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## Supplementary Information

### Occurrence and fate of nitrification and urease inhibitors in the aquatic environment

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69	Mean values with standard deviation are given if more than 50% of the results were >LOQ.	
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83 **Text S1:** Method development and validation

84 LC-ESI-MS/MS analysis with electrospray ionization in positive ionization mode was applied for  
85 the detection of nitrification and urease inhibitors (NUIs).

86 General interface parameters were: ion spray voltage: 5.5 kV; heater temperature: 600 °C; collision  
87 gas: medium; ion source gas 1/2: 60/70 psi and curtain gas: 40 psi. Analyst 1.6.1 software was used  
88 to record and evaluate the obtained chromatographic data. Liquid chromatography (LC) analysis  
89 was carried out using a 1290 HPLC system from Agilent Technologies (Waldbronn, Germany)  
90 equipped with a solvent cabinet, a micro vacuum degasser, a binary pump, a high performance  
91 autosampler with two 54 vial plates and a temperature-controlled column compartment.

92 Several reversed phase (RP) and hydrophilic interaction liquid chromatography (HILIC) columns  
93 were tested. However, some representatives of the analyzed NUIs are very polar compounds, which  
94 couldn't be retained by reversed phase liquid chromatography columns under the applied  
95 conditions. Promising results were obtained for *1H*-1,2,4-triazole and DCD with hydrophilic  
96 interaction liquid chromatography (HILIC), but the technique was only suitable for the most polar  
97 analytes, produced rather broad peaks and needed long re-equilibration time. Furthermore, for  
98 sufficient retention, HILIC technique requires are high percentage of organic solvent (acetonitrile)  
99 in the injected sample to prevent peak broadening. Such a solvent exchange is common when  
100 applying solid phase extraction (SPE), but results in higher limits of quantification due to the  
101 dilution with acetonitrile when a direct injection of water samples is applied (see below). As a  
102 consequence the above mentioned separation with a Hypercarb (Thermo Scientific, Waltham, USA)  
103 column was chosen. With this method the first target analyte *1H*-1,2,4-triazole elutes after 4,8 min.  
104 This is at least one minute after the void volume of the column, which was proven by the injection  
105 of thiourea.

106 After establishing a chromatographic method several SPE cartridges were tested for sample clean-  
107 up and pre-concentration of the analytes. For all tested materials (C18, styrene-divinylbenzene  
108 (SDB), activated carbon) the obtained recoveries

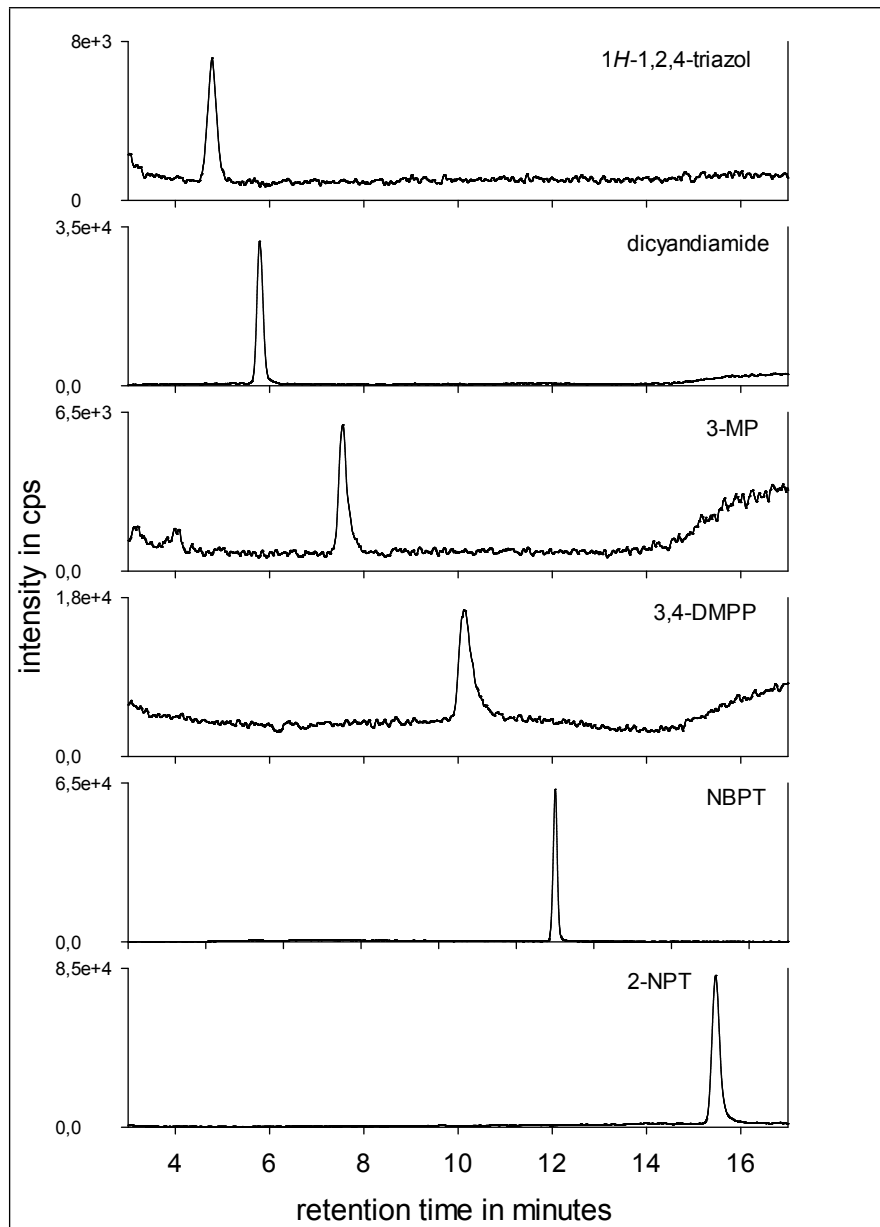
109 were insufficient, especially for 1*H*-1,2,4-triazole and DCD. To differentiate between break-through  
110 of the analytes and insufficient elution, the filtrates were collected and measured. The comparison  
111 of peak areas obtained from the eluate and from an external standard with the same concentration  
112 showed, that SPE sorbents material with hydrophobic interactions were not able to extract the  
113 compounds adequately. For 1*H*-1,2,4-triazole and DCD a break-through of more than 80% was  
114 observed. Schermerhorn et al. (2005) used a mixed-mode cationic exchange material for the  
115 extraction of 1*H*-1,2,4-triazol from water samples at neutral pH, although the compound is present  
116 uncharged at pH 7.<sup>1, 2</sup> A relative recovery of 94%, after correction with an isotopically labeled  
117 internal standard was reported, which allows no conclusion about the real extraction efficiency  
118 (absolute recovery). Applying this SPE protocol an absolute recovery of 38% was obtained for 1*H*-  
119 1,2,4-triazole and only 2% for DCD.

120 Based on unsatisfactory results of the SPE experiments, it was decided to directly inject the aqueous  
121 samples into the LC-MS/MS system.

