

Supporting information for:
**Novel Probe for In Situ Measurement of Freely
Dissolved Aqueous Concentration Profiles of
HOCs at the Sediment-Water Interface**

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1 Pore water probe mooring schematic

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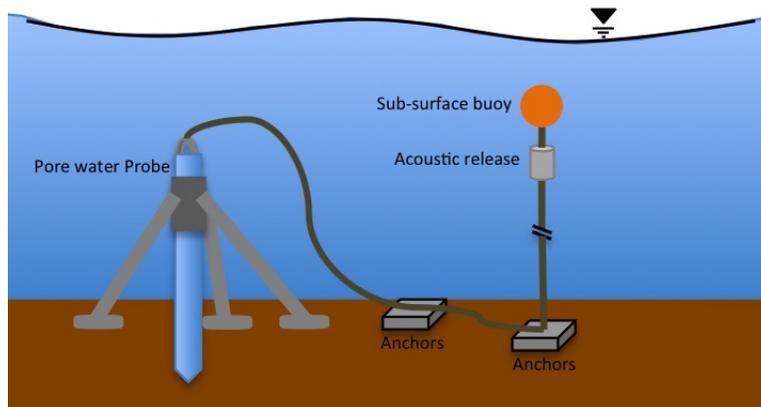


Figure S1: The pore water probe frame is attached to two anchors in series, and the end of the rope is attached to an acoustic release with additional 60 m of rope inside the canister, and finally to a buoy that is held below the water surface to avoid entanglement with boat traffic. When the probe is retrieved, an acoustic deck on board is used to release the acoustic transponder and the additional rope is released that allows the sub-surface buoy to come up to the water surface, and the whole mooring is carefully pulled onboard the boat.

2 Passive samplers

3 PE was obtained from Brentwood Plastics (St. Louis, MO, USA). PE strips were cleaned by
4 solvent washing sequentially overnight with pesticide-grade hexane, acetone, and deionized
5 (DI) water.^{S1} Samplers used in the field were impregnated with PCB155 in two batches of 1-L
6 80:20 methanol: DI water solution spiked with 100 μ L of 100 ppm PCB155 standard solution
7 in hexane (Ultra Scientific, N. Kingstown, RI, USA).^{S2,S3} PRC batch solutions were placed
8 on a shaker for two weeks for equilibrium to be reached. Afterwards, samplers were rinsed
9 with DI water, gently wiped dry with Kimwipe, and stored in a clean 1-L amber jar at 4°C
10 prior to deployment. Lab blanks were extracted to check for consistent PRC concentrations
11 in passive samplers. Standard deviations in PRC concentration in the lab blanks were < 5%
12 of total PRC concentration.

13 PE-water partition values for DDT and its metabolites were measured in the laboratory.
14 PE strips were first impregnated with DDT its metabolites (4,4'-DDT, 2,4'-DDT, 4,4'-
15 DDD, 2,4'-DDD, 4,4'-DDE, 4,4'-DDE, DDMU) in a methanol-water solution spiked with
16 a standard cocktail with the seven DDT analytes in hexane (Ultra Scientific). Specifically,
17 a 200 mL solution of 80:20 methanol:DI water was spiked with 25 μ L of DDT metabolite
18 standard in hexane, and six strips of PE (2.5 cm by 2.5 cm, 0.03 g each) were added.^{S2} This
19 solution was allowed to equilibrate on a shaker for 1 month, and strips were wiped dry with
20 Kimwipe. In triplicate, one DDT-impregnated PE strip was added to a 1-L amber bottle
21 filled with DI water and sodium azide (1 g/L) to avoid bacteria growth. This was allowed to
22 equilibrate for 3-months on a shaker at 150 rpm. PE and water were extracted separately
23 and analyzed to measure the DDT metabolite concentrations in each phase.^{S4} Partition
24 values were calculated by dividing PE concentrations by water concentrations. Measured
25 partition values are presented in Table S1.

26 PE-water partition values (K_{PE}) values measured from other studies are also included
27 Table S1 for comparison. All the DDT metabolite K_{PE} values are statistically the same as
28 the other studies^{S5,S6} (t-test $p > 0.05$) except for 2,4'-DDE. Therefore, we have a high degree

29 of confidence in the K_{PE} values for the DDT metabolites, except for 2,4'-DDE, which may
 30 be inflated based on comparison with the other two studies. However, since the measured
 31 2,4'-DDE in the overlying water and porewater were either below detection limits or less
 32 than 1% of total DDT concentrations, and potential error in the K_{PE} for 2,4'-DDE did not
 33 affect results.

