## **Supplementary Information**

## Changes in Dissolved Organic Matter during the Treatment Processes of a Drinking Water Plant in Sweden and Formation of Previously Unknown Disinfection Byproducts

Michael Gonsior<sup>a\*</sup>, Philippe Schmitt-Kopplin<sup>b,c</sup>, Helena Stavklint<sup>d</sup>, Susan D. Richardson<sup>e</sup>, Norbert Hertkorn<sup>b</sup> and David Bastviken<sup>f</sup>

<sup>a</sup> Chesapeake Biological Laboratory , University of Maryland Center for Environmental Science, Solomons, USA

<sup>b</sup> Helmholtz Zentrum München, Analytical BioGeoChemistry, Neuherberg, Germany

<sup>c</sup> Technische Universität München, Analytical Food Chemistry, D-85354 Freising-Weihenstephan, Germany

<sup>d</sup> Tekniska verken i Linköping AB, Sweden

<sup>e</sup> University of South Carolina, Department of Chemistry and Biochemistry, Columbia, SC, USA

<sup>f</sup> Linköping University, Department of Thematic Studies – Environmental Change, Linköping, Sweden

\*Corresponding author. phone: +14103267245, fax: +14103267302, Email address:

gonsior@umces.edu

## Summary

Table S1	Page S2
Table S2	Page S3
Table S3	Page S4
Figure S1	Page S5
Figure S2	Page S6

spectrum	Figure	NS	AQ [s]	D1 [s]	WDW2	PR2
<sup>1</sup> H NMR						
before / after Cl	6	512	5	5	EM	1
<sup>13</sup> C NMR						
before / after Cl	S2	9361 / 9456	1	19	EM	35 (2)
DEPT 45/135 <sup>13</sup> C NMR						
before / after Cl	S2	16384	1	2	EM	12.5
DEPT 90 <sup>13</sup> C NMR						
before / after Cl	S2	32768	1	2	EM	12.5

**Table S1:** Acquisition parameters of NMR spectra, shown according to figures. NS: number of scans (for 2D NMR: F2); AQ: acquisition time [ms]; D1: relaxation delay [ms]; NE: number of F1 increments in 2D NMR spectra; WDW1, WDW2: apodization functions in F1/ F2 (EM: line broadening factor [Hz]; PR2: coefficients used for windowing functions WDW2.

δ( <sup>1</sup> H) [ppm]	10 - 7.0	7.0 - 5.3	4.9 - 3.1	3.1 - 1.9	1.9 - 0.0	H <sub>olefinic</sub> / H <sub>aromatic</sub>	10 - 5.3 $(\underline{H}C_{sp}^{2})$
key substructures	<u>H</u> ar	<u><b>H</b></u> C=C, <u><b>H</b>CO<sub>2</sub></u>	<u>H</u> CO	<u> Н</u> С-N, <u>Н</u> С-С-Х	<u>H</u> C-C-C-	1 aromatic	
before Cl	2.4	2.1	19.2	28.7	47.6	0.9	4.5
after Cl	3.0	2.7	19.3	29.1	46.0	0.9	5.7

 Table S2:
 <sup>1</sup>H NMR section integrals (percent of non-exchangeable protons) and key substructures of DOM before and after chlorination.

δ( <sup>13</sup> C) ppm	220-18	7 187	-167	167-145	145-108	108-90	90-47	47-0	H/C ratio	O/C ratio	
Key substructures	<u>C</u> =O	<u>C</u>	XC	<u>C</u> ar-O	<u>C</u> ar-C,H	О <sub>2</sub> <u>С</u> Н	О <u>С</u> Н	С <u>С</u> Н			
before chlorination	1.8		9.8	3.9	8.4	5.6	33.7	36.8	1.31	0.70	
after chlorination	1.7	9	Э.1	3.8	10.8	5.6	32.1	35.5	1.28	0.67	
NMR mixing model	C=O	CO	ОН	C <sub>ar</sub> -O	C <sub>ar</sub> -H	O <sub>2</sub> CH	ОСН	CH <sub>2</sub>			
H/C ratio	0		1	0	1	1	1	2			
O/C ratio	1		2	1	0	2	1	0			
DOM (depth)	CH total	CH <sub>2</sub> total	CH <sub>3</sub> total	ratio $(d_1 / c_1 / b_1 / a_1)$ HCar-C / O-HC-O / HC-O / HC-C			ratio (b <sub>2</sub> / a <sub>2</sub> ) H <sub>2</sub> C-O / H <sub>2</sub> C-C		ratio (b <sub>3</sub> / a <sub>3</sub> ) H <sub>3</sub> <u>C</u> -O / H <sub>3</sub> <u>C</u> -C		
before chlorination	36	30	34	15.9	/ 1.0 / 33.9 / 49.1		6.8	6.8 / 93.2		9.1 / 90.9	
after chlorination	35	33	32	14.8 / 1.4 / 34.5 / 49.3		10.7	10.7 / 89.3		5.7 / 94.3*		

