 contaminants.

## 92 **2. Materials and methods**

### 93 *2.1 Bacterial strain and growth conditions*

94 An alkane sensitive bacterial strain *Acinetobacter baylyi* ADP1 was used for all treatments in this  
95 study. This strain was inoculated in minimal medium with 20 mM sodium succinate as the sole  
96 carbon source, shaking at 150 rpm and 30°C for 16 h. To prepare 1.0 L mineral medium, 1.0 g of  
97 (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, 2.5 g of KH<sub>2</sub>PO<sub>4</sub>, 0.1 g of MgSO<sub>4</sub>·7H<sub>2</sub>O, 0.005 g of FeSO<sub>4</sub>·7H<sub>2</sub>O, 0.25 g of  
98 nitrilotriacetic acid, 0.55 g of NaOH and 1 mL of Bauchop and Elsdon solution were mixed well  
99 in 1.0 L deionized water and autoclaved.<sup>34</sup> Once harvested by centrifugation at 4000 rpm for 4 min,  
100 the cell pellets were washed three times with sterile deionized water and finally re-suspended in  
101 fresh mineral medium for use in the experiments. Unless stated otherwise, all chemicals in this  
102 study were of analytical grade and purchased from Sigma Aldrich (UK).

### 103 *2.2 Alkane exposure experiment*

104 Seven pure alkanes (decane, dodecane, tetradecane, hexadecane, nonadecane, eicosane and  
105 tetracosane), and mineral oil were selected as the classic alkane mixture. They were dissolved in  
106 dimethyl sulfoxide (DMSO) to prepare the 10 g/L stock solution. To investigate the impacts of Na<sup>+</sup>,  
107 K<sup>+</sup>, Mg<sup>2+</sup> and Ca<sup>2+</sup> on bacterial selections and accumulations towards alkanes, four cation-

108 providing chemicals (NaCl, KCl, MgCl<sub>2</sub> and CaCl<sub>2</sub>) were dissolved in mineral medium to prepare  
109 100 mM stock solutions, respectively.

110 Bacterial suspensions (9 mL) were mixed with 1 mL mineral medium or cation stock solution to  
111 reach a final cationic concentration of 10 mM. The alkane stock solution was mixed with bacterial  
112 suspensions with or without extra cations to reach a final concentration of 100 mg/L. All the  
113 samples were incubated at 30°C for 1 h, and further centrifuged at 5,000 rpm for 5 min to remove  
114 the medium. The cell pellets were re-suspended in deionized water and washed three times.

### 115 *2.3 Capillary assay of alkane chemotaxis*

116 To confirm bacterial selections and accumulations towards alkanes, capillary assays were  
117 performed based on the protocol of Li et al. with certain modifications.<sup>32</sup> Capillary tubes (internal  
118 diameter of 0.2 mm and length of 10 cm) were immersed into 1 mL of chemotaxis attractant stock  
119 solution (alkanes or mineral oil) for 10 min until the liquid was drawn up to ~1 cm of the length  
120 of the tube. The capillary was then inserted into the bacterial suspensions and incubated for 1 h at  
121 30°C.

122 Quantitative polymerase chain reaction (qPCR) was introduced to quantify the 16S rRNA copy  
123 numbers of chemotactic bacteria in triplicate. After 1-h incubation, the capillary tube was removed  
124 and the 1-cm exterior from the open end was directly plunged into qPCR buffer. The 10 µL qPCR  
125 buffer included 1 µL of primer 314F (5'-CCTACGGGNGGCWGCAG-3'), 1 µL of primer 802R  
126 (5'-TACNVGGGTATCTAATCC-3'), 3 µL molecular water and 5 µL iTaq™ Universal SYBR®  
127 Green Supermix (BioRad, USA). The qPCR thermos cycling program of 16S rRNA was: initial  
128 denaturation at 94°C for 3 min; 34 amplification cycles of 94°C for 45 s, 52°C for 45 s, 72°C for  
129 45 s, and fluorescence data acquisition at 80°C for 15 s. The standard curves were obtained with

130 serial dilutions of quantified plasmid DNA containing the fragment of 16S rRNA.<sup>35</sup>

#### 131 *2.4 Raman micro-spectroscopy measurement*

132 Prior to Raman micro-spectroscopy measurement, 10  $\mu$ L of washed cell pellets were transferred  
133 onto an aluminium foil coated slide and air-dried at room temperature in a fume cupboard. Raman  
134 spectra of all the samples were obtained using an InVia confocal micro-Raman system (Renishaw,  
135 Gloucestershire, UK). This system was equipped with a 100 mW 785 nm excitation laser diode.  
136 The entrance slit of the spectrometer was 50  $\mu$ m combined with a 1200 lines per mm ( $1 \text{ cm}^{-1}$  spatial  
137 resolution) diffraction grating, allowing the dispersion of Raman signals onto a Master Renishaw  
138 Pelletier cooled charged couple detector (CCD). Locations of sample detection were visualized via  
139 an attached microscope (Leica Microsystems, Milton Keynes, UK) with  $\times 50$  objective (0.75  
140 numerical aperture;  $\approx 1 \mu\text{m}$  spatial resolution). The Raman system was calibrated using a Renishaw  
141 silicon calibration source for wavenumber shifts before sample analysis. All sample spectra were  
142 obtained using 50% laser power (13 mW at sample), 10 s acquisition time, and one accumulation  
143 within a spectral range from 800 to 2000  $\text{cm}^{-1}$ . All treatments were carried out in triplicate, and at  
144 least twenty replicates were performed and analysed for each sample.