34 Since the temperature dependency of the partition value is mostly determined by the
 35 decrease in HOC solubility in water as the temperature decreases, the temperature rela-
 36 tionship determined by Jonker et. al.^{S7} for silicone-rubber samplers should be consistent
 37 for PE, and the adjustment to the laboratory-measured $\log K_{PE,i}$ would change by $\log 0.04$,
 38 which is well within the error expected.^{S8} Even more conservative estimates, suggest the
 39 temperature dependence is expected to be between 0.1 and 0.2 log units.^{S8,S9} The effects of
 40 temperature on partition values should not affect the shape of the concentration gradients
 41 and the calculated fluxes.

42 All PE samples were analyzed for DDT and PCB analytes using previously described
 43 cleanup and analytical methods.^{S10}

Table S1: PE Partition Coefficients, Average $\log K_{PE}$ (L water/kg PE) measured at 20°C, standard deviation in parenthesis.

Chemical	K_{PE} (this study)	K_{PE} ^{S5}	K_{PE} ^{S6}
DDMU	5.0 (0.2)	5.4 (0.3)	5.5 (0.1)
2,4'-DDE	6.6 (0.5)	5.7 (0.4)	5.8 (0.1)
4,4'-DDE	5.9 (0.2)	5.8 (0.3)	5.9 (0.1)
2,4'-DDD	5.1 (0.3)	5.0 (0.2)	4.8 (0.03)
4,4'-DDD	5.0 (0.4)	4.9 (0.2)	4.8 (0.04)
2,4'-DDT	5.6 (0.4)	5.8 (0.5)	5.9 (0.05)
4,4'-DDT	5.2 (0.3)	5.6 (0.3)	5.6 (0.05)
PCB69	5.35 ^{S3}		
PCB103	5.78 ^{S3}		
PCB155	6.1 ^{S2}		

44 After retrieval, passive samplers were individually rinsed with DI water and wiped dry
 45 with Kimwipe to remove any sediment particles, biofilm, or organisms. Portions of passive
 46 samplers that were covered by the attachment plate (Fig. 2) were cut off. Samplers were

47 individually placed in 40 mL amber pre-cleaned glass vials that were pre-labeled, and stored
48 at 4°C prior to and during shipment.

49 Determination of Aqueous Concentration

50 In general, a simplified first order equation is used to model passive sampler uptake rate
 51 (equation 1).^{S1,S2}

$$C_{PE,i}(t) = K_{PE,i} \cdot C_{w,i} \cdot (1 - e^{-k_{e,i}t}) \quad (1)$$

52 where $C_{PE,i}$ is the target chemical concentration in PE after deployment period t (ng/g);
 53 $K_{PEW,i}$ is the target chemical PE-water partition coefficient (L/kg); $C_{w,i}$ is the target chem-
 54 ical concentration in water (ng/L); $k_{e,i}$ is the target chemical exchange rate coefficient (1/d);
 55 t is the deployment period (days). Equation 2 results by rearranging eq. 1, where the
 56 adjustment factor, $f_{adj} = \frac{1}{1 - e^{-k_{e,i}t}}$. The exchange rate coefficient $k_{e,i}$ encompasses relevant
 57 environmental factors, such as hydrodynamics, temperature, and biofouling that control
 58 the uptake of HOCs through the aqueous boundary layer. Use of performance reference
 59 compounds utilizes the assumption that the uptake rate of chemicals is equivalent to the
 60 elimination rate. Thus, the elimination rate of the PRC is used to calculate the uptake rate
 61 of chemicals measured. If the PRC is not the isotopically labeled form of the compound of
 62 interest, then the exchange rate of the compound is related to the exchange rate of the PRC
 63 based on the empirical relation in equation 4 .

$$C_{w,i} = \frac{C_{PE,i}(t)}{K_{PE}} \cdot f_{adj} \quad (2)$$

$$C_{PE,PRC}(t) = C_{PE,PRC}(t = 0) \cdot e^{-k_{e,PRC}t} \quad (3)$$

$$\frac{K_{PE,i} \cdot k_{e,COC}}{K_{PE,PRC} \cdot k_{e,PRC}} \propto \left(\frac{D_{w,i}}{D_{w,PRC}} \right)^{2/3} \propto \left(\frac{M_{PRC}}{M_i} \right)^{1/3} \quad (4)$$

64 where $D_{w,i}$ and $D_{w,PRC}$ are the aqueous diffusivity of the chemical and PRC, respectively,
 65 and M is the molar volume.