122 To protect the LC auto-sampler and the analytical column samples from suspended solids, water  
123 samples were filtered using 0.45 µm-PTFE-membrane filters (Millipore, Billerica, U.S.A.) and  
124 1 mL BD Plastipack™ syringes (Becton Dickinson, Madrid, Spain) were used. Losses during  
125 filtration due to sorption were negligible.

126 For the assessment of compound specific limits of quantification (LOQs) two different approaches  
127 were applied for quality assurance. This procedure was chosen as qualifier transitions of 1*H*-1,2,4-  
128 triazol and DCD have poor signal intensities and can be used for confirmation purposes only at  
129 relatively high concentrations. First, LOQs were determined according to the calibration curve  
130 procedure according to German industry standard DIN 32 645 (level of confidence 95%) with an  
131 equidistant 10 level calibration. Secondly, the LOQ calculated according to DIN 32 645 was  
132 disallowed when the signal-to-noise ratio was <10. In that case, the lowest calibration point  
133 fulfilling this criterion was chosen as LOQ (Table S1).

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138 **Figure S1:** Extracted chromatograms of 50 ng/L standards in ultra-pure water (direct injection).  
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146 **Table S1:** Limits of quantification obtained by German industry standard DIN 32 645 and signal-  
 147 to-noise ratio.

values in µg/L	1 <i>H</i> -1,2,4-Triazol	DCD	3,4-DMPP	3-MP	NBPT	2-NPT
LOQ according to DIN 32645	0.03	0.02	0.01	0.01	0.01	0.01
S/N > 10 at	0.1	< 0.01	0.25	0.1	0.03	0.05

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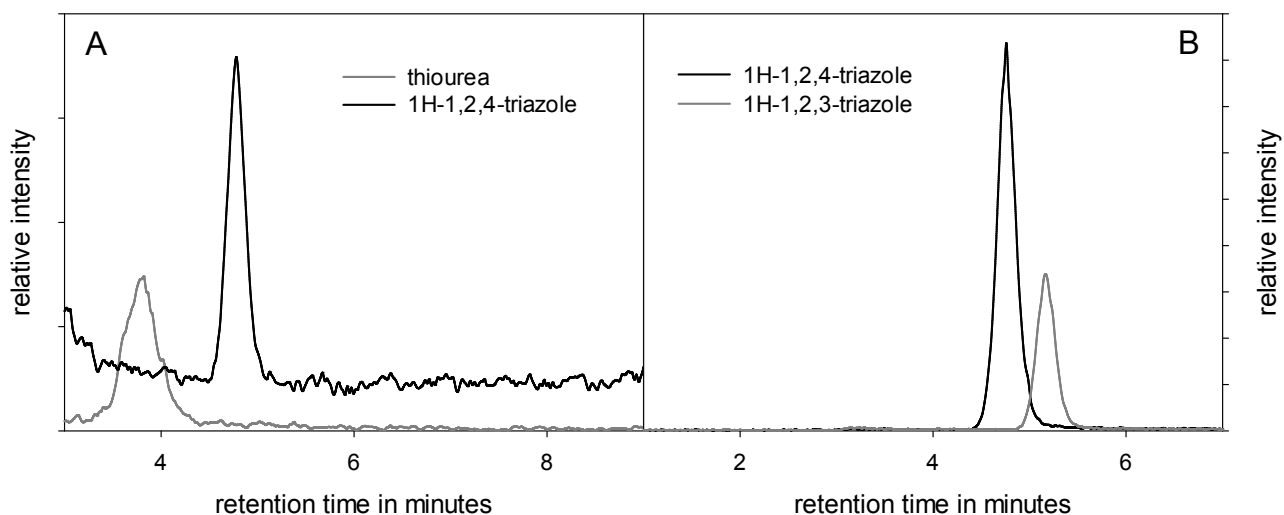
149 Matrix effect were assessed by comparing spiked potable water and surface water with standards of  
 150 the same concentration in ultra-pure water. Absolute recoveries for most compounds were in range  
 151 of 80% and 110%, except for 2-NPT.

152 MS/MS parameters for individual compounds were obtained by direct infusion into the MS  
 153 interface. The two most intensive MS/MS transitions between precursor ion and product ions were  
 154 used for identification and quantification in multiple reaction monitoring (MRM) mode. Precursor  
 155 and product ions, declustering potential, collision energy, and cell exit potential derived from  
 156 optimization of compound specific MS settings as well as general interface parameters are  
 157 displayed in Table S2.

158 **Table S2:** MS/MS parameters and limits of quantification in µg/L (in brackets) of target analytes  
 159 and internal standards, MS-system SCIEX Triple Quad™ 5500.

Analyte	Precursor ion	Product ions	DP <sup>a</sup>	CE <sup>b</sup>	CXP <sup>c</sup>
1 <i>H</i> -1,2,4-triazole (0.1)	70.0	43.0	150	27	8
		28.1	150	41	10
DCD (0.02)	85.0	68.0	66	27	6
		42.9	66	21	20
3,4-dimethylpyrazol (0.25)	97.0	56.1	116	23	10
		42.0	116	37	18
3-methylpyrazol (0.1)	83.0	42.0	156	25	8
		56.1	156	23	10
NBPT (0.03)	168.1	95.0	56	25	10
		74.0	56	15	10
2-NPT (0.05)	217.0	199.9	51	13	16
		121.0	51	23	18
<sup>15</sup> N <sub>4</sub> -DCD	89.0	71.0	86	25	12
1 <i>H</i> -1,2,4-triazole- <sup>13</sup> C <sub>2</sub> , <sup>15</sup> N	73.0	44.0	16	13	14

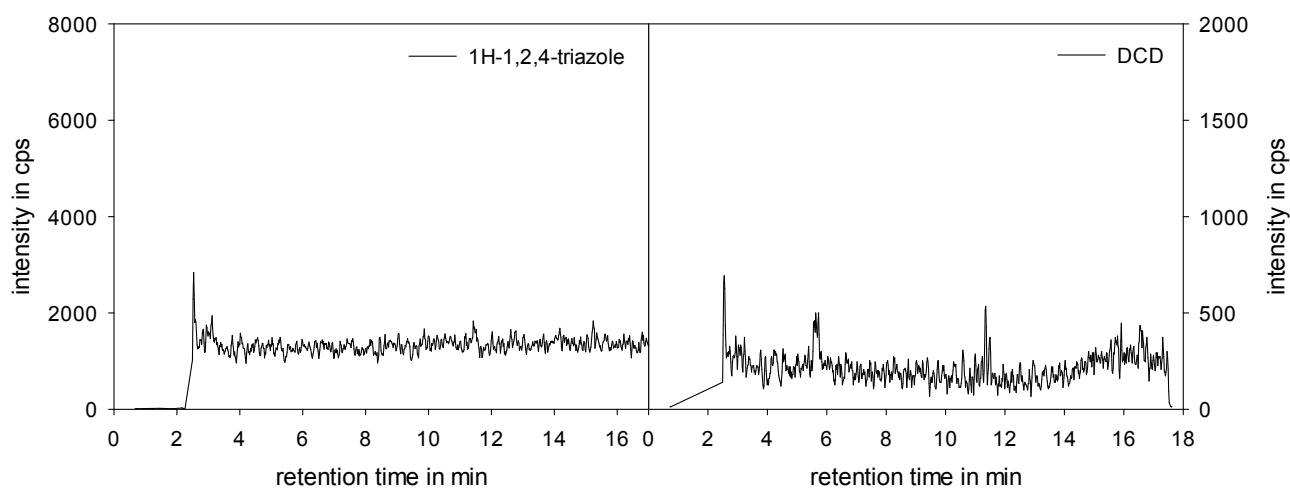
160 <sup>a</sup> DP = declustering potential in volt, <sup>b</sup> CE = collision energy in electron volt, <sup>c</sup> CXP = cell exit potential in volt



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163 **Figure S2:** Quality measures during method development. Extracted chromatograms of thiourea  
164 and 1H-1,2,4-triazole proving retention of the latter (A) and extracted 1H-1,2,4-triazole and 1H-  
165 1,2,3-triazole proving their different retention times (B).