#### 145 *2.5 Computational analysis of Raman spectra*

146 Unless specifically stated otherwise, all the Raman spectral data were analyzed using the IRootLab  
147 toolbox for MatLab (version R2013b, MathWorks, USA).<sup>36</sup> Baseline correction and vector  
148 normalization was applied before principal component analysis (PCA) and linear discriminant  
149 analysis (LDA). PCA was employed to reduce the dimensionality of the multivariate data and  
150 allow the visualization of the natural variance within the data set. To accomplish inter-class  
151 separation and minimize intra-class differences, LDA was used to extract inter-category

152 discriminating features. The separation of individual spectral categories from negative control and  
153 pure alkane classes was measured by exporting PCA-derived data. Post-exposure to different types  
154 of alkanes or treated with different extra ions, the dispersion of individual Raman spectra to that  
155 of negative controls (no alkanes and no extra ions added) and pure alkane was calculated on the  
156 values of principal component 1 (PC1) and PC2, and visualized as dispersion indicator ( $D_I$ ) score  
157 plots (see the Supporting Information (SI) for calculation details). In  $D_I$  score plots, the increasing  
158  $D_I$  value between two categories is proportional to dissimilarity (i.e. higher values indicate greater  
159 dissimilarity from controls). The description and transformation of PCA data of each treatment  
160 were completed using Python (Version 3.0) (see the SI for the relevant code).

### 161 **3. Results and discussion**

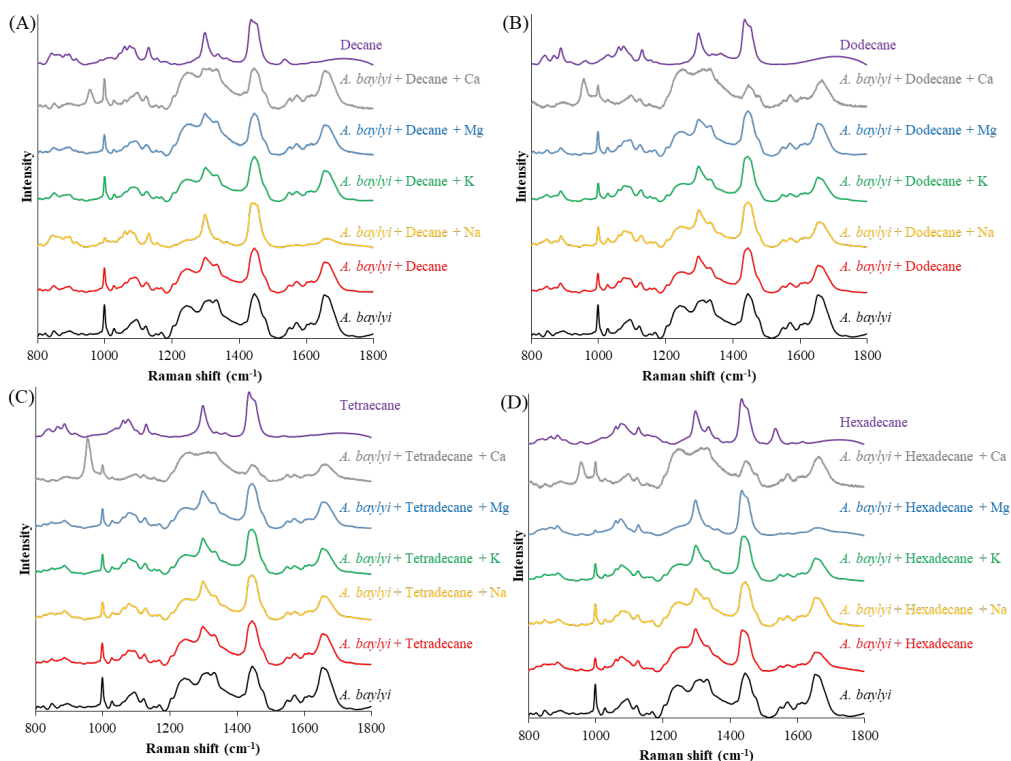
#### 162 *3.1 Raman spectral characterization of alkane exposed *A. baylyi**

163 Raman spectra characterized the peaks of pure alkanes and alkane mixture. Since decane, dodecane,  
164 tetradecane, hexadecane, nonadecane, eicosane and tetracosane were from the group of linear  
165 alkanes, major peaks in their Raman spectra were caused by similar chemical bonds (Figure1). In  
166 Figure S1 and Table 1, notable peaks observed in all Raman spectra of n-alkane molecules included  
167  $890\text{ cm}^{-1}$  ( $\text{CH}_3$  rock),  $1060\text{ cm}^{-1}$  (C–C symmetric stretching),  $1079\text{ cm}^{-1}$  (C–C stretching),  $1136$   
168  $\text{cm}^{-1}$  ( $\text{CH}_2$  stretching),  $1296\text{ cm}^{-1}$  ( $\text{CH}_2$  twisting) and  $1436\text{ cm}^{-1}$  ( $\text{CH}_2$  bending).<sup>37</sup> Although Raman  
169 spectra of pure n-alkane molecules were similar, spectra of decane and hexadecane showed  
170 specific peaks at  $1535\text{ cm}^{-1}$ , which might be attributed to the linkages of C and H deformation.<sup>38</sup>  
171 In the spectrum of dodecane, two distinct peaks at  $961$  and  $1033\text{ cm}^{-1}$  were assigned to  $\text{CH}_2$  rock.<sup>39</sup>  
172 Because of the solid state of nonadecane, eicosane and tetracosane,  $\text{CH}_2$  bending band shifted to  
173  $1440$  and  $1463\text{ cm}^{-1}$ , indicating an increase in intermolecular interactions and vibrational motion.<sup>40</sup>  
174 In the alkane mixture, Raman spectra of mineral oil showed similar peaks at  $890$ ,  $1060$ ,  $1136$  and