66 Assessment of equilibrium in PE

67 We estimated that 90 days should be sufficient for near-equilibrium in deployed PE based
68 on previous measurements with an earlier prototype of the pore water probe which involved
69 attaching PE to stainless steel plates^{S3} attached to an anchor and deployed for 2 months at
70 the same location in Pallanza Bay (unpublished study). These PE samplers were spiked with
71 2, 3', 4, 5-tetrachlorobiphenyl (PCB 69) and 2, 2', 4, 5', 6-pentachlorobiphenyl (PCB 103). The
72 calculated average (\pm st. dev.) exchange rate coefficient (k_e) for PE just above the sediment
73 surface was approximately $0.015 \pm 0.004 \text{ day}^{-1}$ for the two PRCs (n=4). This k_e is a time-
74 averaged uptake rate during the sampling period, and was calculated by solving for k_e in eq.
75 3 after a deployment period of $t = 60$ days, where $C_{PE,PRC}(t)/C_{PE,PRC}(0)$ is the fraction
76 of PRC concentration remaining in the PE after the deployment period.^{S1} The uptake rate
77 of the target compounds (DDT metabolites) was calculated by adjusting the PRC depletion
78 rates by the differences in molar volumes and PE-water partition values (eq. 4).^{S1}

79 The calculated values of k_e for DDT metabolites ranged between 0.014 and 0.2 day^{-1} ,
80 where DDE compounds with higher K_{PE} values had a smaller k_e compared to DDD, which
81 had lower K_{PE} and higher k_e values. The equilibrium fraction of each DDT metabolite
82 compound in the PE ($f_{eq} = C_w(t = 0)/C_w(t = \infty)$) can be calculated as $(1 - e^{-k_e,DDT \cdot t})$
83 based on eq. 1. Therefore, the fraction to equilibrium for target DDT metabolite compounds
84 for a deployment of 93 days would range between 73–100% for a k_e range between 0.015 – 0.1
85 day^{-1} .

86 This estimate can be checked three ways:

87 **Method 1.** The PRC depletion rate of PCB155 used as a PRC in this study (93 day and
88 130 day deployment period) was lower than expected compared to the preliminary study (60
89 day deployment). During the April-July 2014 deployment period, PRC depletion in samplers
90 above the sediment was $11 \pm 3\%$, while below the sediment this was $21 \pm 8\%$.

91 The PRC exchange rate coefficient, $k_{e,PRC}$ in equation 3 was calculated to be between
92 $0.002 \pm 0.001 \text{ day}^{-1}$ for samplers on the pore water probe above and below the sediment.

93 Using the PRC relationship in equation 4, the average exchange rate ($k_{e,i}$) for 4,4'-DDD
94 0.03 ± 0.01 , which is three times slower than exchange rates calculated for the 2012 60-day
95 study. These values are also significantly less than values measured at a different site using
96 the PE samplers on stainless steel plates.^{S3} Therefore, we suspect that the depletion rates
97 measured in this study are too low, and perhaps there were other processes that hindered the
98 depletion of the heavier PRC from the PE. Nevertheless, using the conservative depletion
99 rates measured in this study, the DDD metabolites, which account for most of the pore water
100 concentration were within 90% of equilibrium for the 93-day deployment period. Since the
101 major metabolites were near equilibrium, the calculated aqueous concentration using either
102 method (with and without f_{adj}) showed similar results, where all values are within a factor of
103 two (Fig. S2). More hydrophobic metabolites with greater values of $\log K_{PE}$, like 4,4'-DDE,
104 were further from equilibrium where f_{adj} was on average 4.7 ± 2.4 .

105 **Method 2.** Another method of modeling uptake rate of target compounds into the PE
106 deployed in the sediment pore water is by modeling the diffusion of target compounds into
107 PE from diffusion out of the sediment pore water.^{S11} Parameters used in this model included
108 DDT diffusivity in PE,^{S6} K_{PE} (measured in this study), and sediment-water partition coef-
109 ficients K_d . Sediment-water partition coefficients were measured by combining wet sediment
110 (30 g archived grab sample from Pallanza Bay), PE (0.2 g of 51 μm thick PE), DI water (1
111 L), and sodium azide (1 ng/L) in a 1 L amber glass jar and allowing the system to come to
112 equilibrium on a shaker for at least one month (n=15). K_d values for DDT and its metabo-
113 lites were calculated by dividing sediment concentrations (C_{sed}) by PE concentrations (C_{PE})
114 and multiplying by PE partition-water partition values (K_{PE}). These measured values are
115 shown in Table S2.