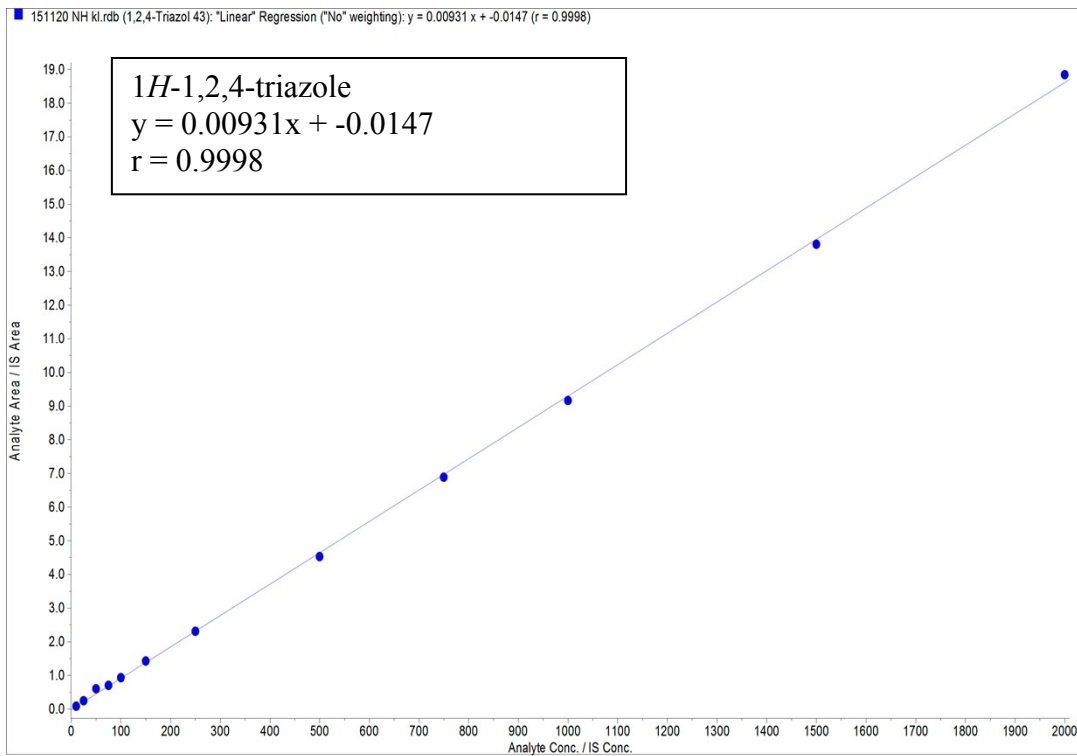
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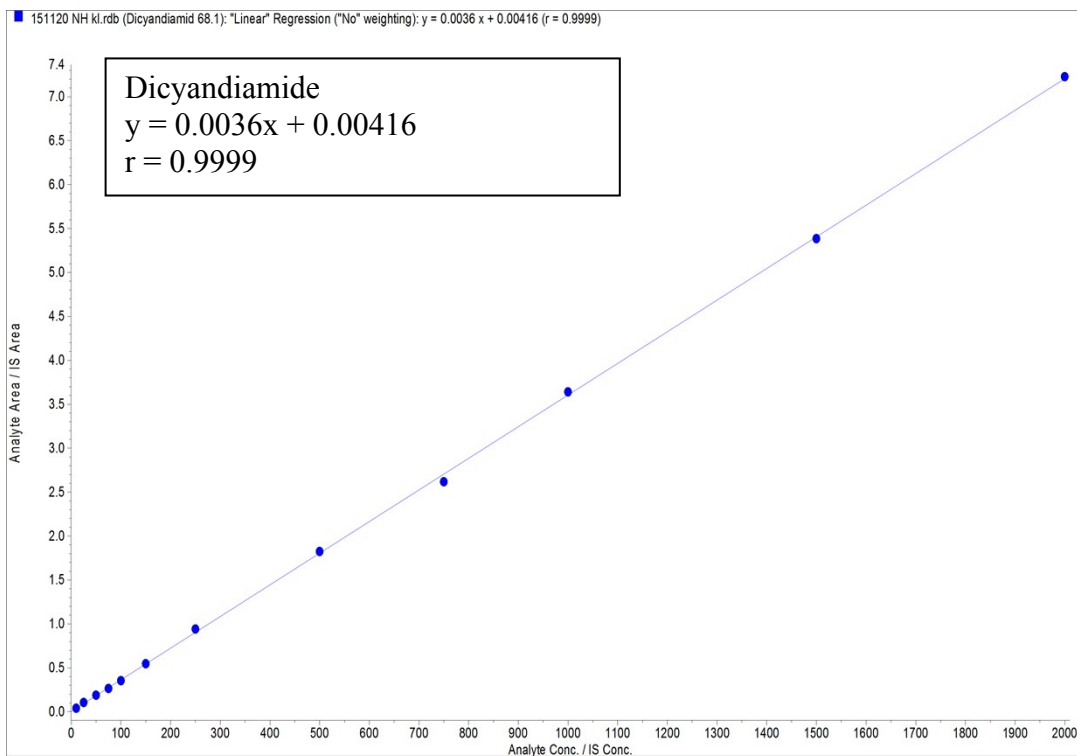
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169 **Figure S3:** Extracted chromatograms of 1H-1,2,4-triazole (left) and DCD (right) for a blank  
170 sample. Note, the different scaling when comparing it with 50 ng/L standard in Figure S1.