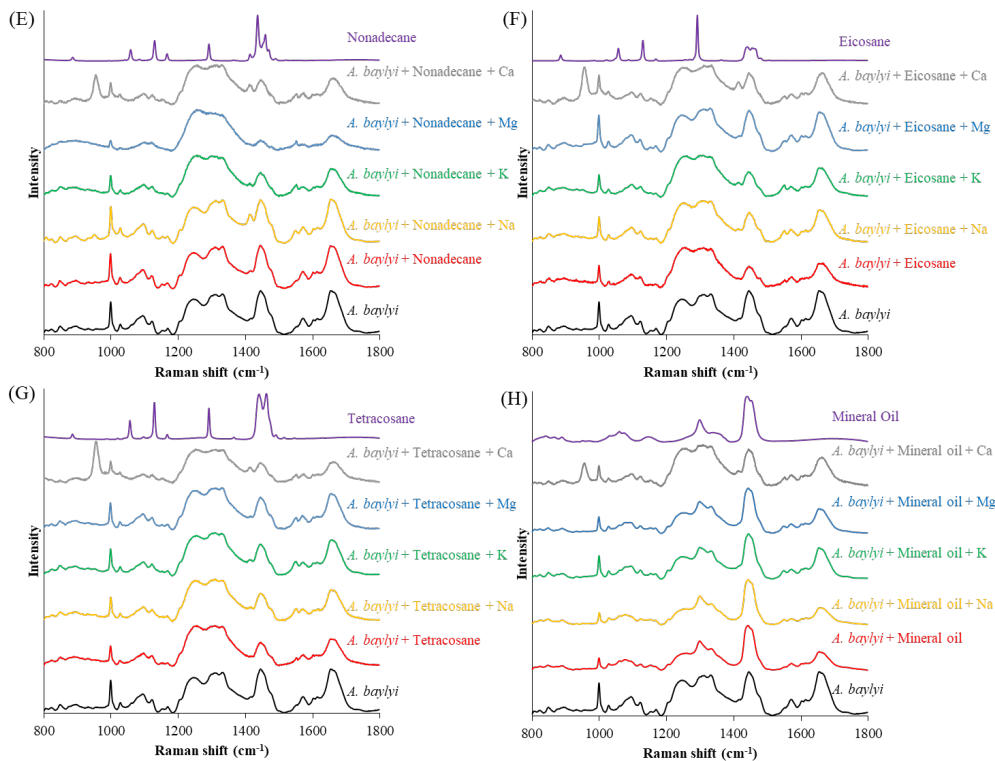


175 1436 cm<sup>-1</sup> (Figure 1H).

176 To study spectral alterations of alkane exposed bacteria, Raman spectra of original *A. baylyi* were  
177 measured as control. Raman peaks at 1238 and 1311 cm<sup>-1</sup> were assigned to amino acids, and the  
178 band of 1441 cm<sup>-1</sup> was generated from the glycine in bacterial cells (Table 1).<sup>41</sup> The band at 1002  
179 cm<sup>-1</sup> represents phenylalanine, and the peak at 1663 cm<sup>-1</sup> was attributed to the protein on cell  
180 membranes.<sup>42</sup> *Acinetobacter baylyi* ADP1 are ubiquitous bacteria observed in natural soil  
181 environments. They are frequently found to be able to utilize alkanes with carbon lengths ranging  
182 from 12 up to 36.<sup>13</sup> Previous work has shown the strong accumulation towards dodecane and  
183 tetradecane in *A. baylyi* cells.<sup>35</sup> As a consequence, Raman micro-spectroscopy is able to test the  
184 bacterial selections and accumulations towards short-, medium- and long-chain n-alkanes.



185



186

187 **Figure 1.** Raman spectra of *A. baylyi* towards decane (A), dodecane (B), tetradecane (C),  
 188 hexadecane (D), nonadecane (E), eicosane (F), tetracosane (G), and mineral oil (H). Spectra in  
 189 black are for pure bacterial cells, red for the exposure of alkane, yellow for the treatment with extra  
 190 sodium, green for the extra potassium, blue for the extra magnesium, grey for the extra calcium  
 191 and purple for pure alkane.

192 Post-exposure to decane, dodecane, tetradecane, hexadecane and mineral oil, Raman spectral  
 193 alterations at 1296 and 1436  $\text{cm}^{-1}$  generated from alkanes indicated their accumulation  
 194 around/inside bacterial cells. However, no significant Raman alterations from alkanes were found  
 195 in the spectra of nonadecane, eicosane or tetracosane exposures. Sensitive sensing behaviours of  
 196 *A. baylyi* enabled the accumulation of alkane molecules in/on bacterial cells towards decane to  
 197 hexadecane, whereas no positive response was found for nonadecane, eicosane and tetracosane.  
 198 *Acinetobacter* can move towards the area where alkane molecules were available.<sup>12</sup> Once adhering