116 The dimensionless parameter $T = \frac{t \cdot D_{PE}}{l^2}$ for 4,4'-DDD in a 51 μm thick PE deployed for
117 130 days is 168. D_{PE} (m^2/s) is the diffusivity of the target compound in PE, t = deployment
118 time period (s), and l is the PE thickness (m). By this approach, the calculated fraction of
119 equilibrium f_{eq} for DDT metabolites ranged between 0.80 – 0.97 for 130 days in 51 μm PE

Table S2: Average measured K_d (L/kg) in sediment samples collected from Pallanza Bay, standard deviations in parenthesis, n=15 (triplicate analysis of 5 different sediment samples)

Chemical	K_d (L/kg)
DDMU	3.4 (0.1)
2, 4'-DDE	4.4 (0.6)
4, 4'-DDE	4.3 (0.2)
2, 4'-DDD	4.0 (0.5)
4, 4'-DDD	3.9 (0.3)
2, 4'-DDT	4.9 (0.5)
4, 4'-DDT	4.4 (0.6)

120 sheets. For the DDD metabolite compounds which account for over 80% of the total freely
 121 dissolved DDT concentration in the pore water, $f_{eq} > 90\%$. The f_{eq} for all DDT metabolites
 122 was $> 80\%$.

123 **Method 3.** A third method of evaluating equilibrium in PE in situ measurements is
 124 by comparing concentrations measured during different deployment periods or by comparing
 125 samplers with different surface to volume ratio measurements.^{S12} Samplers deployed for
 126 longer time periods should measure higher concentrations that are closer to equilibrium
 127 until samplers converge when equilibrium is reached. Similarly, samples with higher surface
 128 to volume ratios should measure concentrations closer to equilibrium than samples with
 129 lower surface to volume ratios until equilibrium is reached. In Table S3, we compared
 130 calculated $C_w = C_{PE}/K_{PE}$ from different PE measurements at Site P2 in Pallanza Bay
 131 deployed in the overlying water. Measured DDD concentrations seem to be at equilibrium,
 132 since concentrations were consistent during April-July sampling period when samplers were
 133 either exposed one-sided or double-sided to the overlying water. Other DDT metabolites
 134 were not consistently measured in the overlying water.

135 Measured pore water concentrations measured at Site P2 during three different deploy-
 136 ment period and deployment lengths are shown in Table S4. This data supports conclusions
 137 that PE samplers are at equilibrium for DDD metabolites. Whether 4, 4'-DDE with higher
 138 K_{PE} value was at equilibrium is not entirely conclusive based on this comparison, but these
 139 concentrations represent a small fraction of total DDT concentrations.

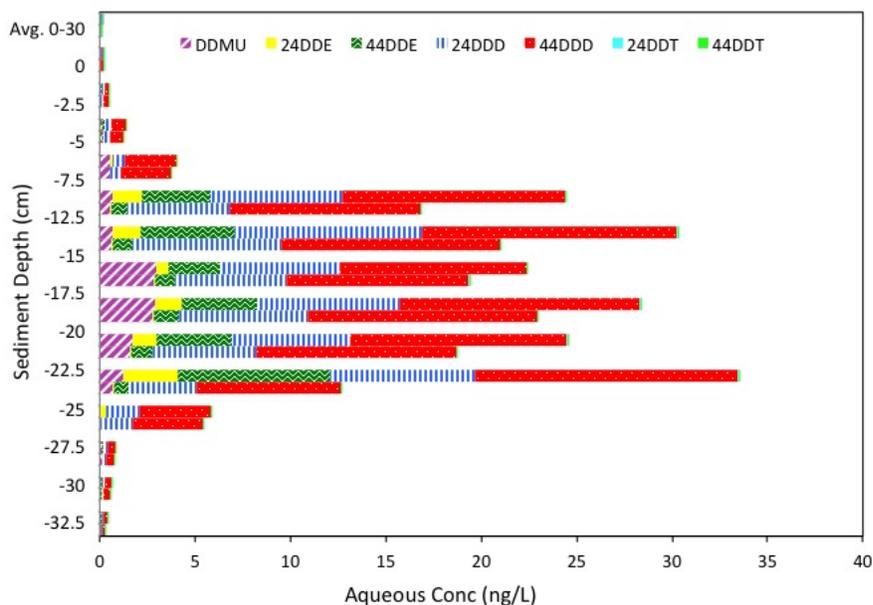
Table S3: Calculated $C_w = C_{PE}/K_{PE}$ from PE deployed in the overlying water for different deployment lengths (60 day (2012), 93 days (Apr-Jul 2014), 130 days (Jul-Nov 2014), and sampling configuration (single and double-sided exposure). Depth refers to height above sediment. Samplers attached to the pore water probe had only one side exposed to overlying water and pore water because samplers were wrapped tightly around the pore water probe. Double-sided samplers were attached to pore water probe frames with zipties so that both sides of the sampler were freely exposed to the surrounding water.

