

1 **Development and validation of an *in situ* high-resolution**  
2 **technique for measuring antibiotics in sediments**

3 Yanying Li<sup>a,b</sup>, Qiuyu Rong<sup>c</sup>, Chao Han<sup>d</sup>, Hanbing Li<sup>e</sup>, Jun Luo<sup>b</sup>, Liying Yan<sup>b</sup>, Degao  
4 Wang<sup>a</sup>, Kevin C. Jones<sup>b,c,\*</sup> and Hao Zhang<sup>c,\*</sup>

5 <sup>a</sup> College of Environmental Science and Engineering, Dalian Maritime University,  
6 Dalian, Liaoning 116023, P. R. China

7 <sup>b</sup> State Key Laboratory of Pollution Control and Resource Reuse, School of the  
8 Environment, Nanjing University, Nanjing, Jiangsu 210023, P. R. China

9 <sup>c</sup> Lancaster Environment Centre, Lancaster University, Lancaster, LA1 4YQ, U.K.

10 <sup>d</sup> Nanjing Institute of Geography & Limnology, Chinese Academy of Sciences,  
11 Nanjing, Jiangsu 210008, P. R. China

12 <sup>e</sup> Department of Environmental Science, Faculty of Environment and Life, Beijing  
13 University of Technology, Beijing 100124, China

14

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16 \* corresponding authors

17 E-mail: [h.zhang@lancaster.ac.uk](mailto:h.zhang@lancaster.ac.uk); Tel: +44 1524 593899. (Hao Zhang)

18 [k.c.jones@lancaster.ac.uk](mailto:k.c.jones@lancaster.ac.uk); Tel: +44 1524 510230. (Kevin C. Jones)

19

20 **Abstract**

21 Important biogeochemical processes occur in sediments at fine scales. Sampling  
22 techniques capable of yielding information with high resolution are therefore needed to  
23 investigate chemical distributions and fluxes and to elucidate key processes affecting  
24 chemical fates. In this study, a high-resolution diffusive gradients in thin-films (DGT)  
25 technique was systematically developed and tested in a controlled sediment system to  
26 measure organic contaminants, antibiotics, for the first time. The DGT probe was used  
27 to resolve compound distributions at the mm scale. It also reflected the fluxes from the  
28 sediment pore-water and remobilization from the solid phase, providing more dynamic  
29 information. Through the fine scale detection, a reduction of re-supply was observed  
30 over time, which was concentration and location dependent. Compared to the Rhizon  
31 sampling method, antibiotic concentrations obtained by DGT probes were less than the  
32 pore-water concentrations, as DGT measures the labile fraction of the compounds. The  
33 DGT probe was also tested on an intact sediment core sampled from a lake in China  
34 and used to measure the distribution of labile antibiotics with depth in the core at the  
35 mm scale.

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37 **Key words:** antibiotics, sediment profile, high-resolution, fluxes

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39 **Highlights:**

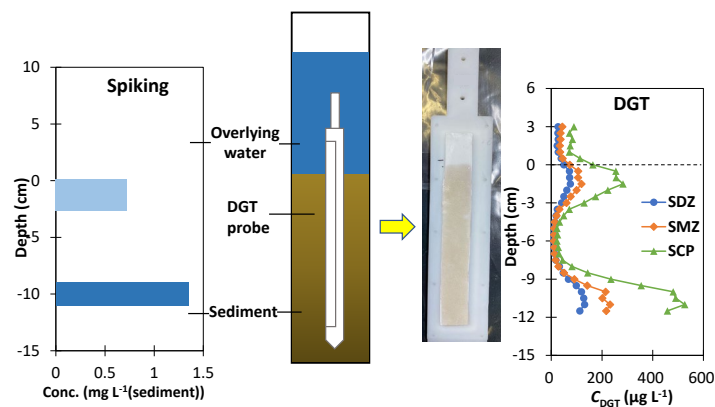
40 An new *in situ* high resolution sampling probe using DGT technique for studying  
41 antibiotics was developed and tested in controlled sediment systems

42 The resolving power was at the mm scale, better than the Rhizon sampler and  
43 conventional slicing.

44 Vertical distribution of antibiotics in sediments and their transport were determined

45 The DGT probe was applied to detect antibiotics in an intact sediment core.

46 **Graphical abstract**



47

## 48 1. Introduction

49 Sediments can be both sinks and secondary sources of organic contaminants to the  
 50 water column. Sorption of compounds onto settling particles can transport them to the  
 51 bottom sediments, and as compounds become incorporated into sediments and subject  
 52 to burial, they are removed from the water column [1]. However, if environmental  
 53 conditions change (e.g. following seasonal redox changes, or physical disturbance of  
 54 the sediments), sediment bound compounds can be released back into the overlying  
 55 water [2]. Removal/re-release of chemical compounds at the sediment-water interface  
 56 and in surface layers of the sediments can be dynamic processes in space and time [3].  
 57 Once incorporated into deposited sediments, the vertical concentration profiles can  
 58 provide information on the input history of contaminants and yield information on the  
 59 *in situ* degradation of the contaminants [4].

60 The distribution of contaminants (both vertical and horizontal) in the sediments is often  
 61 considered to be heterogeneous. Discrete particles, such as phytoplankton and faecal  
 62 pellets, may act as ‘hot spots’ in sediments [5]. Physical and chemical properties of  
 63 sediments – such as redox, pH and organic carbon vary over fine scales, which can  
 64 affect chemical speciation, microbial activity and decomposition processes [6].

65 Therefore, to understand mechanistic aspects of the behaviour of contaminants taking  
66 place in the sediments, it is necessary to study biogeochemical processes at the  
67 appropriate fine scale. This requires a high-resolution sampling technique; given how  
68 quickly sediment conditions can change following disturbance or sampling, it is also  
69 essential that the high-resolution sampling should be performed *in situ* with minimal  
70 disturbance.

71 Antibiotics are widely used in human therapy, animal husbandry and aquaculture [7].  
72 They are not completely absorbed by the body, with up to 80% of the administered  
73 antibiotics excreted as original parent compound or metabolites through feces and urine  
74 of patients and livestock into water bodies and soils – then ultimately reaching  
75 sediments of aquatic systems [8]. A large fraction of antibiotics used in aquaculture also  
76 enter waterbodies and reach sediments directly. Antibiotics in the environment can  
77 inhibit the growth of some beneficial microorganisms, change the microbial  
78 communities and affect metabolism in ecosystems [9]. Additionally, there is concern  
79 about the widespread occurrence of antibiotic residues in the environment, with the  
80 potential for development and transfer of resistance, leading to the threat to humans  
81 [10]. Antibiotics have been detected in surface waters, ground waters, soils and  
82 sediments, up to 100s  $\mu\text{g L}^{-1}$  or  $\mu\text{g kg}^{-1}$  [11-14]. In sediments, concentrations in the  
83 range of 10s – 10000s  $\mu\text{g kg}^{-1}$  have been reported around the world [15-17]. There are  
84 uncertainties over the persistence, potential for re-mobilisation and re-release, and  
85 bioavailability of antibiotics in sediments. Therefore, tools and studies to investigate  
86 the distribution and behavior of antibiotics in sediments are needed.