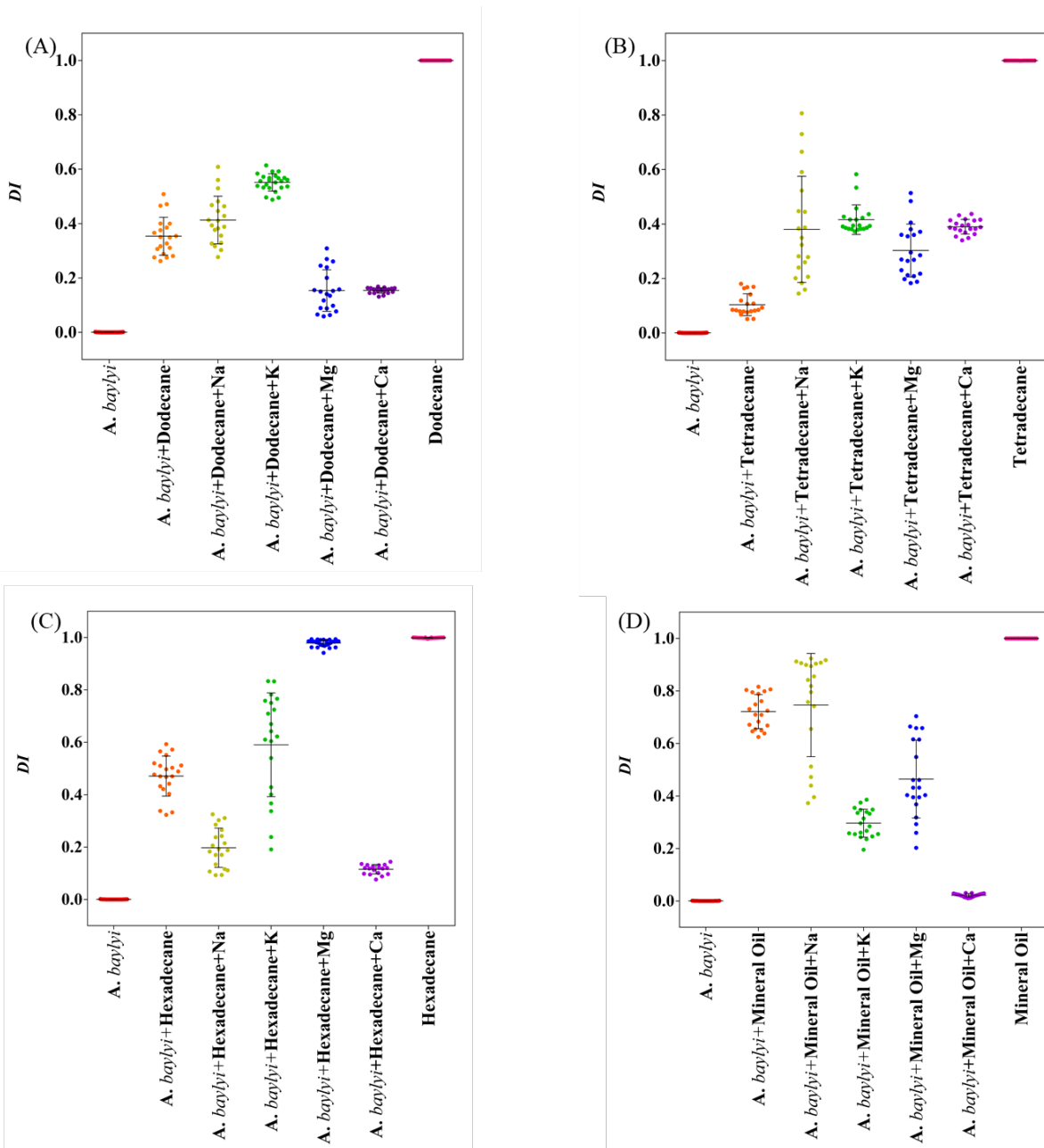
199 to the alkane droplets, *Acinetobacter* excretes bio-surfactants to emulsify alkane molecules on  
 200 membranes within a second, contributing to the massive accumulation of molecules on or inside  
 201 bacterial cells.<sup>43-45</sup>

202 In Figure S2, the accumulation of 16S rRNA copies in capillary tubes of alkane were less than 19  
 203 carbons range from  $8.8 \times 10^9$  to  $2.33 \times 10^{11}$  copies per capillary, significantly higher than those in  
 204 the treatments of nonadecane, eicosane and tetracosane. Higher copies illustrated the more  
 205 bioavailable alkanes in capillary tubes for *A. baylyi* to collect. The results from capillary assays  
 206 confirmed the bioavailability of alkane was related to the bacterial accumulation.

207 **Table 1.** Variations and assignments of Raman bands of alkanes and *A. baylyi*

Band (cm <sup>-1</sup> )	Tentative bands assignment	Origin
890	Terminal methyl CH <sub>3</sub> rock	All alkanes
1060	C-C symmetric stretch	All alkanes
1079	C-C stretch	All alkanes
1136	CH <sub>2</sub> stretch	All alkanes
1296	CH <sub>2</sub> twist	All alkanes
1436	CH <sub>2</sub> bend	All alkanes
1535	C-H deformation	Decane and Hexadecane
961	CH <sub>2</sub> rock	Dodecane
1033	CH <sub>2</sub> rock	Dodecane

1440	CH <sub>2</sub> bend shift	Nonadecane, eicosane and tetracosane
1463	CH <sub>2</sub> bend shift	Nonadecane, eicosane and tetracosane
1002	Benzene ring breathing	Bacterial phenylalanine
1238	/	Bacterial amino acids
1311	/	Bacterial amino acids
1441	Deformations of CH <sub>2</sub>	Bacterial glycine
1663	C=C stretching	Bacterial proteins



209  
 210 **Figure 2.** *DI* scores of the effects of four nutrient cations on bacterial selections and accumulations  
 211 of *A. baylyi* towards dodecane (A), tetradecane (B), hexadecane (C), and mineral oil (D)

212

213 Post-exposed to decane in each treatment, Raman spectra showed distinguishable peak alterations,  
214 resulting in significant differences in  $D_I$  score plots (Figure S3A). In original MM, the 0.177  $D_I$   
215 value indicated the moderate bacterial accumulation towards decane in *A. baylyi*. In the treatment  
216 with Na,  $D_I$  values were close to that of pure decane, demonstrating the remarkable enhancement  
217 for its accumulation in *A. baylyi*. In contrast, the low  $D_I$  values found in K (0.114), Mg (0.066) and  
218 Ca (0.012) treatments indicated the decreased bioavailability of decane. The accumulation of  
219 decane in *A. baylyi* generated relevant Raman alterations at 1296 and 1436  $\text{cm}^{-1}$  in corresponding  
220 spectra (Figure 1A). The outer-membrane protein OmpS is responsible for the signal transport of  
221 short-chain alkanes such as octane, nonane and decane in bacterial cells.<sup>20</sup> Extra Na in the ambient  
222 environment around bacteria may accelerate the signal transport inside cytoplasm, or enhance the  
223 emulsification of decane molecules on the cell membrane to increase its bioavailability. Post-  
224 exposed to dodecane,  $D_I$  values of Na (0.501) or K (0.583) were higher than those in the original  
225 MM (0.423), showing the increased bioavailability in *A. baylyi* with these two cations (Figure 2A).  
226 It is reported that *A. baylyi* is able to grow with dodecane as the sole carbon source in liquid  
227 medium.<sup>46</sup> The strong bacterial affinity of dodecane results in the fast molecules' transport through  
228 the cell of *A. baylyi*, which is well characterized by Raman spectra.<sup>32</sup>