deployment period	60 days (double-sided)	93 days (single-sided)	93 days (double-sided)	130 days (single-sided)
depth (cm)	5-40	2.5-25	0-30	2.5-25
n	4	10	4	10
DDMU	< 0.01	0.03 (0.004)	0.01 (0.002)	0.02 (0.0008)
2,4'-DDE	< 0.01	< 0.01	< 0.01	< 0.01
4,4'-DDE	0.01 (0.002)	< 0.01	< 0.01	< 0.01
2,4'-DDD	0.02 (0.003)	0.02 (0.004)	0.02 (0.006)	0.05 (0.01)
4,4'-DDD	0.03 (0.004)	0.03 (0.004)	0.03 (0.008)	0.05 (0.02)
2,4'-DDT	< 0.01	< 0.01	< 0.01	< 0.01
4,4'-DDT	0.01 (0.001)	0.03 (0.006)	< 0.01	< 0.01

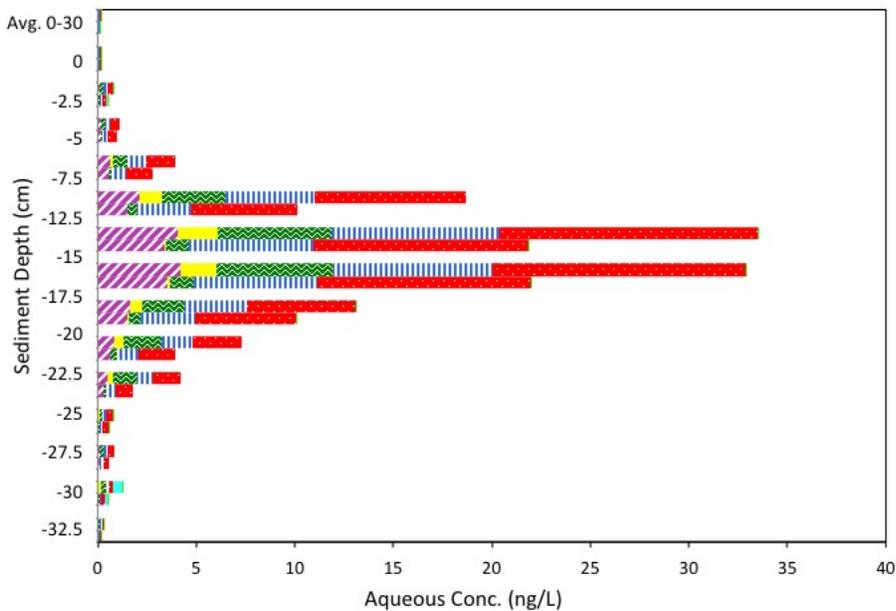
Table S4: Calculated $C_w = C_{PE}/K_{PE}$ from PE deployed in the pore water for different deployment lengths (Phase 1 = 60 days, Phase II = 93 days, Phase III = 130 days. Samplers were placed approximately two cm within the sediment. Samplers were attached to a stainless steel plate (63 days) or to the pore water probe (93 days and 130 days), so that samplers were only exposed on one side to the surrounding pore water DDT concentration.

	60 days (n=4)	93 days (n=1)	130 days (n=2)
DDMU	< 0.01	0.05	0.06
2,4'-DDE	0.004 (0.007)	< 0.01	< 0.01
4,4'-DDE	0.01 (0.002)	0.02	0.03
2,4'-DDD	0.1 (0.010)	0.1	0.1
4,4'-DDD	0.3 (0.019)	0.3	0.2
2,4'-DDT	< 0.01	< 0.01	< 0.01
4,4'-DDT	< 0.04	< 0.04	< 0.04

140 Based on these three methods of evaluating equilibrium in passive samplers, we conclude
 141 that the PE samplers were at or near equilibrium for DDD compounds for the 130 day de-
 142 ployment period, and the calculated freely dissolved aqueous concentrations in the overlying
 143 water and pore water concentrations are accurate within the error of the PE-water partition
 144 values, which is the largest source of uncertainty in the calculated concentrations.