87 The conventional methods to study the distribution of antibiotics in sediments are  
88 centrifugation to remove the pore-water or slicing the sediment core followed by  
89 organic solvent extraction of the residues. For example, Li et al. [18] sliced sediment  
90 cores at 1 cm intervals, then used the traditional extraction approach. The spatial  
91 heterogeneity of contaminants at the micro-scale becomes lost with such methods.  
92 These commonly used *ex situ* methods disturb the original sediment environment,  
93 including the redox conditions and the equilibrium between solution and solid phase  
94 [19]. The Rhizon sampler was developed to collect pore-water from soils by vacuum  
95 filtration through a thin porous tube attached to a syringe, although it has not been used  
96 for sampling organic chemicals yet. Compared to centrifugation, the Rhizon sampler is  
97 a nondestructive *in situ* method which minimizes disturbance of the soil/sediment [20].  
98 However, it can only provide a resolution of 2 – 3 cm, and the small volumes collected  
99 (<10 mL) may not be sufficient to detect contaminants present at trace levels. A multi-  
100 section peeper was developed to measure the distribution of hydrophobic compounds  
101 in sediment porewater, with a resolution of 2 cm [21]. It relied on a laboratory-derived  
102 sampling rate (which varied between studies), to estimate the chemical concentrations.  
103 DGT (diffusive gradients in thin-films) is an *in situ* high-resolution dynamic technique,  
104 which accumulates labile contaminants in pore-waters and their resupply from the solid  
105 phase [22]. It provides time weighted average (TWA) concentrations, as a function of  
106 deployment time. DGT probes have been used to successfully measure the fluxes and  
107 the distribution of metals and nutrients in sediments at high spatial resolutions [23, 24].  
108 This is achieved by diffusion of contaminants followed by uptake into a binding gel,

109 which can then be sliced to small strips at fine scale (mm) and extracted. By combining  
110 the DGT technique with direct computer-imaging densitometry (CID) analysis or laser-  
111 ablation (LA) techniques, high resolution of 100s  $\mu\text{m}$  have been achieved and new  
112 knowledge of biogeochemical processes of inorganics have been obtained [25]. Until  
113 now, only 3 studies have used DGT to detect organic contaminants (pesticides, PFASs  
114 and antipsychotic drugs) in intact sediment cores [26-28]. However, DGT methods for  
115 a wide array of organic compounds – including antibiotics – have been developed and  
116 successfully applied in waters and soils [29]. Several studies have been carried out  
117 using DGT to measure antibiotics in river waters [30], sea waters [31] and soils [32].  
118 These studies on the application of DGT for *in situ* sampling in sediments and the  
119 established DGT methods for antibiotics have laid the foundation for this research.  
120 The aims of this study were therefore: i) to develop, test and validate the use of DGT  
121 probes for measuring antibiotics in sediments; ii) to perform laboratory studies on  
122 prepared sediments kept under controlled conditions; iii) to test the working conditions  
123 for DGT probes and to investigate the mobility of antibiotics spiked into sediment  
124 samples and iv) to use the DGT probe on intact lake sediment cores, to test its suitability  
125 for sampling the *in situ* distribution of antibiotics.

## 126 **2. Materials and Methods**

### 127 **2.1 Materials and Preparations**

128 Standard compounds sulfadiazine (SDZ), sulfamethazine (SMZ),  
129 sulfachloropyridazine (SCP), clindamycin (CLD, only for field work) and an internal  
130 standard sulfamethazine- $^{13}\text{C}_6$  (SMZ- $^{13}\text{C}_6$ ) were used for this work (obtained from

131 Sigma-Aldrich). Their selected physico-chemical properties are given in [SI Table S1](#).  
132 Stock solutions (1 mg mL<sup>-1</sup>) of all compounds were made and stored in pure methanol  
133 (MeOH). Acetonitrile (ACN) and MeOH of HPLC grade were purchased from Merck  
134 (Germany).  
135 The configuration of a DGT probe was first described elsewhere [22] and shown in [Fig.](#)  
136 [S1](#). The probes used here consisted of four layers: a polyethersulfone (PES) filter  
137 membrane on the bottom to prevent the binding gel from sticking to the backing plate,  
138 a 0.5 mm thick XAD18 binding gel, a 0.8 mm agarose diffusive gel, and a 0.14mm PES  
139 filter membrane on the top for protecting the gels. These four layers are held together  
140 by an acrylonitrile butadiene styrene (ABS) assembly with an exposure window of 1.8  
141 × 15 cm. The probes were purchased from DGT Research Ltd, UK. They were de-  
142 oxidized overnight, prior to deployment.

## 143 **2.2 Laboratory Experiments**

144 The purposes of the lab experiments were to test the performance (method sensitivity,  
145 deployment time, reliable spatial resolution) of the DGT probes for sampling antibiotics  
146 in sediment cores. To do this, layered sediments (both uncontaminated and spiked) were  
147 carefully prepared in a purpose-built tank, which allowed probes to be deployed *in situ*  
148 for known times, then removed, and pore-waters to be sampled with minimal  
149 disturbance of the sediments.

### 150 ***Experiment Tank Design***

151 The experiments were conducted in a glass tank designed for DGT probe deployment  
152 and pore-water sampling (see [Fig. 1\(a\)](#)). The tank (30 cm (length) × 20 cm (width) ×

153 30 cm (height)) had 9 pore-water sampling ports (4 mm diameter) on the front. Three  
154 upper ports were at -1.5, -5.5 and -9 cm below the sediment surface.

### 155 ***Sediment Preparation***

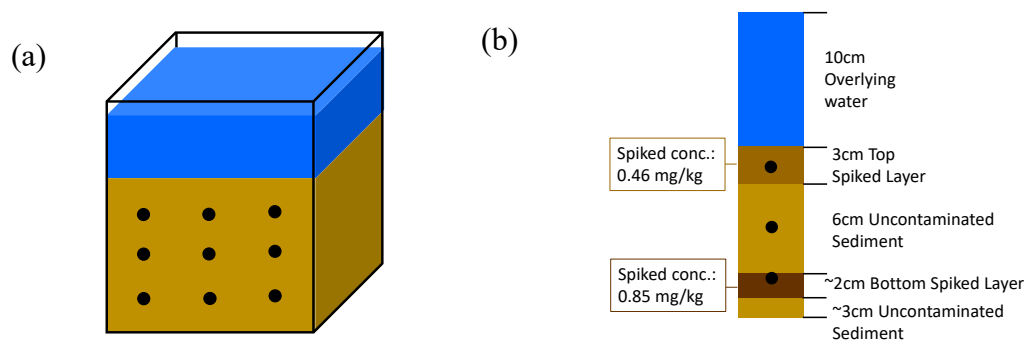
156 The relatively uncontaminated sediment was collected from the center of Lake Chaohu  
157 (Anhui, China) using a gravity corer (Ke Fan, China) [33]. Two different levels of  
158 spiking were performed to obtain sediments with 0.46 mg kg<sup>-1</sup> (low concentration  
159 spiking) and 0.85 mg kg<sup>-1</sup> (high concentration spiking) targeted antibiotics. Each spiked  
160 sediment was then well mixed in glass beakers. The high spiking concentrations were  
161 used in case the concentrations of target compounds were below detection limits, due  
162 to the fast adsorption of the freshly added compounds.