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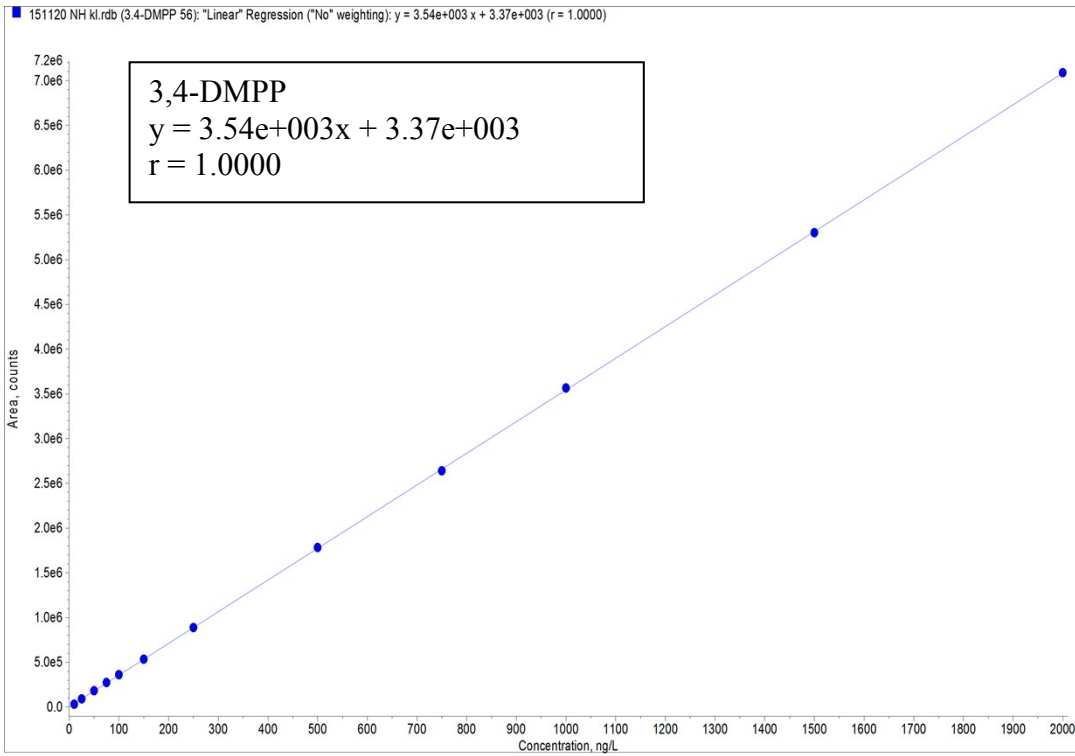
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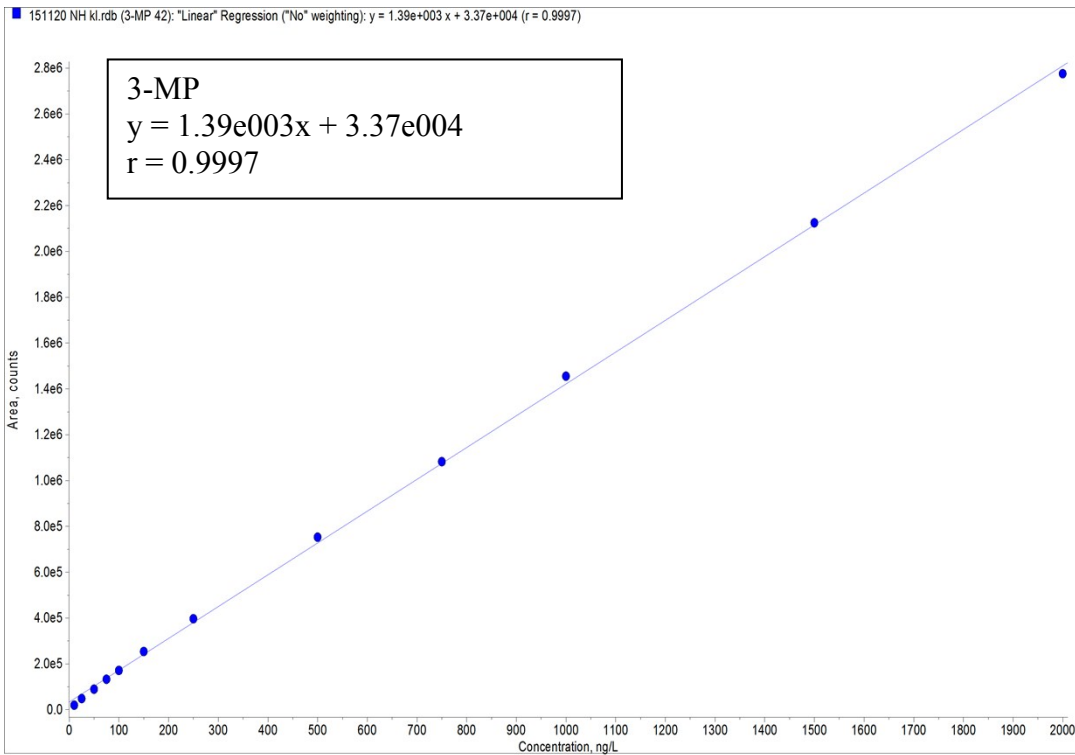
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174 **Figure S4:** Calibration curves, linear equations and regression coefficients for all NUIs under  
 175 investigation.