(a) First deployment



(b) Second deployment

Figure S2: Aqueous concentration in pore water measured during both deployment periods assuming equilibrium (bottom bar of each pair) and with adjustment factor (f_{adj} from equations 2-4) (top bar of each pair). Measured DDT metabolites are 4, 4'-DDD (red, dots), 2, 4'-DDD (blue, vertical stripes), 4, 4'-DDE (green, squiggly), 2, 4'-DDE (yellow, solid), DDMU (purple, diagonal stripes).

145 Calculating diffusive flux in and from sediment

146 When considering diffusion in the overlying water, $D_{z,i}$ is equal to the molecular diffusivity
147 of the compound i (D_i). In the sediment pore water, the diffusivity will be reduced both by
148 less space available for diffusion (proportional to the porosity (ϵ) and longer paths for the
149 components to travel (tortuosity (τ) related to ϵ by $\tau = \epsilon^{-1/3}$). Hence in the pore water D_z
150 will be given by

$$D_z = \frac{\epsilon}{\tau} \cdot D = \epsilon^{4/3} \cdot D \quad (5)$$

151 Porosity was calculated from the water content in the sediment cores and diffusivities (D)
152 of the different DDT metabolites was calculate from molecular weights and adjusted for
153 temperature at $9^\circ C$.^{S13}

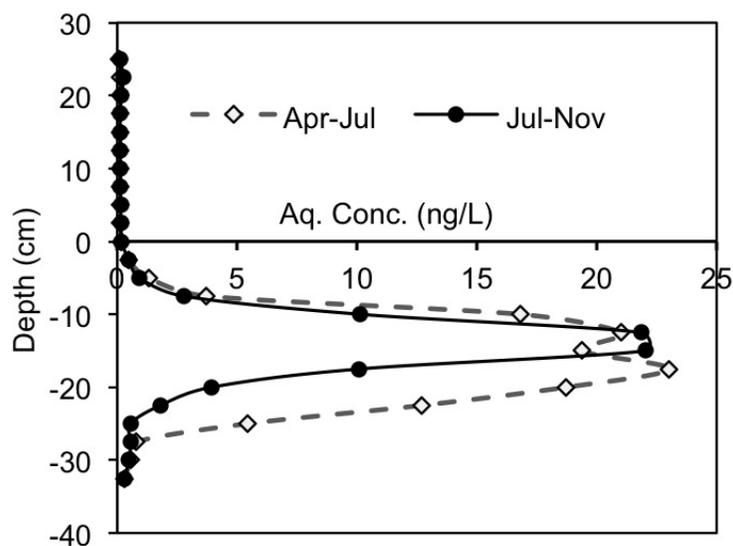
154 The diffusive flux, F can be calculated with the following equation:

$$F = D_{z,i} \left(\frac{C_{w_{z_1}} - C_{w_{z_2}}}{z_2 - z_1} \right) \quad (6)$$

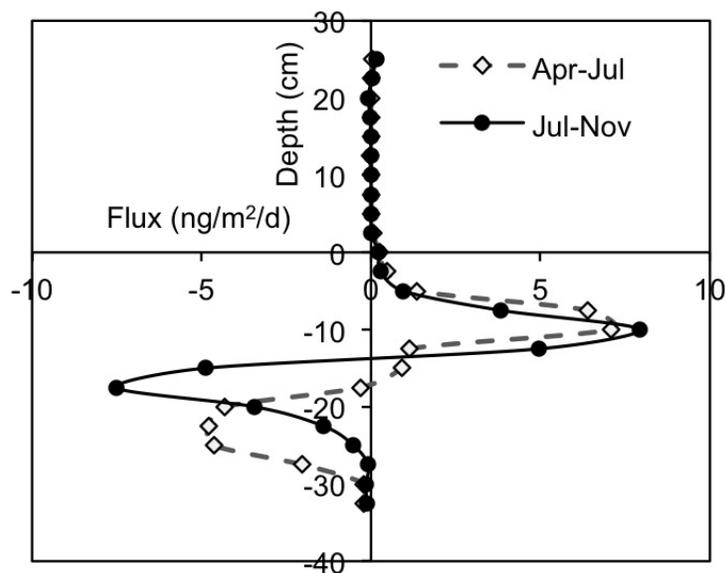
155 Where $D_{z,i}$ is the diffusivity of the compound of interest, i , at depth z . $C_{w_{z_1}}$ and $C_{w_{z_2}}$ is the
156 concentration in the aqueous phase at depths z_1 and z_2 .