163 The sediments were transferred to fill the glass tank as shown in [Fig. 1\(b\)](#). The idea was  
164 to introduce contaminated and clean layers of sediment into pre-defined positions in the  
165 tank and then monitor any dispersal/change of the compound over time, using the  
166 probes. This would test the sensitivity and resolution of the probe, and demonstrate the  
167 applicability of the probe for detecting spatial and temporal changes of compounds *in*  
168 *situ*. Briefly, a layer of uncontaminated sediment was added to fill the bottom 3 cm of  
169 the tank; then a layer (~2 cm) of spiked sediment with high concentration was added;  
170 then another 6 cm of uncontaminated sediment was added; then an upper layer (3 cm)  
171 of spiked sediment with lower concentration was added. The top of the sediment was  
172 then smoothed flat and 6 L freshwater (no target antibiotics detected) was gently added  
173 into the tank, avoiding disturbance to the sediment. The depth of water overlying the  
174 sediment was ~10 cm. Prior to the experiments, the sediment-water system was then



175 left to settle for 2 days in a temperature-controlled room at 20.5 °C (the lowest  
176 temperature can be achieved in the warehouse) .

177 The tank was covered with black plastic bags to avoid possible photo-degradation of  
178 antibiotics and to minimize artificial stimulation of microbial activity.



179 **Fig. 1.** (a): Sketch of the experiment tank; (b): Distribution of the spiked and  
180 uncontaminated sediment introduced into the experimental tank. The black dots are  
181 ports for rhizon samplers

## 182 *DGT Probe Deployment and Pore-water Collection*

183 **Test 1:** Validation of sampling and analysis performance: A DGT probe (with XAD-18  
184 as binding layer resin, material particle size: 75 – 150  $\mu\text{m}$ ) was deployed in a well-  
185 stirred solution (0.01M NaCl, pH = 6.0) with known concentrations (given in Fig. 2 and  
186 Table S4) of three target compounds for 16 h. The binding gel was sliced into 5 mm  
187 strips and analysed following elution procedures described in later section. For the  
188 slicing procedures, the gels were cut from the exposure window with a razor blade and  
189 the diffusive gel and filter membrane were discarded. The binding gels were sliced to  
190 the desired resolutions using a micromanipulator guillotine [34].

191 **Test 2:** Investigation of spatial resolution of the technique: Three probes were carefully  
192 inserted vertically into the tank. After 24 h, they were retrieved and jet washed with  
193 MQ water to remove residual sediment. The binding gels were sliced at 10, 5 and 3 mm

194 intervals, respectively, to compare the spatial resolutions of the technique.

195 **Test 3:** Testing the response of the probes to deployment time: Three probes were  
196 inserted into the sediment simultaneously. The deployment was carried out as in Test 2.  
197 They were deployed for different time periods (12, 24 and 48 h). The binding gels of  
198 the three probes were cut into 5 mm intervals.

199 **Test 4:** Comparison of DGT and pore-water sampling: One probe was deployed in  
200 sediment tank for 24 hours. Overlying water and pore-water was also sampled (~3 ml  
201 per sample) during DGT deployment using 5 cm Rhizon samplers, which were inserted  
202 horizontally into the sampling ports at three levels (see [Fig. 1](#)).

203 **Test 5:** Exploring the behaviour of target compounds in sediments over fine scale:  
204 Probes were deployed for 24h at different time series (from Day 1 to Day 15) to  
205 visualize the change of distribution and labile concentration of target compounds.

### 206 **2.3 DGT Application in Intact Lake Sediment Cores**

207 One sediment core (diameter 8 cm, length 60 cm) was collected by gravity corer from  
208 the north west of Lake Chaohu, Anhui Province, China with PVC tubes in January 2020  
209 and stored in an incubator (15°C) before deployment of DGT probes. The sampling  
210 location was close to the entrance of a river which runs through Hefei city, in an area  
211 reported previously to be contaminated with various antibiotics, including 3 target  
212 compounds mentioned above [35]. After stabilizing the cores (by keeping them in a 15 °C  
213 water tank and covered with black plastic bags) in the laboratory for 4 days, DGT  
214 probes (XAD 18 – DGT for antibiotics measurement and Chelex-DGT for Fe/Mn redox  
215 condition measurement) were deployed for 48 h. The binding gels were sliced at 5 mm

216 resolutions.

## 217 **2.4 Sample Treatment and Chemical Analysis**

218 ***DGT samples:*** Each gel slice was eluted by 3 mL ACN with 30 min ultrasonic  
219 extraction. 0.5 mL of the extractant was sub-sampled and evaporated under a gentle  
220 stream of nitrogen to dryness, then reconstituted in a mixed solution of MeOH and MQ  
221 water (v/v = 1:9). 0.15 mL of overlying water/pore-water was diluted with MQ water  
222 and MeOH to the ratio MeOH: MQ = 1:9.

223 ***Lake sediment samples:*** The two sediment cores collected from the lake were then  
224 sliced at 1 cm intervals; each segment was transferred into a 50 mL centrifuge tube and  
225 centrifuged at 3000 rpm for 30 min. The pore-water obtained by centrifugation was  
226 then pre-concentrated by SPE cartridges (details in the [SI](#)). Then ~5 g of the sediment  
227 remaining after centrifugation was taken and mixed with 20 mL ACN. The mixture was  
228 shaken for 2 h, then centrifuged at 3000 rpm for 20 min. All the extractions and  
229 supernatants were evaporated to dryness under a gentle stream of nitrogen and  
230 reconstituted in 1 mL with MeOH:MQ = 1:9.

231 All solutions were filtered through 0.2  $\mu$ m PTFE syringe filters (Anpel, China) prior to  
232 the analysis. The samples were analyzed by UPLC-MS/MS (details in the [SI](#)).

## 233 **2.5 Calculations for DGT Measurements**

234 The principles of using DGT to sample analytes from soils and sediments has been  
235 described elsewhere [36]. Briefly, the analytes diffuse through the diffusion layer and  
236 are accumulated by the binding gel, leading to a linear concentration gradient in the  
237 diffusion layer. As analytes in the pore-water become progressively depleted, the

238 interfacial concentration (between the DGT probe exposure window and the sediment)  
239 of the analytes declines, which induces desorption from the solid phase. DGT therefore  
240 accumulates the labile analytes (the fraction that can easily dissociate and resupply) in  
241 the sediments and provides a time weighted average (TWA) concentration for the  
242 deployment period. Equation (1) was used to calculate the DGT measured  
243 concentration ( $C_{\text{DGT}}$ ,  $\mu\text{g L}^{-1}$ ) at the interface of the sediment and the DGT probe:

$$244 \quad C_{\text{DGT}} = \frac{M \Delta g}{DA t} \quad (1)$$

245 where,  $M$  is the mass of analyte accumulated on the binding gel (ng),  $t$  is the deployment  
246 time (s),  $D$  represents the diffusion coefficient of the analyte in the diffusion layer ( $\text{cm}^2$   
247  $\text{s}^{-1}$ ),  $A$  is the sampling area ( $\text{cm}^2$ , depends on the desired resolution),  $\Delta g$  is the thickness  
248 of the diffusion layer (cm). The  $D$  values for SDZ, SCP and SMZ at  $15^\circ\text{C}$  (average  
249 temperature during the experiment) are  $3.73 \times 10^{-6}$ ,  $3.32 \times 10^{-6}$ , and  $3.54 \times 10^{-6} \text{ cm}^2 \text{ s}^{-1}$   
250 respectively, according to the previous research [30].