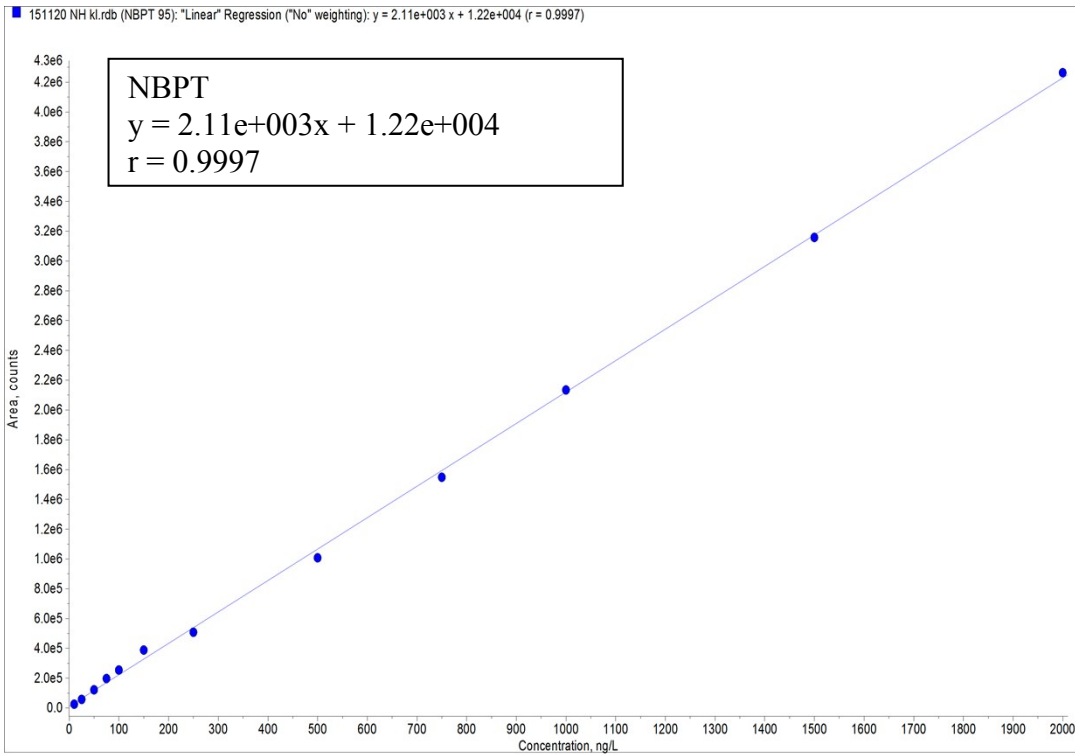
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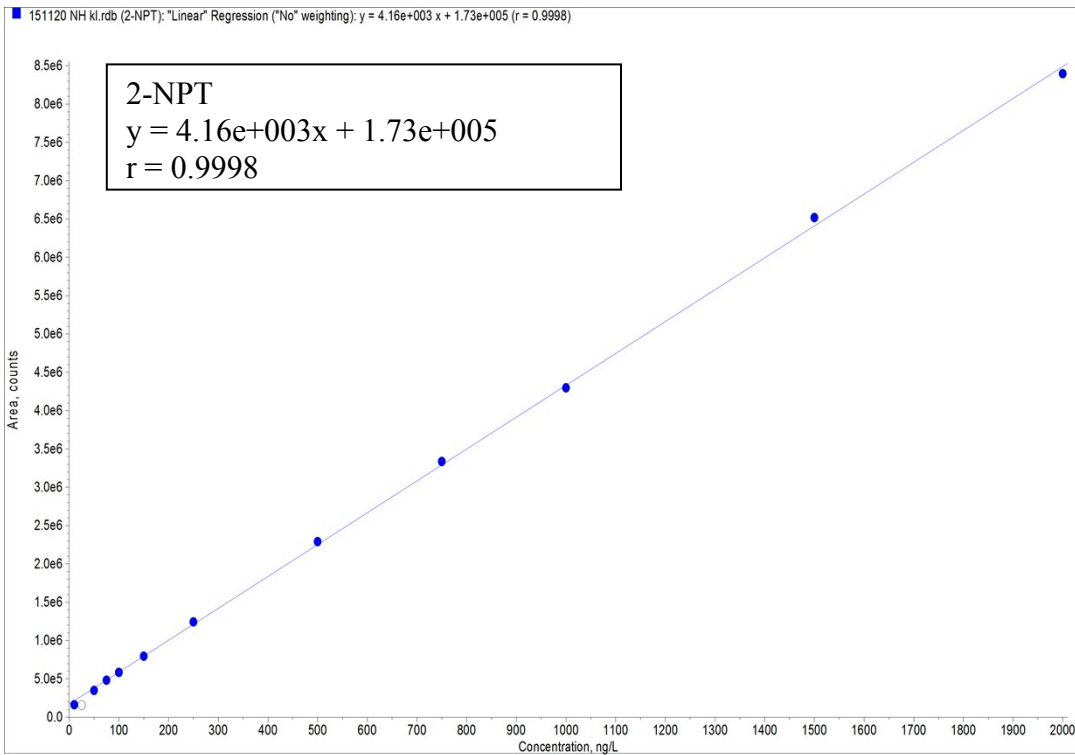
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178 **Figure S4:** continued

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182 **Figure S4:** continued

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189 **Table S3:** Proof of repeatability of the method by ten injection of an analytical standard.

	1H-1,2,4-triazol	DCD	3.4-DMPP	3-MP	NBPT	2-NPT
	peak area					
500 ng/L, injection 1	6,13E+05	3,22E+06	2,44E+06	7,13E+05	4,56E+06	1,65E+07
500 ng/L, injection 2	5,83E+05	3,25E+06	2,31E+06	7,27E+05	3,00E+06	1,51E+07
500 ng/L, injection 3	5,82E+05	3,23E+06	2,31E+06	8,68E+05	3,05E+06	1,52E+07
500 ng/L, injection 4	5,92E+05	3,12E+06	2,37E+06	7,09E+05	3,09E+06	1,50E+07
500 ng/L, injection 5	5,75E+05	3,23E+06	2,35E+06	7,26E+05	4,04E+06	1,51E+07
500 ng/L, injection 6	5,66E+05	3,00E+06	2,28E+06	7,02E+05	4,06E+06	1,46E+07
500 ng/L, injection 7	5,53E+05	3,10E+06	2,35E+06	8,48E+05	3,93E+06	1,45E+07
500 ng/L, injection 8	5,49E+05	3,03E+06	2,35E+06	7,03E+05	4,11E+06	1,40E+07
500 ng/L, injection 9	5,54E+05	3,24E+06	2,33E+06	7,56E+05	3,14E+06	1,38E+07
500 ng/L, injection 10	6,34E+05	3,22E+06	2,42E+06	7,22E+05	3,18E+06	1,41E+07
mean value	5,80E+05	3,16E+06	2,35E+06	7,47E+05	3,62E+06	1,48E+07
standard deviation	2,60E+04	8,89E+04	4,68E+04	5,74E+04	5,48E+05	7,44E+05
standard deviation in %	4.5	2.8	2.0	7.7	15.2	5.0

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192 **Text S2:** Analysis of other micropollutants within monitoring programs 3 and 4.

193 Artificial sweeteners acesulfame and sucralose were analyzed according to Scheurer et al., (2009).<sup>3</sup>

194 Briefly, 50 mL of the respective water sample were enriched at pH 3 with a styrol-divinylbenzene

195 cartridges (Bakerbond SDB 1, 200 mg/6 mL from J.T. Baker, Deventer, The Netherlands). After

196 drying of the sorbent material, analytes were eluted with 3 x 3 mL of methanol. The eluate was

197 blown down to dryness and reconstituted with 500 µL buffer A and B (A: ultra-pure water, B:

198 methanol, both with 2 mM ammonium acetate). Chromatographic separation was achieved with a

199 Zorbax Eclipse XDB-C8 column (2.1 mm x 150 mm, 3.5 µm; Agilent Technologies, Santa Clara,

200 CA, USA). Initial conditions of the gradient program were 90% buffer A, which was held for 2 min,

201 then decreased to 25% within 4 min, held for 5 min and then increased again to the initial

202 conditions within 1 min. After each sample run the column was re-equilibrated for 7 min. Liquid

203 chromatography was carried out using a model 1290 HPLC system from Agilent Technologies. The

204 HPLC system was connected to an API 5500 triple-quadrupole mass spectrometer (Applied

205 Biosystems/ MDS Sciex Instruments, Concord, ON, Canada) with an electrospray interface

206 operated in negative ionization mode. Sucralose-d<sub>6</sub> and acesulfame-d<sub>4</sub> were used for internal

207 standard calibration.

208 Carbamazepine, gabapentin, primidone, metformin, valsartan acid, diatrizoate, hydrochlorothiazide,  
209 and oxypurinol were analyzed by direct injection of 40  $\mu$ L via a MPS-DualHead-WorkStation  
210 (Gerstel, Mülheim a. d. Ruhr, Germany) onto the column. Chromatographic separation was  
211 achieved with a Kinetex EVO C18 column (2.1 mm x 150 mm, 5  $\mu$ m; Phenomenex, Aschaffenburg,  
212 Germany) using ultra-pure water (A) and methanol (B), both containing 1 mM ammonium acetate,  
213 as eluents. Initial conditions of the gradient program were 98% eluent A, which was held for 5 min,  
214 then decreased to 25% within 20 min, held for 9 min and then increased again to the initial  
215 conditions within 1 min. After each sample run the column was re-equilibrated for 7 min. The  
216 HPLC system was connected to an API 5500 triple-quadrupole mass spectrometer  
217 Carbamazepine-d<sub>10</sub>, gabapentin-d<sub>4</sub>, metformin-d<sub>6</sub>, diatrizoate-d<sub>6</sub>, hydrochlorothiazide-<sup>13</sup>C,<sub>d</sub><sub>2</sub> and  
218 oxypurinol-<sup>13</sup>C,<sup>15</sup>N<sub>2</sub> were used for internal standard calibration. Primidone and valsartan acid were  
219 analyzed without internal standards.