157 A comparison of calculated flux from measured concentration profiles during the two
158 deployment periods is shown in Fig. S3(b). The shape of the flux profile is the same
159 between the two measurements. Difference in the concentration and flux profiles could be
160 due to heterogeneity in the sediment at slightly different locations. Another possibility is
161 cross contamination in the pore water below the concentration peak due to the penetration
162 of the probe.

163 Pore water profile measurements at other DDT-contaminated coastal sites in the U.S. and
164 China showed a similar shape where the freely dissolved aqueous concentrations were lower
165 above the sediment than within the pore water near the sediment surface.^{S14,S15} At those
166 sites, the pore water concentrations of the major DDT metabolites were much greater in the
167 other studies than in Pallanza Bay, which resulted in greater calculated diffusive flux from the



(a) Total DDT concentration profile



(b) Total DDT flux profile

Figure S3: Comparison of calculated freely dissolved aqueous concentration (a) and flux (b) of total DDT during first (open diamond, dotted line) and second deployment periods (filled circle solid line).

168 sediment bed by at least an order of magnitude. The peak concentration profile measured at
 169 depth in this study is unique in showing DDT contamination that occurred during a specific
 170 historical period (approximately between 1960s and 1980s based on geochronological dating
 171 of cores, unpublished data). The sharp peak in the historical record of the sediment DDT

172 concentration was also not blurred by sediment resuspension events since Pallanza Bay is
173 rather deep and mainly depositional with minimal resuspension.^{S16}

174 **Sediment Cores**

175 Cores were collected by Carma® Quality Coring. The top 30 cm of the core was sectioned
 176 at 1 - 5 cm intervals, and sectioned sediment samples were placed in separate 125 mL amber
 177 glass jars (I-CHEM 200 series) for storage and shipment. Sediment samples were analyzed for
 178 DDT and its metabolites using previously published methods^{S10} to measure total sediment
 179 concentrations.

Table S5: Sediment core concentration profile ($\mu\text{g}/\text{kg}$ dry).

Depth (cm)	DDMU	2,4'-DDE	4,4'-DDE	2,4'-DDD	4,4'-DDD	2,4'-DDT	4,4'-DDT	ΣDDT
0-1	0.6	0.6	4.7	3.6	3.2	0.4	1.7	15
1-2	1.3	0.7	5.3	4.5	5.6	0.4	1.4	19
2-3	3.0	1.4	7.4	7.5	13	0.4	2.6	35
3-4	1.7	1.5	5.5	6.0	13	0.4	4.7	32
4-6	5.4	7.0	24	14	28	1.2	3.6	82
6-8	16	16	64	31	65	1.8	19	212
8-10	< 0.5	154	424	220	651	0.3	54	1,505
10-15	124	224	833	597	1,330	12	61	3,190
15-20	< 0.5	5.9	12	19	19	0.4	2.4	60.

Table S6: Aqueous Concentration assuming equilibrium (ng/L water) for first deployment period (93 days).