251 The average flux of analytes ( $F$ ,  $\text{ng cm}^{-2}\text{s}^{-1}$ ) from the sediment to the DGT probe during  
252 the deployment time ( $t$ ) and defined sampling area ( $A$ ), can be obtained using the  
253 Equation (2).

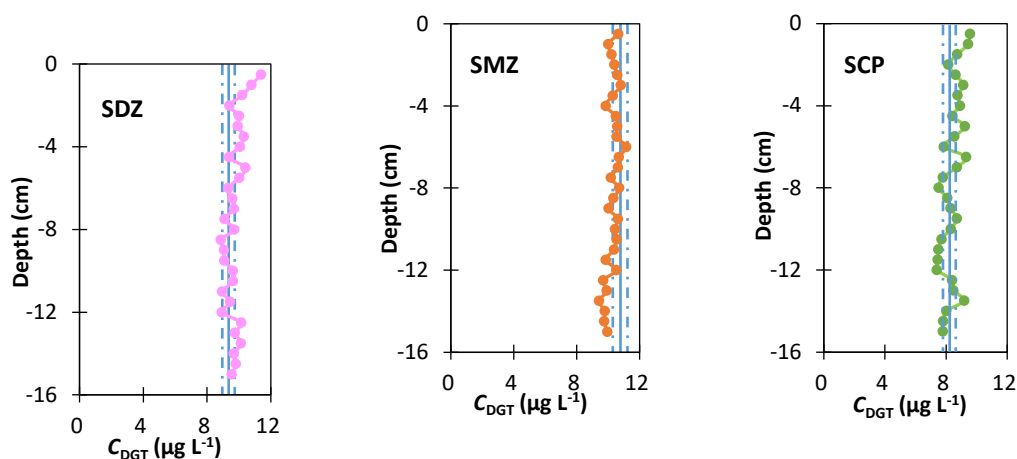
$$254 \quad F = \frac{M}{A t} \quad (2)$$

### 255 **3. Results and Discussions**

#### 256 **3.1 Validation of DGT Probe Performance**

257 The quality (accuracy and precision) of the measurements using DGT probes could be  
258 influenced by errors introduced from gel cutting, elution procedures and instrumental  
259 analysis. Previous studies have shown the homogeneity of the agarose binding gel

260 incorporated with WAX resin at 3 cm intervals [27], while the measurement of trace  
261 metals using DGT probes with a Chelex-100 binding gel had a precision of <10% at a  
262 resolution of 1.25 mm [22]. The relative standard deviation of the measurement of three  
263 target compounds obtained by the DGT probe with XAD-18 binding gel were all <8%  
264 (Fig. 2), and the concentrations derived from DGT were 94 – 102% of the  
265 concentrations in the spiked solution. These results confirmed that accurate and precise  
266 measurements of antibiotics can be made at high spatial resolution of 5mm using DGT  
267 probes.



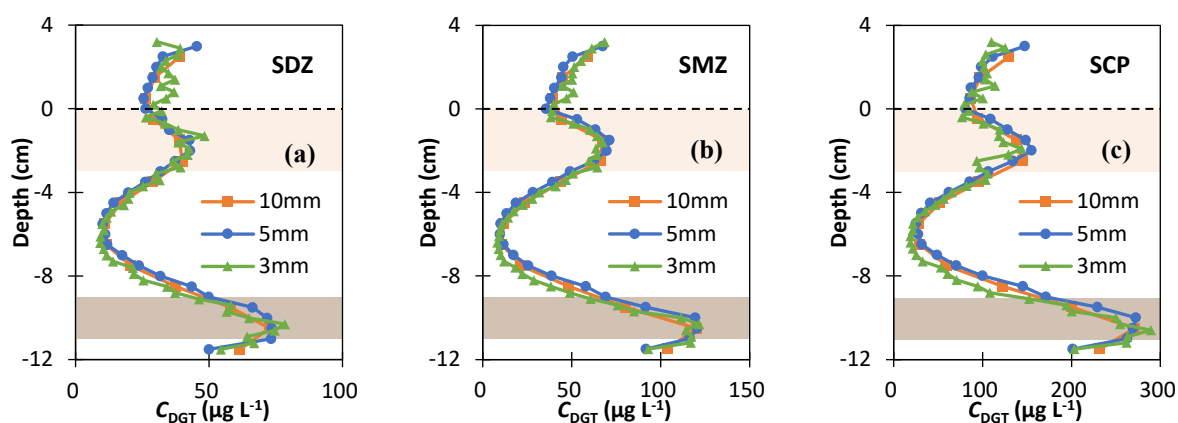
268 **Fig. 2.** Concentration profiles of three target antibiotics measured using a DGT probe  
269 in a well-stirred solution (deployment time = 16h). Blue solid lines are concentrations  
270 of the solution with standard deviations (dotted lines).

## 271 3.2 Performance of DGT Probes in Sediments – Laboratory Testing

### 272 3.2.1 Different spatial resolutions

273 Observing fine scale heterogeneity of contaminants in sediments relies on achieving  
274 appropriate spatial resolution and reliable analysis [37]. As shown in Fig. 3, the 3  
275 vertical concentration profiles obtained with 10, 5 and 3 mm resolutions gave good  
276 agreement for the overall concentrations and distributions of each target compound.

277 This suggests i) the sediment tank test system had been prepared with consistent layered  
278 structure across the locations where the probes were deployed; ii) DGT probes have the  
279 ability to detect accurately the peaks and troughs of antibiotic concentrations at high  
280 spatial resolution in the sediment profiles and iii) resolution of 3 mm can be achieved  
281 to give more details of the concentration profiles.