220 Analysis of ethylenediaminetetraacetic acid (EDTA) was performed according to DIN EN ISO  
221 16588:2004-02.<sup>4</sup> Heptadecanoic acid nitrile was used as an internal standard for gas  
222 chromatography. 1,2-Diaminopropane-*N,N,N',N'*-tetraacetic acid was used as an internal standard  
223 for the whole analytical protocol.

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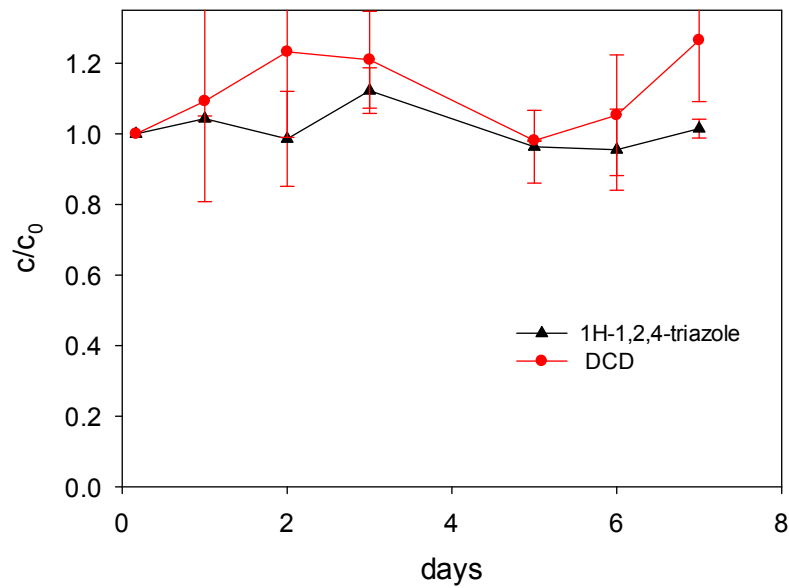
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233 **Table S4:** Characterization of standard soils LUFA 2.1 and LUFA 2.4 used for sorption study.

standard soil	2.1	2.4
sampling date	12.06.2013	10.06.2013
content of organic carbon in % C	0.66 ± 0.10	2.21 ± 0.46
pH in 0,01 M CaCl <sub>2</sub>	5.2 ± 0.3	7.2 ± 0.2
cation exchange capacity in meq/100 g	4.1 ± 0.6	32.2 ± 4.4
particle size distribution in % according to USDA		
<0.002 mm	2.4 ± 0.4	26.5 ± 1.9
0.002 – 0.006 mm	1.6 ± 0.4	8.1 ± 1.0
0.006 – 0.02 mm	3.6 ± 0.4	14.8 ± 1.1
0.2 – 0.063 mm	7.0 ± 0.5	23.0 ± 1.0
0.063 – 0.2 mm	27.2 ± 0.5	19.0 ± 0.3
0.2 – 0.63 mm	55.7 ± 1.5	6.9 ± 2.2
0.63 – 2.0 mm	2.5 ± 0.4	1.7 ± 0.2
soil type	sand	loam

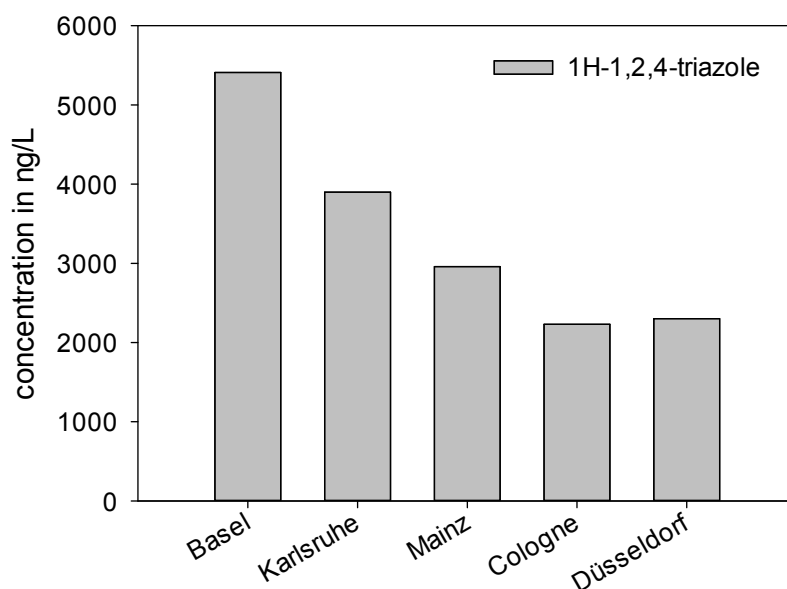
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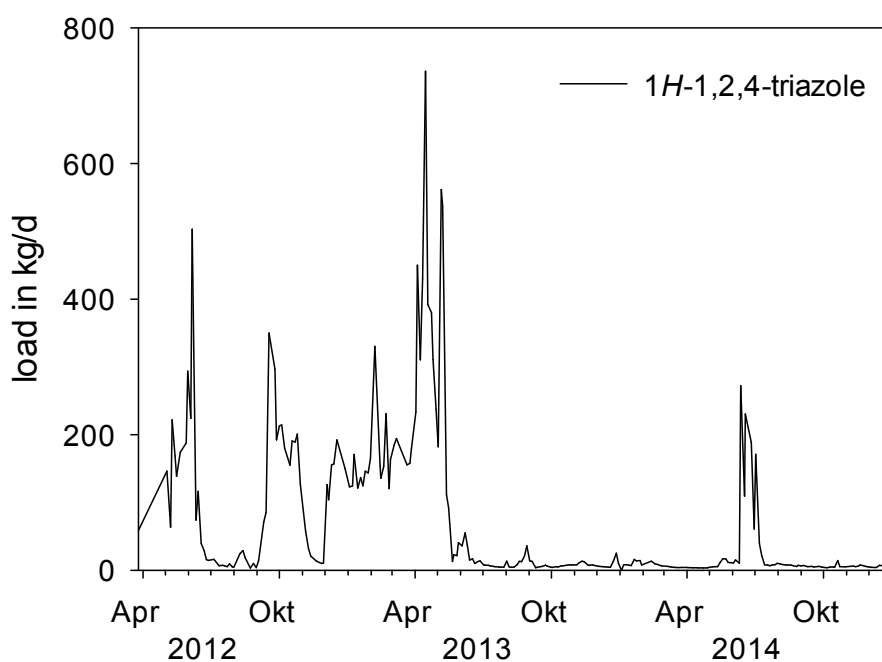
239 **Figure S5:** Behavior of 1H-1,2,4-triazole und DCD in batch tests with activated sludge (modified  
240 Zahn-Wellens test), test duration 7 d.