229 From the results in Figure 2B, all tested cations improved the bioavailability of tetradecane, and  
230 peak alterations at 1060, 1079 and 1296  $\text{cm}^{-1}$  observed in Raman spectra also proved the  
231 accumulation of tetradecane in *A. baylyi* cells (Figure 1C). The bioavailable fraction of tetradecane  
232 affects the signal transmitter protein, OmpS, and the alkane uptake protein, OmpW in bacterial  
233 cells to change their behaviours.<sup>20,47</sup> Extra cations possibly induced more available alkane  
234 molecules for bacteria to enhance their upregulations of relevant proteins. The  $D_I$  value of  
235 bioaccumulated hexadecane in normal MM was 0.547 (Figure 2C), demonstrating its stronger

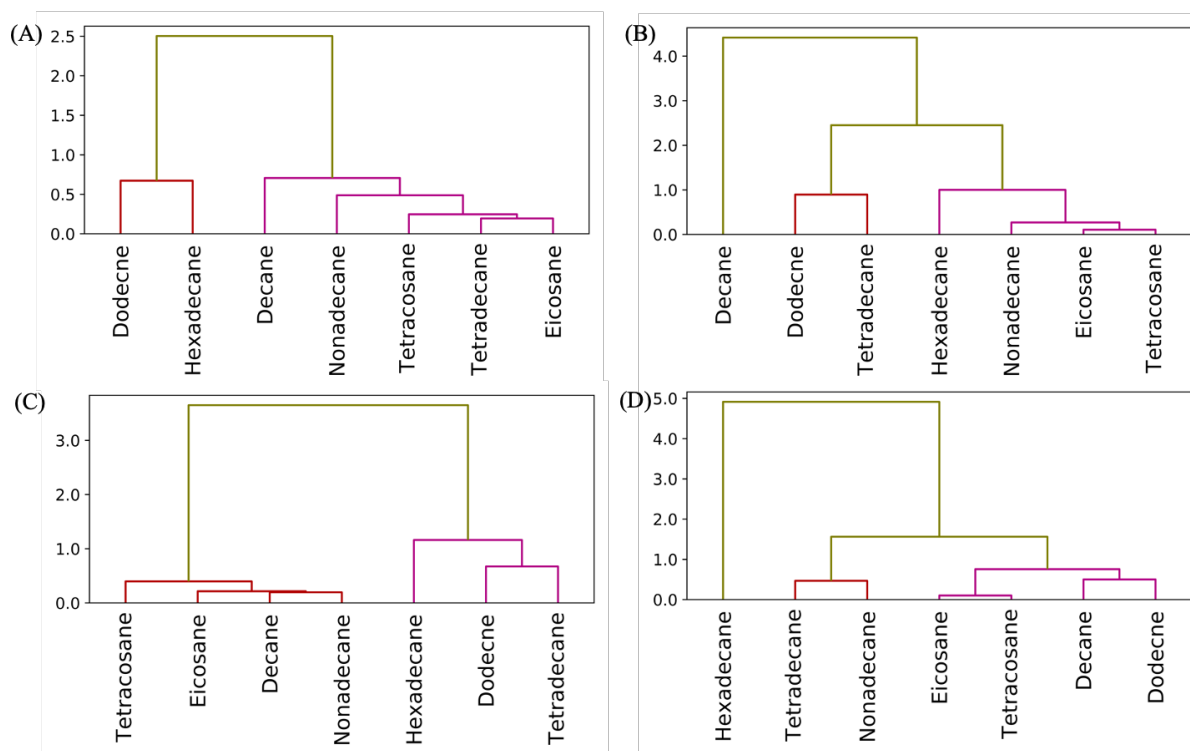
236 bacterial affinity compared to tetradecane. In four cation treatments,  $D_I$  values were in the order:  
237 Mg (0.991) > K (0.788) > Na (0.272) > Ca (0.133). The bioavailability of hexadecane for *A. baylyi*  
238 was increased by K and Mg, but decreased by Na and Ca, compared to that in normal MM (Figure  
239 2C). Due to the influences of cations, the incremental/reduced available fraction of hexadecane  
240 regulates its accumulation in *Acinetobacter*,<sup>48,49</sup> and possibly affects the expression of relevant  
241 sensory proteins at cellular level.<sup>50</sup>

242 In normal and extra Na equipped MM, the accumulation of *A. baylyi* towards nonadecane was  
243 weak, proved by the relatively low  $D_I$  values, 0.001 and 0.002, respectively (Figure S3B). The  
244 presence of extra K, Mg and Ca slightly increased the bioavailability of nonadecane, but no major  
245 alterations were found in relevant Raman spectra (Figure 1E and 2E), indicating that only small  
246 amounts of nonadecane molecules were accumulated by *A. baylyi*. Eicosane and tetracosane had  
247 similar  $D_I$  values in normal medium (Figure S3C and S3D), demonstrating their weak  
248 bioavailability. The four tested cations were limited in changing the available fractions of eicosane  
249 and tetracosane, and their effects on the accumulation behaviour of *A. baylyi* were slight. Although  
250 the uptake and degradation of long-chain alkanes in *Acinetobacter* is well-documented,<sup>51</sup> further  
251 studies are required to understand how their bioavailability and bioaccumulation in different  
252 environmental conditions impact ecological systems. Different from short and medium chain  
253 alkanes, nonadecane, eicosane and tetracosane are waxy solids at room temperature and  
254 atmospheric pressure. In the liquid medium, the low solubility of long chain alkane limits the  
255 interaction with cations. Therefore, compared with liquid alkanes, their bioavailable fractions  
256 remain at a relatively low level with extra cations.

257 The  $D_I$  results in Figure 2D indicated the strong bioavailability of mineral oil. The similar  $D_I$  value  
258 found in Na treatment demonstrated the weak effect on its accumulation in *A. baylyi* cells.