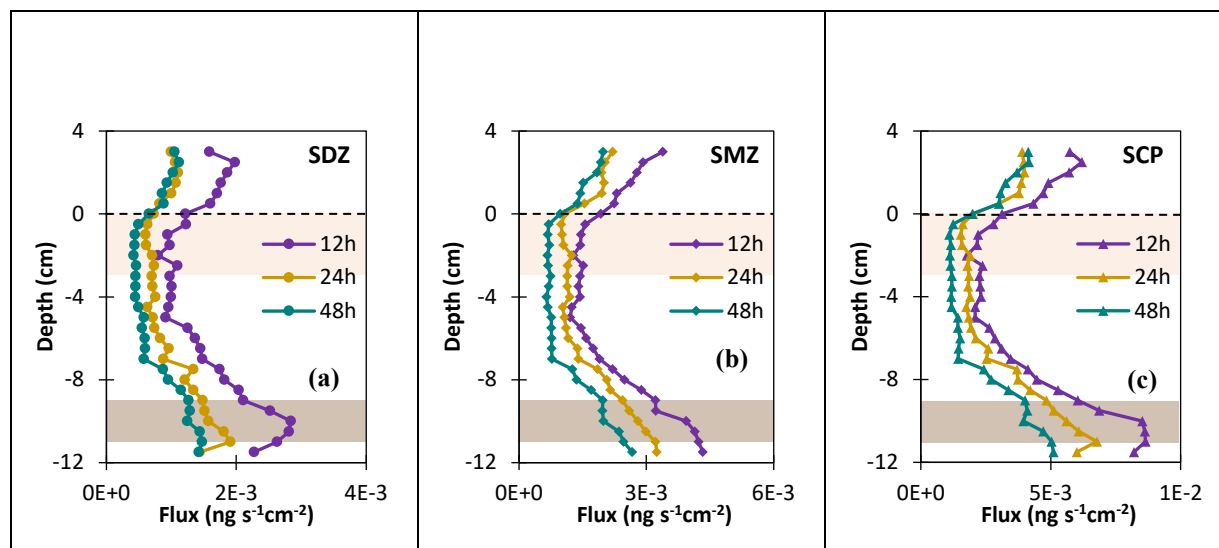
282 **Fig. 3.** DGT profiles for 3 target antibiotics at different resolutions (10, 5 and 3 mm).  
283 The shaded boxes in the background are the initial spiked layers.

### 284 3.2.2 Rates of release from the sediment to the DGT probes

285 DGT samples the labile portion of compound – that in the solution phase, other forms  
286 that can pass through the diffusive layer and a desorbed fraction re-supplied from the  
287 solid phase when the solution phase is depleted by uptake. Test 2 was designed to  
288 investigate the resupply of the antibiotics from the solid phase to DGT probes over time,  
289 by comparing the fluxes after 12, 24 and 48 hours of deployment in the tank.

290 This test was conducted at 15 days after the spiking, the peaks of the upper  
291 contaminated layers (as shown in Fig. 3) disappeared. As presented in Fig. 4, in the  
292 lower concentration areas (~ from sediment surface to -8 cm), the decrease of fluxes  
293 from 24 h to 48 h was larger than that from 12 h to 24 h, indicating a gradual reduction

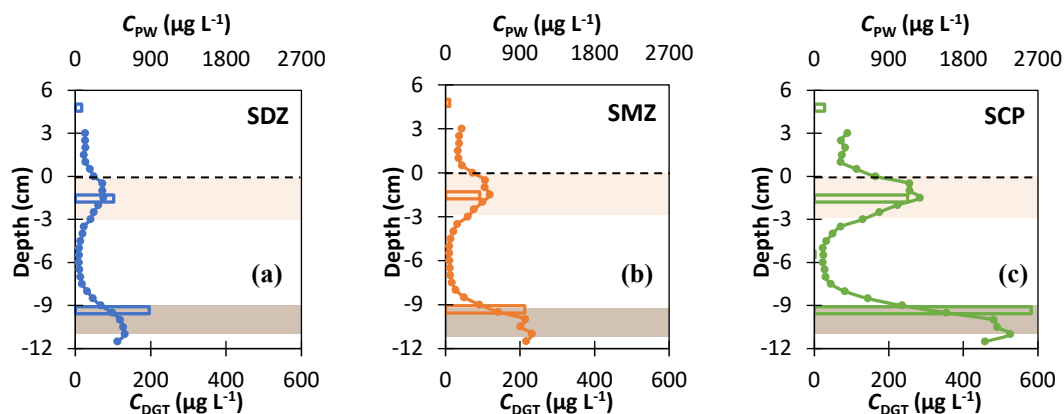
294 of the resupply of the analytes. In the higher concentration areas (deeper spiked layer  
295 from -8 to -11 cm and the overlying water), the opposite situation occurred, showing a  
296 fast reduction of re-supply in the beginning, then the rate of re-supply slowed down.



297 **Fig. 4.** DGT flux profiles for 3 target antibiotics deployed for different lengths of time  
298 (12, 24 and 48 h). The shaded boxes in the background are the original spiked layers.

### 299 3.2.3 Comparison of DGT and Rhizon Pore-water Measurements

300 A DGT probe was deployed in parallel with the *in situ* pore-water sampling of the  
301 Rhizon samplers. The pore-water sampling could not be undertaken with the same  
302 spatial resolution as the DGT sampling (the sampling ports for pore-water were at -1.5,  
303 -6 and -9.5 cm, overlying water was sampled at 5 cm), but the profiles of these two  
304 approaches were similar. The highest pore-water concentrations occurred at -9.5 cm,  
305 the mid-values at -1.5 cm, both in/close to the spiked sediment layers, while the lowest  
306 Rhizon sampler concentrations were detected at -5.5 cm in uncontaminated sediment  
307 layer (Fig. 5). The concentrations measured at -5.5 cm were low (see Fig. 5), but the  
308 detailed data of concentrations measured with the Rhizon sampler at different depths  
309 are presented in the Table S5.



310 **Fig. 5.** Concentration-depth profiles of the 3 target antibiotics collected by DGT probe  
 311 (solid dots) and Rhizon samplers (bar). The shaded boxes in the background are the  
 312 original spiked layers.

313 The DGT probe was able to provide the vertical distribution of target analytes in  
 314 sediments with a much higher resolution. There was good correspondence between the  
 315 profiles using the two methods in this laboratory-controlled system relied on pre-  
 316 existing knowledge of the peak locations. In real environments, where the distribution  
 317 of the analytes is not known in advance, if the spatial resolution of the measurement is  
 318 not high enough, such peaks may be missed.

319 Concentrations obtained by Rhizon pore-water sampling ( $C_{PW}$ ) were 3 – 9 times larger  
 320 than those of labile antibiotics obtained by DGT sampling ( $C_{DGT}$ ). This difference has  
 321 been obtained and explained previously in other studies for other analytes [38, 39].

322 DGT measures the analytes through an accumulating process, the analytes adjacent to  
 323 the DGT sampler surface are depleted by DGT accumulation. The re-supply of analytes  
 324 from that adsorbed on the solid phase is required to sustain the flux from pore-water to  
 325 the DGT probe. This may be kinetically limited in some cases. DGT only measures  
 326 labile species that can be desorbed from the sediment and diffused through the



327 sampler's diffusive gel layer. Some compounds may be complexed and too large to  
328 diffuse through the pores of the gel layer. Therefore, it is reasonable to have smaller  
329 labile concentrations measured by DGT compared to the total dissolved concentrations  
330 by pore-water sampling.

331 In the overlying water (above 1 cm), the labile concentrations of 3 antibiotics measured  
332 by DGT were – on average – 26 (SDZ), 37 (SMZ) and 78 (SCP)  $\mu\text{g L}^{-1}$ , again much  
333 less than the 78, 51 and 124  $\mu\text{g L}^{-1}$  measured by Rhizon sampling of the water column.