Depth	DDMU	2, 4'-DDE	4, 4'-DDE	2, 4'-DDD	4, 4'-DDD	2, 4'-DDT	4, 4'-DDT	ΣDDT
25	0.02	< 0.0005	< 0.001	0.02	0.02	< 0.005	< 0.04	0.06
22.5	0.04	< 0.0005	< 0.001	0.02	0.02	0.005	< 0.04	0.08
20	0.03	< 0.0005	< 0.001	0.03	0.03	0.01	< 0.04	0.1
17.5	0.03	< 0.0005	< 0.001	0.02	0.03	< 0.005	< 0.04	0.08
15	0.03	< 0.0005	< 0.001	0.03	0.03	< 0.005	< 0.04	0.09
12.5	0.03	< 0.0005	< 0.001	0.02	0.03	< 0.005	< 0.04	0.08
10	0.03	< 0.0005	< 0.001	0.02	0.02	< 0.005	< 0.04	0.07
7.5	0.04	< 0.0005	< 0.001	0.02	0.02	< 0.005	< 0.04	0.08
5	0.03	< 0.0005	< 0.001	0.02	0.02	< 0.005	< 0.04	0.07
2.5	0.03	< 0.0005	0.003	0.03	0.03	< 0.005	0.05	0.1
0	0.03	< 0.0005	0.01	0.04	0.08	< 0.005	< 0.04	0.2
-2.5	0.05	< 0.0005	0.02	0.1	0.3	< 0.005	< 0.04	0.5
-5	0.13	0.002	0.05	0.3	0.7	< 0.005	0.06	1
-7.5	0.5	0.01	0.005	0.6	3	< 0.005	< 0.04	4
-10	0.6	0.07	0.8	5	10	0.01	< 0.04	17
-12.5	0.6	0.07	1	8	11	0.02	< 0.04	22
-15	3	0.06	1	6	10	0.02	< 0.04	19
-17.5	3	0.1	1	7	12	0.01	0.05	23
-20	2	0.08	1	5	11	0.02	< 0.04	19
-22.5	0.7	0.06	0.8	4	8	0.01	< 0.04	13
-25	< 0.02	0.03	0.01	2	4	0.01	< 0.04	5
-27.5	0.1	0.001	0.04	0.2	0.4	< 0.005	0.06	0.8
-30	0.07	< 0.0005	0.03	0.1	0.3	< 0.005	< 0.04	0.6
-32.5	0.05	< 0.0005	0.01	0.06	0.2	< 0.005	< 0.04	0.3
FB(n=3)	< 0.02	< 0.0005	< 0.001	< 0.008	0.004 ± 0.001	0.02 ± 0.003	0.02 ± 0.004	0.04

Table S7: Aqueous concentration assuming equilibrium (ng/L water) for second deployment period (130 days).

Depth	DDMU	2, 4'-DDE	4, 4'-DDE	2, 4'-DDD	4, 4'-DDD	2, 4'-DDT	4, 4'-DDT	ΣDDT
25	< 0.02	< 0.0005	0.01	0.04	0.04	0.01	< 0.04	0.1
22.5	0.02	0.001	0.01	0.08	0.1	0.01	< 0.04	0.2
20	0.02	< 0.0005	0.01	0.06	0.04	0.01	< 0.04	0.1
17.5	< 0.02	< 0.0005	0.01	0.04	0.04	0.01	< 0.04	0.1
15	< 0.02	< 0.0005	0.01	0.05	0.03	0.01	< 0.04	0.1
12.5	< 0.02	< 0.0005	0.01	0.04	0.03	0.01	< 0.04	0.09
10	< 0.02	< 0.0005	0.01	0.05	0.04	0.01	< 0.04	0.1
7.5	< 0.02	< 0.0005	0.01	0.05	0.04	0.01	< 0.04	0.1
5	< 0.02	< 0.0005	0.01	0.05	0.04	0.01	< 0.04	0.1
2.5	< 0.02	< 0.0005	0.01	0.05	0.04	0.01	< 0.04	0.1
0	< 0.02	< 0.0005	0.01	0.06	0.05	0.01	< 0.04	0.1
-2.5	0.06	0.001	0.03	0.2	0.2	0.01	< 0.04	0.5
-5	0.1	0.002	0.06	0.3	0.5	< 0.005	< 0.04	1
-7.5	0.5	0.01	0.2	0.8	1.3	< 0.005	< 0.04	3
-10	1	0.03	0.5	3	5	< 0.005	< 0.04	10
-12.5	3	0.09	1	6	11	< 0.005	< 0.04	22
-15	4	0.08	1	6	11	< 0.005	< 0.04	22
-17.5	2	0.04	0.6	3	5	< 0.005	< 0.04	10
-20	0.6	0.02	0.3	1	2	< 0.005	< 0.04	4
-22.5	0.3	0.01	0.1	0.5	0.9	< 0.005	< 0.04	2
-25	0.05	0.002	0.04	0.2	0.3	0.01	< 0.04	0.6
-27.5	0.07	0.002	0.04	0.2	0.3	< 0.005	< 0.04	0.6
-30	0.03	0.003	0.05	0.1	0.2	0.13	< 0.04	0.5
-32.5	< 0.02	0.001	0.01	0.06	0.07	< 0.005	< 0.04	0.1
FB (n=3)	< 0.02	< 0.0005	< 0.001	< 0.008	0.03 ± 0.001	0.01 ± 0.0002	0.03 ± 0.0006	0.07

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