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242

243 **Figure S6:** Concentration of 1*H*-1,2,4-triazole in the River Rhine, samples taken at the same day in  
 244 April 2013.



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246 **Figure S7:** Mass load of 1*H*-1,2,4-triazole in the River Rhine at the sampling point in Cologne  
 247 based on river discharge and measured concentration; samples taken between March 2012 and  
 248 December 2014.

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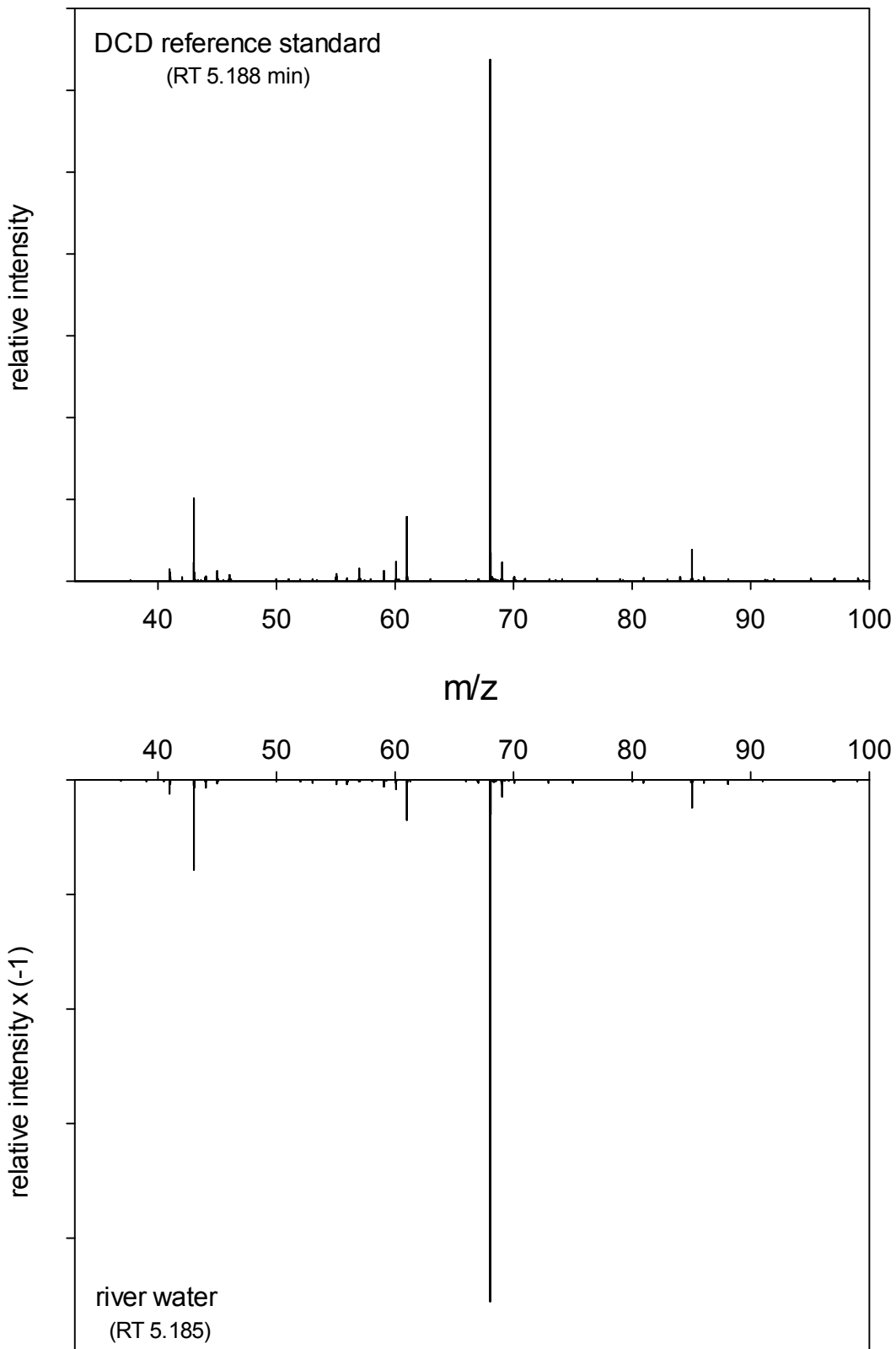
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253 **Text S3:** HRMS measurement for confirmation of high DCD concentrations

254 For further confirmation purposes of high DCD concentrations within monitoring program 2, LC  
255 high resolution mass spectrometry measurements (HRMS) were performed with a Q-TOF 6540  
256 (Agilent Technologies, Waldbronn, Germany). MS/MS spectra of a surface water sample close to  
257 the potential discharger was compared with that of a reference standard of DCD. Further  
258 confirmation of was given by the Federal Institute of Hydrology (BfG, Koblenz, Germany) by  
259 applying a LC/HRMS method with a different chromatography and MS system. The MS/MS  
260 spectra which were obtained at the retention time of DCD were similar and showed the same  
261 product ions as used in the quantitation method (Figure S8). The spectra corresponds also well with  
262 the one available from the National Institute of Standards and Technology (NIST) database,  
263 (<http://webbook.nist.gov/cgi/cbook.cgi?ID=C461585&Mask=200#Mass-Spec>) although a different  
264 fragmentation mechanism is used. According to Schymanski et al., 2014 an identification level 1,  
265 can be assigned to the observation of DCD in the surface water sample.<sup>5</sup>



266

267 **Figure S8:** High resolution MS/MS mass spectra of a DCD reference standard and a river water  
268 sample close to the point of discharge (monitoring program 2) obtained with a Agilent Q-TOF  
269 6540. Additional confirmation by BfG Koblenz, see Text S3.



270 **Table S5A:** Correlation coefficients of micropollutants considering all sampling points of  
 271 monitoring program 3 (River Main and its tributaries). Numbers in bold indicate three  
 272 micropollutants with the highest sum of correlation coefficients.