334 Except for the non-labile fraction as explained above, this can mainly be ascribed to the  
335 large DBL (diffusive boundary layer) which affects DGT water sampling in the static  
336 water-sediment tank [40]. The DBL thickness was measured around 0.44 – 0.45 mm  
337 for metals [40] and 0.75 mm on average for some organic contaminants [41] in static  
338 solutions. If we take 0.5 mm as the DBL thickness for the target compounds,  $C_{\text{DGT}}$   
339 values for them were 40 (SDZ), 57 (SMZ) and 119 (SCP)  $\mu\text{g L}^{-1}$ , much closer to the  
340 measured  $C_{\text{PW}}$  values.

341 The  $R$  value ( $C_{\text{DGT}}/C_{\text{PW}}$ ) has often been used as an indicator of the extent of the  
342 depletion of pore-water concentrations at the DGT interface, to reflect the resupply  
343 efficiency from the solid phase to pore-water, in response to the depletion by DGT [36].

344 Higher  $R$  values indicate a greater tendency to resupply. If  $R \geq 0.90$  (indicating that the  
345 analytes can be re-supplied rapidly from the solid phase to porewater and then to the  
346 DGT probe), this means there is weak binding of the compound to the sediment. If  $R <$   
347 0.1, it normally indicates there is no resupply from the solid phase, supply of the  
348 analytes solely depends on the diffusion from the pore-water. If the  $R$  value is between

349 0.1 and 0.9, the analytes in the pore-water are partially sustained, indicating the  
350 resupply from the solid phase is slower than the depletion of DGT uptake.

351 Typical  $R$  values of the 3 antibiotics in this study at the two different depths are shown  
352 in [Table 1](#). The values were different between compounds and varied with sampling  
353 depth, being higher in the more reactive upper layer. This is the first time that DGT was  
354 used to investigate antibiotics in sediments, so these  $R$  values are the first set for  
355 antibiotics in sediments. Chen et al. [42] measured  $R$  values of  $\sim 0.13$  for SMZ in  
356 controlled soil experiments, a value close to that in the deeper layer in this study. Here  
357 the compounds were freshly added, while in the real environment  $R$  values might be  
358 even lower, as compounds age and become more associated with sediment particles  
359 over time [43].

360 **Table 1.**  $R^a$  values derived for the 3 antibiotics in the controlled sediment study.

Depth (cm)	SDZ	SMZ	SCP
-1.5	0.16	0.29	0.25
-9.5	0.11	0.15	0.14

361 <sup>a</sup>:  $R = C_{\text{DGT}} / C_{\text{PW}}$

### 362 *Time Series Measurements*

363 The concentration profiles of the 3 antibiotics in [Fig. 6](#) and [Fig. S2](#) demonstrate the  
364 diffusional transport of these compounds from spiked sediment to the adjacent  
365 uncontaminated sediment layers and overlying waters over time. Comparing to [Fig.](#)  
366 [1\(b\)](#), the target compounds spread up and down from the 2 discrete layers of  
367 contaminated sediments that had been introduced into the tank initially. During the  
368 experiment, DGT measured declines in the maximum peak ( $\sim -11$  cm) and secondary

369 peak ( $\sim -1.5$  cm), with more rapid decreases from the secondary peak. This may be  
370 because compounds in the upper spiked layer dispersed both to the adjacent unspiked  
371 layers and overlying waters, while diffusion from the deeper contaminated layer was  
372 only through pore-water. According to the Einstein relationship (Eq. 3) [44], the  
373 diffusive distance can be roughly estimated.

$$374 \quad L^2 = 2D_s t_e \quad (3)$$

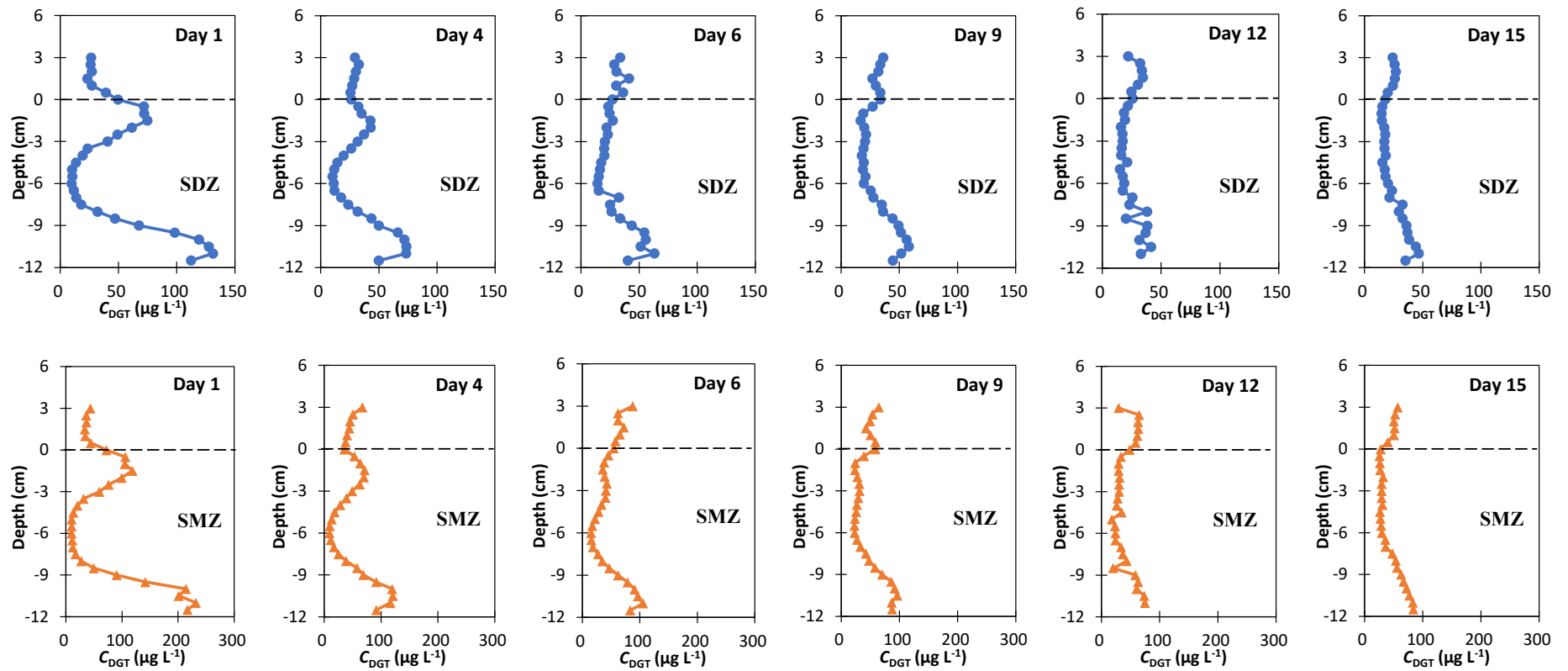
375 Where  $L$  is the diffusive distance (1D),  $D_s$  represents the diffusion coefficient in  
376 sediments, and  $t_e$  is time. The diffusive distances of the 3 target compounds were 0.76  
377 – 0.80 cm after 24 h, while after 15 days, the diffusive distances were 2.94 – 3.11 cm.  
378 However, the antibiotics could be detected at greater depths after 24 hours,  
379 demonstrating that some transportation of compounds in the sediments had occurred by  
380 processes in addition to diffusion. This is evidence that there was some physical  
381 disturbance/mixing of some of the original contaminated sediment (probably the finest  
382 fraction/colloids) when the system was set up and the tank was settling down. After that  
383 (after 24 h) transport is probably mainly controlled by diffusion, with no additional  
384 physical mixing/disturbance.