259 Although the bioavailability was reduced with K and Mg, the Raman peak at 1296  $\text{cm}^{-1}$  ( $\text{CH}_2$   
 260 twisting of mineral oil molecule) was observed in relevant spectra, showing its substantial  
 261 bioaccumulation (Figure 1G). However, the extra Ca significantly reduced the bioavailability of  
 262 mineral oil, leading to the unchanged Raman spectra pre- and post-exposed to mineral oil (Figure  
 263 1G and 2D). As a mixture of different alkanes, mineral oil has strong bioavailability for bacteria.  
 264 Also, extra cations can affect various components in mineral oil, resulting in the instability of  
 265 bioavailable mineral oil. Therefore, by evaluating the stability of bioavailable hydrocarbons, the  
 266 Raman spectral assay provides a tool for the risk assessment of oil contamination.

### 267 3.3 Impacts of cations on alkane bioaccumulation



268  
 269  
 270 **Figure 3.** Hierarchy clustering analysis for the bioaccumulation of *A. baylyi* towards alkanes in  
 271 the normal medium (A), with extra Na (B), K (C), and Mg (D). Dendrograms represent alkanes



272 cluster, and y axis indicates the differences in bioavailability of alkanes (high value demonstrates  
273 great dissimilarity).

274 From the results in Figure 3, extra cations exhibited different influences on the bioavailability of  
275 short chain (decane and dodecane), medium chain (tetradecane, hexadecane and nonadecane), and  
276 long chain alkane (eicosane and tetracosane). In the normal medium, dodecane and hexadecane  
277 from one cluster indicated their adequate available fractions for bacteria to access (Figure 3A), but  
278 the noticeable variations demonstrated the diverse bacterial accumulations towards these two  
279 alkanes (Figure S4A). The strong selections and accumulations towards dodecane and hexadecane  
280 are attributed to the emulsan produced from *Acinetobacter* to improve the bacterial emulsification,  
281 and then increase the accumulation of alkane molecules.<sup>26,52</sup>

282 In the Na treatment, the highest impact factor (0.89) and tiny variations of decane-exposed  
283 bacterial cells reflected its stable bioavailable fraction for bacterial cells to collect (Figure 4 and  
284 Figure S4B). The sole cluster of decane in the dendrogram also indicated the specific effect of  
285 sodium on the bioaccumulation of this alkane (Figure 3B). Dodecane and tetradecane fell in one  
286 cluster, showing the positive response of *A. baylyi* towards these two alkanes (Figure 3B). Owing  
287 to the high halotolerant ability of *A. baylyi*, we believe that this bacterial strain is sensitive to the  
288 change in the concentration of Na in its surrounding environments.<sup>34</sup> The increase in external Na  
289 concentration affects the bacterial adherence and emulsification ability, leading to the uneven  
290 bioaccumulation of alkane molecules on/inside cell membranes,<sup>53</sup> which is also indicated by the  
291 significant variations of dodecane, tetradecane and hexadecane.

292 Alkane clusters in Figure 3C illustrated two different ways that K influenced the bioavailability of  
293 alkanes. The positive effect of extra K on bacterial selections and accumulations towards dodecane,  
294 tetradecane and hexadecane contributed to high  $D_I$  values: 0.583, 0.416 and 0.788, respectively

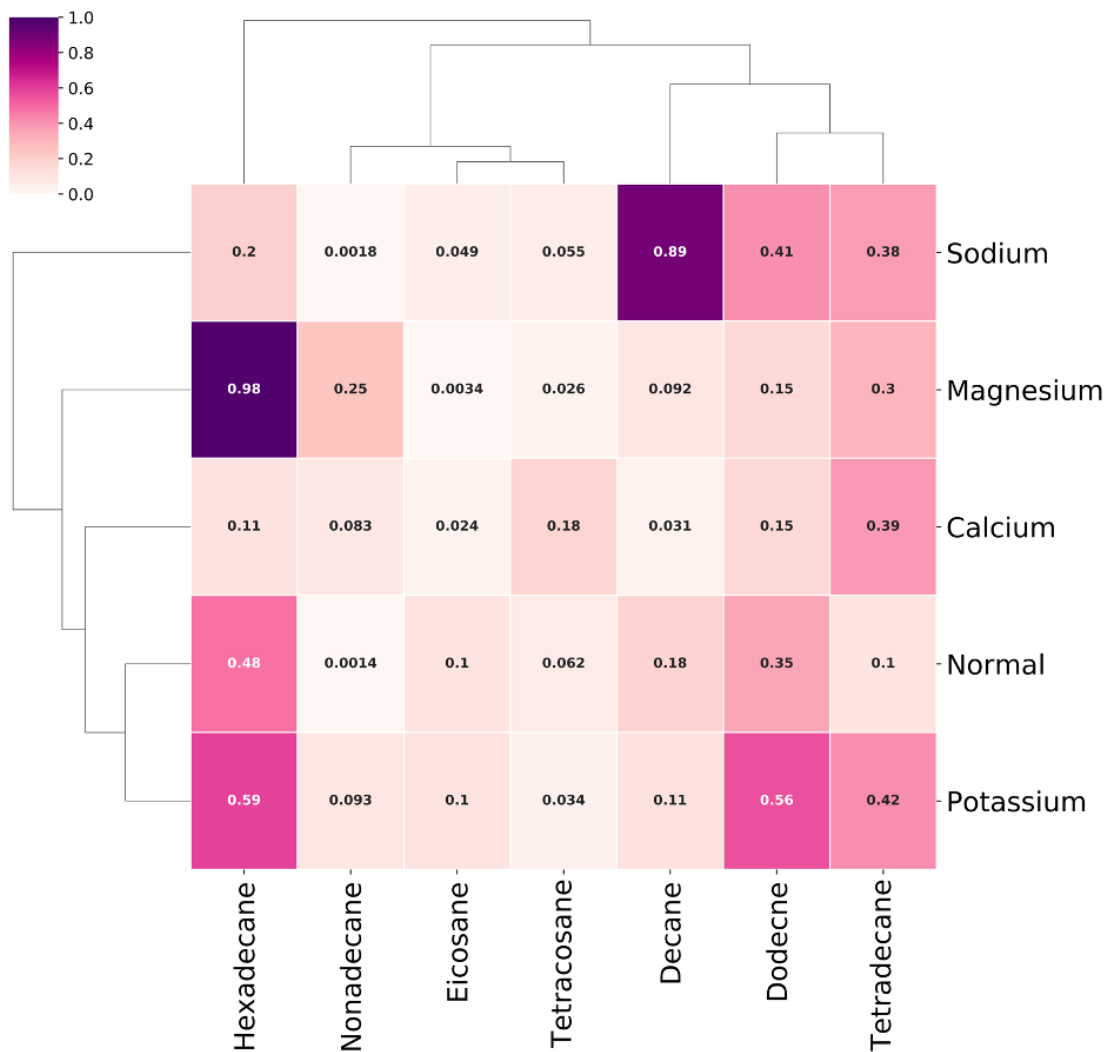
295 (Figure S4C). The bioavailability of the other 4 tested alkanes was weakly related to the impact of  
296 K, proved by the factor shown in Figure 5. The external K enhanced the bacterial accumulation of  
297 medium-chain n-alkanes,<sup>54</sup> and improved the transfer of information signals among bacterial cells  
298 to collect available alkane molecules.<sup>55</sup> The bioaccumulation of dodecane by *A. baylyi* increased  
299 smoothly, but bioavailable tetradecane and hexadecane molecules were not well-distributed in the  
300 medium with extra K, as shown by the noticeable variations of these two alkanes. As a  
301 consequence, K is essential for bacteria to accumulate alkane molecules, and can also influence  
302 the heterogeneity of bioavailable alkanes' distribution.