	CBZ	GAB	PRIM	MET	VAC	DCD	EDTA	SUC	ACE	DTZ	OXY	sum
CBZ	X	0.82	0.72	0.70	0.83	0.21	0.15	0.75	0.65	0.49	0.57	5.89
GAB	0.82	X	0.66	0.76	0.70	0.17	0.23	0.77	0.66	0.38	0.54	5.69
PRIM	0.72	0.66	X	0.52	0.82	0.40	0.33	0.73	0.60	0.67	0.79	<b>6.24</b>
MET	0.70	0.76	0.52	X	0.55	0.30	0.3	0.59	0.70	0.26	0.55	5.23
VAC	0.83	0.70	0.82	0.55	X	0.39	0.20	0.89	0.53	0.73	0.78	<b>6.42</b>
DCD	0.21	0.17	0.40	0.30	0.39	X	0.62	0.41	0.10	0.42	0.70	3.72
EDTA	0.15	0.23	0.33	0.30	0.20	0.62	X	0.28	0.31	0.36	0.52	3.30
SUC	0.75	0.77	0.73	0.59	0.89	0.41	0.28	X	0.41	0.61	0.73	6.17
ACE	0.65	0.66	0.60	0.70	0.53	0.10	0.31	0.41	X	0.31	0.42	4.69
DTZ	0.49	0.38	0.67	0.26	0.73	0.42	0.36	0.61	0.31	X	0.75	4.98
OXY	0.57	0.54	0.79	0.55	0.78	0.70	0.52	0.73	0.42	0.75	X	<b>6.35</b>

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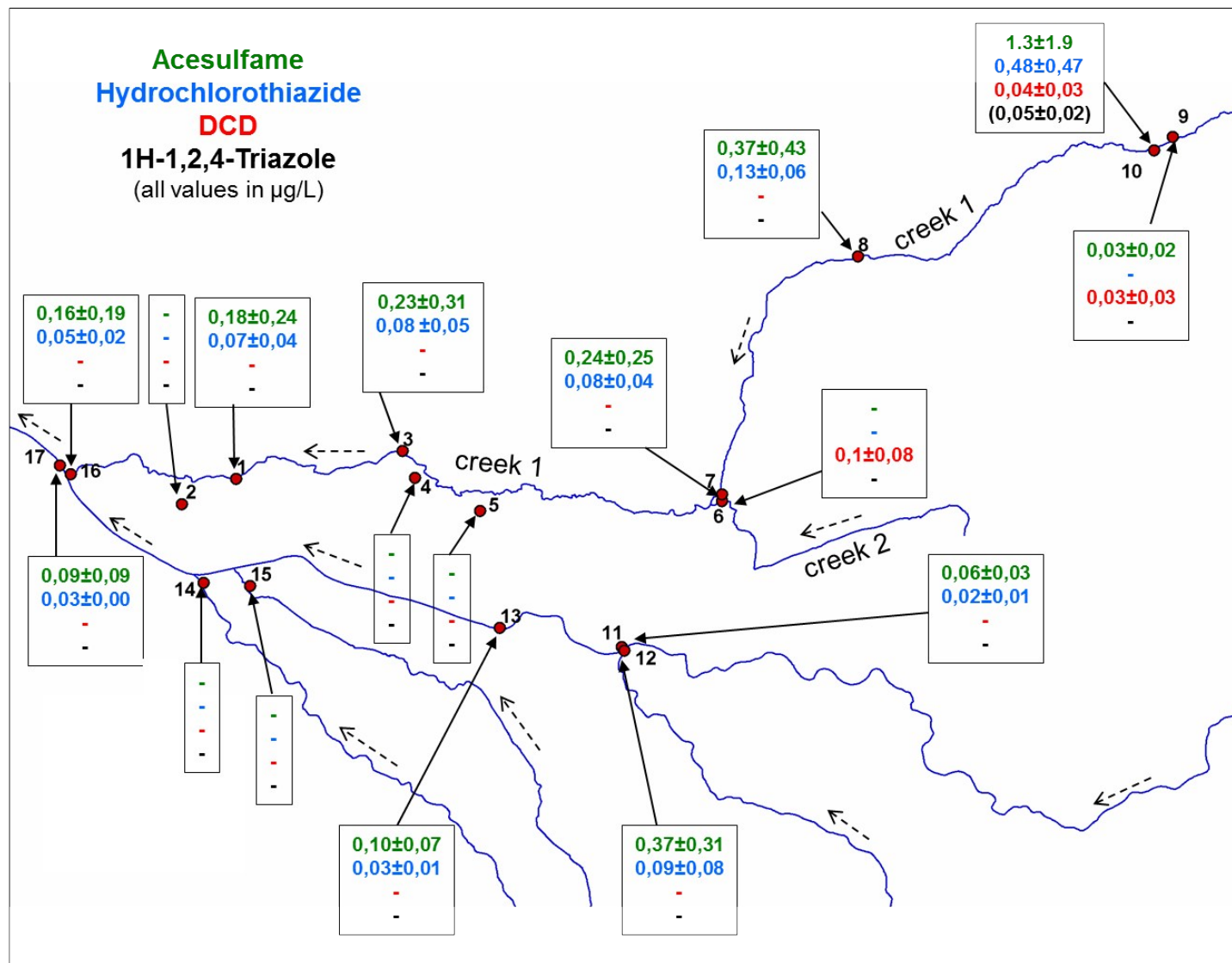
274 **Table S5B:** Correlation coefficients of micropollutants considering sampling points of monitoring  
 275 program 3 (River Main only, tributaries were excluded). Numbers in bold indicate three  
 276 micropollutants with the highest sum of correlation coefficients.

	CBZ	GAB	PRIM	MET	VAC	DCD	EDTA	SUC	ACE	DTZ	OXY	sum
CBZ	X	0.13	0.50	0.30	0.53	0.02	0.04	0.03	0.51	0.12	0.50	2.68
GAB	0.13	X	0.21	0.07	0.33	0.05	0	0.43	0.28	0	0.30	1.80
PRIM	0.50	0.21	X	0.06	0.75	0	0.02	0.05	0.65	0.10	0.40	2.74
MET	0.30	0.07	0.06	X	0.16	0.21	0.03	0.32	0.03	0.16	0.33	1.67
VAC	0.53	0.33	0.75	0.16	X	0.04	0	0.14	0.52	0.07	0.67	<b>3.21</b>
DCD	0.02	0.05	0	0.21	0.04	X	0.08	0.13	0.12	0.15	0.05	0.85
EDTA	0.04	0	0.02	0.03	0	0.08	X	0.06	0	0.01	0.03	0.27
SUC	0.03	0.43	0.05	0.32	0.14	0.13	0.06	X	0	0.23	0.13	1.52
ACE	0.51	0.28	0.65	0.03	0.52	0.12	0	0	X	0.33	0.37	<b>2.81</b>
DTZ	0.12	0,00	0.10	0.16	0.07	0.15	0.01	0.23	0.33	X	0	1.17
OXY	0.50	0.30	0.40	0.33	0.67	0.05	0.03	0.13	0.37	0	X	<b>2.78</b>

277

278 **Table S6:** Concentrations of 1H-1,2,4-triazole and DCD in several WWTP effluents as well as  
 279 population equivalents of the WWTPs.

WWTP	population equivalents (treatment capacity)	1H-1,2,4-triazole concentration in µg/L	DCD
WWTP 1	55,000	< 0.5	0.37
WWTP 2	120,000	< 0.5	0.45
WWTP 3	250,000	< 0.5	2.9
WWTP 4	7,500	< 0.5 / <0.5	< 0.1 / < 0.1
WWTP 5	725,000	0.84	1.0
WWTP 6	14,300	< 0.5	0.18
WWTP 7	15,000	< 0.5	< 0.1
WWTP 8	15,000	< 0.5	0.27



1

2 **Figure S9:** Concentrations of acesulfame, hydrochlorothiazide, DCD, and 1H-1,2,4-triazole measured in the catchment of a water utility (monitoring  
 3 program 4). Mean values with standard deviation are given if more than 50% of the results were >LOQ. Dashed arrows indicate flow directions.

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