385 The changes of distribution and concentration of the 3 target compounds in pore-waters  
386 showed a similar pattern (see in [Fig. S3](#)). In addition to diffusion to and adsorption by  
387 adjacent sediments, degradation could also make a little contribution to the changes in  
388 concentration. Biodegradation of freshly added SMX has been observed over 24 days  
389 after spiking in a non-sterile sediment [45]. Because the tank was covered with black  
390 plastic, any photo-degradation should have been minimal.

391 In summary, the changes in the concentrations/profiles of target antibiotics after 24  
392 hours of the spiking are believed to be mainly due to adsorption to the sediment particles,  
393 since the concentration decrease in the spiked layer was much larger than the increases  
394 in unspiked layers during the experiment. Diffusive transfers from the original  
395 contaminated layers also contributed to the reduction of compounds from the spiked  
396 layers, likely together with a small degree of biodegradation.

### 397 *Comparison of Chemical Behaviours*

398 The 3 antibiotics all have high aqueous solubility ( $77 - 7000 \text{ mgL}^{-1}$ ) and low/moderate  
399 partition coefficients to sediments ( $\log K_{ow}$  from -0.09 to 0.31) [30]. They can therefore  
400 readily enter the sediment pore-water and overlying water column. The concentration  
401 profiles of these 3 compounds obtained by DGT probes had the same trend as shown in  
402 the figures above. Although the spiked concentrations of antibiotics were the same, the  
403 labile concentrations measured by DGT probes were in the order of  $\text{SDZ} < \text{SMZ} < \text{SCP}$   
404 at all depths (see Fig.5). This order is the opposite to what has been observed in acid  
405 soils (pH ranging from 4.5 to 5.1) [46], while the affinity of SCP was less than SMZ in  
406 neutral soils [47]. The sediment used in this work was basic with pH of 8.5; this could  
407 be the reason for the mobility order of these three antibiotics being opposite to that in  
408 acid soils.



409 **Fig. 6.** DGT profiles of SDZ (blue dots) and SMZ (orange triangle) measured at different times  
 410

411 ***Comments on detection limits and sampling times***

412 Compound-specific detection limits can be derived using eq. 1. In this study, the  $D$   
413 values of the target antibiotics were between  $3.32 \times 10^{-6}$  and  $3.73 \times 10^{-6}$  (at  $20.5^\circ\text{C}$ ), the  
414 thickness of the diffusion layer was 0.094 cm, the exposure area ( $A$ ) was  $0.9 \text{ cm}^2$  (for  
415 the 5 mm intervals). The IDLs of the target compounds were  $\sim 50 \text{ ng L}^{-1}$  and final sample  
416 volumes ( $V_0$ ) was typically 1 mL. Therefore, the MQLs (method quantification limit)  
417 of DGT sampling were about  $16 - 18 \text{ ng L}^{-1}$  for deployment of 24 h as in this study.

418 The minimum sampling time for field deployment means the time needed for the  
419 amount of target compounds adsorbed by the binding gel to exceed the IDLs. In the  
420 present study, taking SDZ as an example, the minimum sampling time ( $t_{\min}$ ) can be  
421 calculated by:

$$422 \quad t_{\min} = \frac{\Delta g \times M_{IDL}}{D \times A \times C_{DGT}} \quad (3)$$

423  $C_{DGT}$  can be estimated by  $C_{PW}$  and an empirical value of  $R$  (measured in the range  
424 between 0.1 – 0.3 here).  $M_{IDL}$  is the mass accumulated on the binding gel (ng),  $M_{IDL} =$   
425  $IDL \times 1000 / V_0$ . If the IDL was  $50 \text{ ng L}^{-1}$  (see above), and  $C_{PW}$  values in a sediment  
426 core are typically  $10 - 100 \text{ ng L}^{-1}$  [48, 49] then the  $t_{\min}$  needed will range from 5 hours  
427 to 2 days for resolution of 5mm.

428 DGT method sensitivities can be improved (e.g. by a factor of 10), by simply reducing  
429 the final sample volumes, increasing the deployment times, reducing the diffusion layer  
430 thickness and increasing the exposure areas. The MQLs using the Rhizon sampler were  
431  $\sim 16 \text{ ng L}^{-1}$  too (assuming the pre-concentration factor = 3). In the real environment, if  
432 the concentrations of the target compounds are lower, more pore-water is required to

433 obtain a higher pre-concentration factor for the Rhizon samplers. Another widely used  
434 method to obtain pore-waters samples is slicing the sediment core layers followed by  
435 centrifugation. However, this *ex situ* approach would certainly disturb the sediment  
436 structures and change the redox conditions and pH, which control chemical speciation  
437 and availability. The advantages of the DGT technique are directly sampling *in situ* in  
438 field conditions, purer matrix and greater analytical sensitivity.

### 439 **DGT application in intact lake sediment cores**

440 To validate the application of DGT probes for antibiotics in the real environment,  
441 probes were deployed in the top 13 cm of the intact sediment cores collected from the  
442 field. In the sediment core collected in Chaohu Lake, only SMZ of the 3 target  
443 antibiotics was detectable, concentrations of the other two target compounds were  
444 <MDL. The SMZ depth profiles of DGT measured concentrations ( $C_{DGT}$ ) and pore-  
445 water concentrations obtained by slicing and centrifugation ( $C_{PW}$ ), are shown in [Fig. 7](#)  
446 [\(a and b\)](#). Another commonly used antibiotic, clindamycin (CLD) was also detected in  
447 the sediment core. The results of concentration profiles of CLD are presented in [Fig. 7](#)  
448 [\(c and d\)](#). To assess the redox condition of the sediments, additional information on the  
449 DGT measured Fe and Mn profiles is presented in [Fig. S4](#).

450 The baseline DGT (labile) concentrations of SMZ were quite constant ( $\sim 50 \text{ ng L}^{-1}$ )  
451 with depth. Two peaks were present at different depths. A slight increase in  $C_{DGT}$  was  
452 observed around -5.5 cm, while a maximum concentration of  $515 \text{ ng L}^{-1}$  was observed  
453 at -10.5 cm, followed by subsequent decrease below that depth. The  $C_{PW}$  profile showed  
454 a similar trend to  $C_{DGT}$  with two peaks, and the concentration gradually increased to

455 maximum value of 32.5 ng L<sup>-1</sup> at -11 cm.