303 The dendrogram illustrated the positive relation of medium-chain alkane including dodecane,  
304 tetradecane and hexadecane to Mg (Figure 3D). The highest  $D_I$  score was from hexadecane,  
305 showing its strong bioaccumulation in *A. baylyi*, and the small variation proves that extra Mg was  
306 able to stabilize the bioavailability of this alkane. The bioavailability of tetradecane and  
307 nonadecane was also strongly correlated to the effect from external Mg (Figure 4). Extracellular  
308 Mg stabilized the bioaccumulation of *A. baylyi* towards tetradecane, hexadecane and  
309 nonadecane.<sup>56</sup> However,  $D_I$  values in Figure S4D showed that the impacts of Mg on the  
310 bioavailability were not uniform for all n-alkanes. Hence, we deduce that the transportation  
311 proteins of hexadecane in *A. baylyi* are more likely to interact with Mg than that of other n-alkanes.

312 The low  $D_I$  values indicated that extra Ca significantly reduce the bioavailability of alkanes (Figure  
313 S4E). The limited bacterial accumulations towards n-alkanes also resulted in the low impact factor  
314 of Ca in Figure 5. A more than 100 nM concentration of extracellular Ca can prohibit bacterial  
315 movements,<sup>57</sup> and bacterial cells tend to aggregate by interacting with the anionic components in  
316 bacterial cells.<sup>58</sup> Successful bacterial accumulation requires the efficient movement towards  
317 specific molecules, but extra Ca disables such ability of bacteria. External Ca possibly blocks

318 transferring channels of alkanes' signals in bacterial cells, leading to the inactive bioaccumulation  
319 of them.

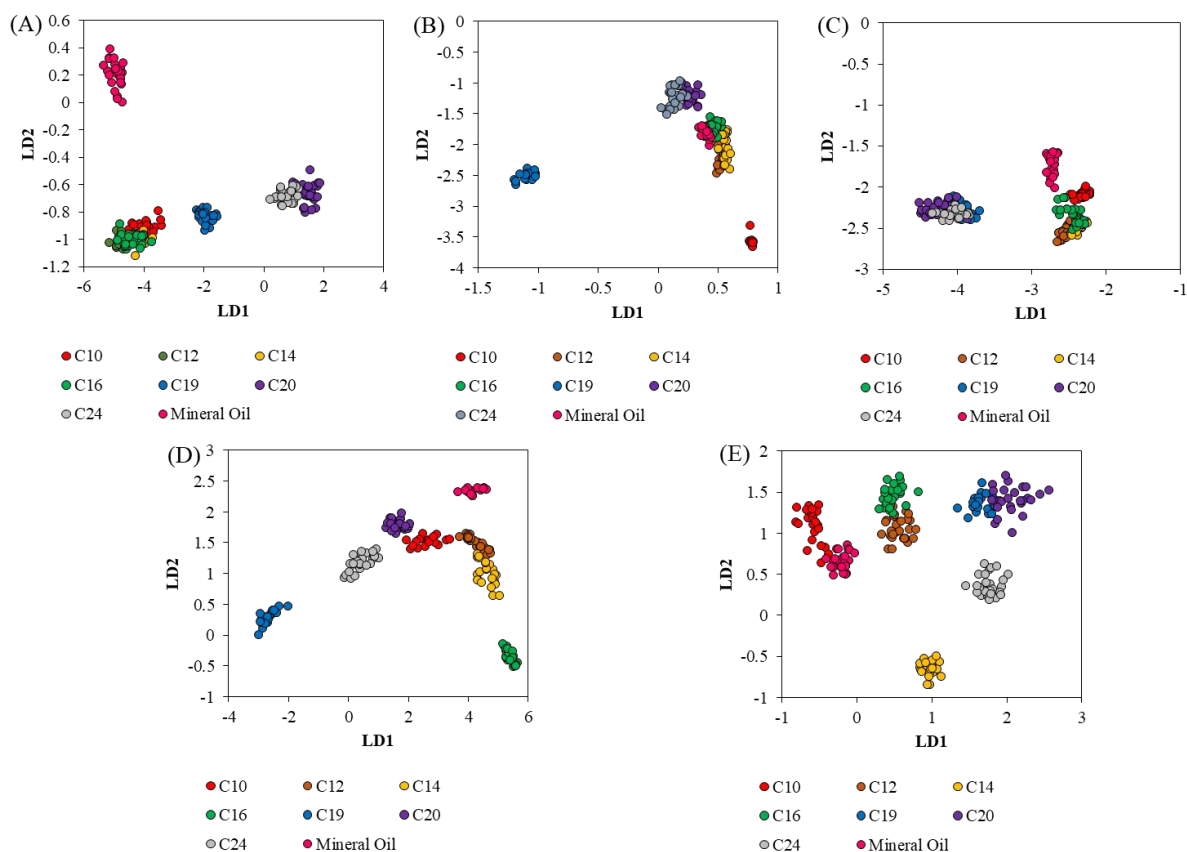
320 We investigated impact factors of cations that account for the alkanes' bioaccumulation, and  
321 compared them with those in the normal medium. The heatmap from hierarchy clustering analysis  
322 (Figure 5), shows the specific accumulation behaviour of *A. baylyi* towards alkanes is attributed to  
323 different impacts from each cation. The bioavailability of decane and hexadecane is highly  
324 sensitive to extra Na and Mg, respectively. Bioavailable fractions of dode-, tetra- and hexadecane  
325 are increased for bacteria to accumulate with external K. Because of cationic impacts, bacterial  
326 activities cannot remain at stable levels, especially on oil contaminated sites that involve  
327 complicated physicochemical conditions. Therefore, with the help of Raman micro-spectroscopy,  
328 we were able to visualize the effects of cations on bacteria, to uncover the underlying limitations  
329 to bacterial remediation of oil contaminated sites. Although further studies are required before  
330 applying this instrument in real environmental conditions, this method has already provided  
331 positive results in this study, to indicate it can help improve bioremediation of oil contamination.



332

333 **Figure 4.** Cluster heatmap for the accumulation behaviours of *A. baylyi* with different cations. The  
 334 impact factor of the cation was characterized into five quartiles, and is illustrated with intensity.  
 335 The dendrogram on the left represents cation clusters, and the one on the top refers to the alkane  
 336 clusters.

337 3.4 Cationic effects on bacterial accumulation in alkane mixture



338  
 339 **Figure 5.** LDA plots of alkane chemotactic selection of *A. baylyi* in alkane mixture (mineral oil)  
 340 with no extra ions (A), and in the presence of Na (B), K (C), Mg (D), and Ca (E)

341 The PCA-LDA of Raman spectra was performed to determine the available components in the  
 342 mineral oil for bacterial cells to interact with. In normal MM, the majority of alkane molecules  
 343 collected by *A. baylyi* in mineral oil were mainly short- and medium-chain alkanes, including  
 344 decane, dodecane, tetradecane and hexadecane (Figure 5A). Although mineral oil consists of  
 345 alkane molecules from C<sub>10</sub> to C<sub>35</sub>,<sup>59</sup> the major components in its bioavailable fraction are short-  
 346 and medium-chain n-alkanes.

347 Extra Na enabled the LDA group of mineral oil aggregates with dodecane, tetradecane and

348 hexadecane in LDA plot (Figure 5B), indicating the increased bacterial accumulation of these three  
349 alkanes in alkane mixture.<sup>53</sup>  $D_I$  scores also confirmed that extra Na prompts the bioavailability of  
350 carbon lengths of 12, 14 and 16. The LD1 values of Raman spectra indicated that external K  
351 enabled *A. baylyi* to capture large amounts of decane and hexadecane molecules in mineral oil  
352 (Figure 5C). Tetradecane exhibited strong affinity to *A. baylyi* with extra K as well. Hence, K may  
353 accelerate the movement of *A. baylyi* towards decane and hexadecane molecules, but reduce the  
354 accumulation of other n-alkanes in mineral oil.<sup>54</sup> The close distance of LDA groups indicated a  
355 number of dodecane molecules accumulated by *A. baylyi* in mineral oil with extra Mg (Figure 5D).  
356 Since extra Mg is able to stabilize the phosphorylated CheY, it can assist bacterial cells to access  
357 chemical molecules faster in alkane mixture.<sup>60</sup> From the LDA in Figure 5E, Ca weakened bacterial  
358 accumulation behaviours towards alkane molecules in mineral oil. Unlike the LDA plot of the  
359 normal medium, separated LDA groups of pure n-alkanes demonstrated the limited bacterial  
360 movement and reduce bioavailability of mineral oil due to extra Ca in aquatic environments.

361 Mineral oil is a complex alkane mixture comprising saturated n-alkanes and aromatic  
362 hydrocarbons.<sup>59</sup> Diverse cations challenge the bacterial accumulation of alkane molecules and  
363 affect their bioavailability, leading to complex impacts on ecological systems.<sup>61,62</sup> Therefore, the  
364 impacts of cations on the bioavailability of alkanes are significant for the treatment of oil  
365 contamination. Raman micro-spectroscopy has been used successfully to measure chemotaxis-  
366 related behaviours of *A. baylyi* towards hydrocarbons.<sup>32</sup> This application confirms that extra  
367 cations can change the dominant components in alkane mixture, thereby leading to different  
368 accumulation behaviours of bacteria towards alkane molecules in alkane mixtures.

369 This study applied Raman micro-spectroscopy to investigate the impacts of nutrient cations on the  
370 bioavailability of alkanes and alkane mixture in the liquid phase. To the best of our knowledge,

371 this is the first study to use Raman spectral methods to show the bacterial selections and  
372 accumulations of alkanes under different environmental conditions. The cation-alkane patterns  
373 observed the unstable/variable bioavailability of alkanes in the presence of extra cations, and  
374 differentiated bacterial accumulation behaviours in complex environments. Varying  
375 physicochemical conditions disturb the stable response of bacteria towards hydrocarbons.  
376 Therefore, this study provides a new perspective and method to gain information on the change in  
377 bioavailability of alkanes for the potential treatment of oil contamination incidents. Further work  
378 is required to elucidate the combining effects of nutrients for bacterial accumulation of  
379 hydrocarbons in environmental samples from oil-contaminated sites. In the future, this Raman  
380 spectral assay is helpful to evaluate the bioavailability and biotoxicity of environmental pollutants  
381 in natural ecological systems.

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