456 It is an unusual phenomenon that  $C_{DGT}$  values at all depths were much higher than  $C_{PW}$ .

457 Fe and Mn are oxygen sensitive elements and their concentration profiles can be used

458 to assess the redox condition of the sediment. The concentration profile of labile Fe (II)

459 and Mn (II) measured in the same sediment core is shown in SI Fig. S4, a DGT probe

460 with Chelex-100 in binding layer for trace metals was deployed *in situ* and measured at

461 5 mm resolution. The results demonstrated that the reduction of Fe and Mn oxides/

462 hydroxides to labile Fe (II) and Mn (II) occurred in the upper layer of the sediment,

463 intensified at -4.0 cm, below which the sediment was in an anoxic condition. Slicing

464 and centrifugation of the sediment were both conducted in the open air, which destroyed

465 the anoxic condition of the sediment, SMZ may be rapidly adsorbed to the precipitated

466 iron oxides, the degradation of the target compounds might also be accelerated [50].

467 Therefore, these conventional *ex situ* extraction approaches may seriously

468 underestimate the compound levels detected.

469 A similar situation was observed for CLD as shown in Fig. 7 (c and d). The profiles of

470 concentrations measured by two methods have comparable shape, with a gradual

471 upward trend above -9.0 cm, then increased sharply at -9 to -10 cm. The  $C_{DGT}$  profile

472 has more variations as it was measured at higher resolution. The change of the redox

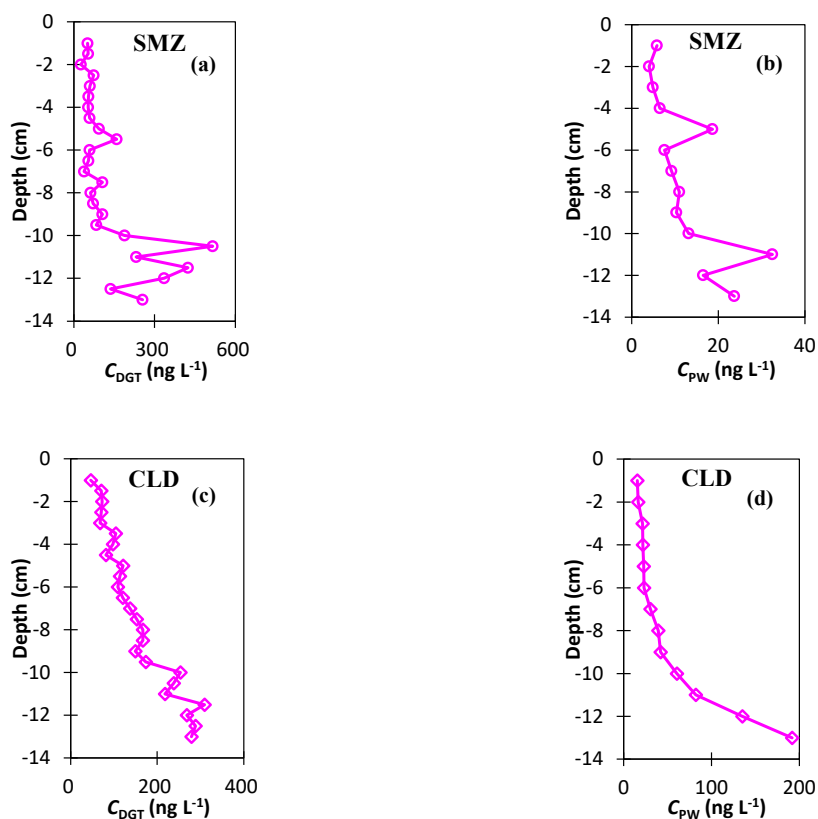
473 condition also influenced the measurement of CLD concentrations in pore-water,

474 proving that an *in situ* sampling approach, such as DGT, is more suitable for the

475 measurement of compounds in sediments.

476





477 **Fig. 7.** SMZ and CLD concentration distribution profiles in Lake Chaohu sediments by  
 478 DGT and pore-water measurements (a: SMZ measured by DGT; b: SMZ measured by  
 479 pore-water extraction; c: CLD measured by DGT; d: CLD measured by pore-water  
 480 extraction.

### 481 **Environmental Implication and Future Research Directions**

482 This is the first systematic study to validate and demonstrate the use of the DGT probe  
 483 for high spatial resolution (sub-mm/mm scale) sampling of antibiotics in sediments.  
 484 Deploying probes over 12 – 48 hours, detection limits of 10s ng L<sup>-1</sup> were readily  
 485 achieved by this approach and it can be improved by optimizing the deployment  
 486 conditions. Finer spatial scale measurement can be achieved with suitable detection  
 487 systems to investigate compound distribution *in situ* at the mm scale. This makes it  
 488 possible to conduct studies on the influence of redox changes and processes in the  
 489 surface microlayer of the sediment-water interface, where the exchange, binding and

490 breakdown of organic compounds can occur. Monitoring *in situ* rates of dissipation,  
491 transformation and exchange is now possible.

492 As a dynamic sampling technique, DGT is able to provide kinetic information in  
493 sediments including fluxes, labile pool size and kinetic resupply. In a previous study,  
494 kinetic information was determined for pesticides in sediments.[32] In the future, DGT  
495 probes can be used together with the DIFS (DGT-induced fluxes in soils and sediments)  
496 model, to obtain better knowledge of such processes for antibiotics.

497 The microcosm study showed that the 3 test antibiotics can rapidly re-mobilise and  
498 migrate from sub-surface contaminated layers into clean sediments and overlying water  
499 bodies by diffusion transport, adsorption processes and possibly certain degree of  
500 biodegradation. In fact, DGT has been developed for several other groups of organic  
501 compounds (including personal care products, endocrine disrupting chemicals,  
502 pesticides, psychiatric pharmaceuticals and polyfluoroalkyl substances),[51-56] so its  
503 application to investigate fine-scale biogeochemical processes in sediments and the  
504 sediment-water interface can also be extended to other compound classes. Investigation  
505 on the effect of bioperturbation and dynamic transport may also possible using DGT  
506 technique in different conditions of the sediments.

507 **CRedit authorship contribution statement**

508 Yanying Li: Investigation, Data analysis, Writing – original draft. Qiuyu Rong:  
509 Investigation, Validation. Chao Han: Methodology, Sample collection. Hanbing Li:  
510 Validation, Methodology. Jun Luo: Supervision, Funding acquisition. Liying Yan:  
511 Investigation, Data analysis. Degao Wang: Supervision. Kevin C. Jones:

512 Conceptualization, Supervision, Writing – review & editing, Funding acquisition. Hao  
513 Zhang: Conceptualization, Supervision, Writing – review & editing, Funding  
514 acquisition.

#### 515 **Declaration of Competing Interest**

516 The authors declare that they have no known competing financial interests or personal  
517 relationships that could have appeared to influence the work reported in this paper.

#### 518 **Data availability**

519 The authors do not have permission to share data.

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#### 524 **Supporting Information**

525 Supplementary data associated with this article can be found in the online version at  
526 Additional information as noted in the text including analytical methods, five tables and  
527 four figures.

528

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