**Table S3.** (Top): <sup>13</sup>C NMR section integrals (percent of total carbon) and key substructures of DOM before and after chlorination. Middle: Substructures used for NMR-derived reverse mixing model with nominal H/C and O/C ratios given. Bottom: percentage of methin, methylene and methyl carbon related to total protonated <sup>13</sup>C NMR integral as derived from <sup>13</sup>C DEPT NMR spectra of DOM according to carbon multiplicity (left 3 columns) and relative proportions of these CH<sub>n</sub> units binding to oxygen versus carbon chemical environments.

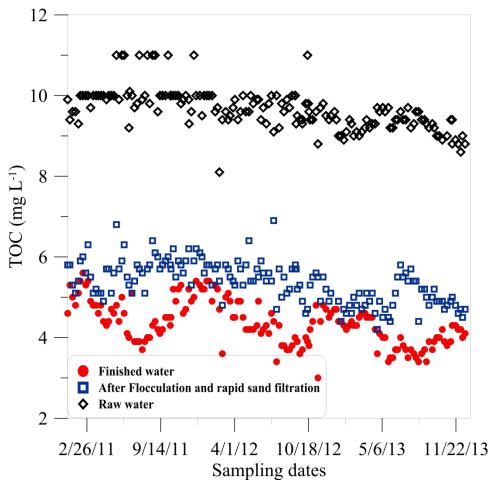
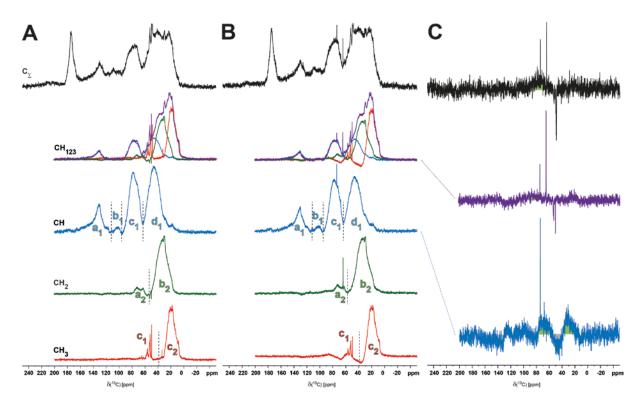


Figure S1: Total organic carbon concentration of the raw water, after flocculation/rapid sand filtration, and of the processed water, Råberga drinking water treatment plant, Linköping, Sweden.



**Figure S2:** <sup>13</sup>C NMR spectra of Raberga slow sand SPE-DOM (<sup>12</sup>CD<sub>3</sub>OD solution,  $B_0 = 11.7 \text{ T}$ ) obtained by solid phase extraction (PPL) before (**A**) and after (**B**) chlorination; <sup>13</sup>C DEPT NMR spectra: (top) superimposed protonated carbon NMR resonances (CH<sub>123</sub>; DEPT-45 <sup>13</sup>C NMR spectra), multiplicity-edited <sup>13</sup>C NMR spectra are (second from top) CH; methin, with indices a<sub>1</sub>-d<sub>1</sub> denoting following chemical environments: H<u>C</u><sub>ar</sub>-C / O-H<u>C</u>-O / H<u>C</u>-C), (second from bottom) CH<sub>2</sub>; methylene, with indices a<sub>2</sub> and b<sub>2</sub> denoting following chemical environments: H<sub>2</sub><u>C</u>-O / H<sub>2</sub><u>C</u>-C, and (bottom) CH<sub>3</sub>; methyl, with indices a<sub>3</sub> and b<sub>3</sub> denoting following chemical environments: H<sub>3</sub><u>C</u>-O / H<sub>3</sub><u>C</u>-C. The respective <sup>13</sup>C NMR section integrals are provided in Tab. 4. (**C**) difference <sup>13</sup>C NMR spectra as derived from (top) single pulse <sup>13</sup>C NMR spectra, (middle) DEPT-45 <sup>13</sup>C NMR spectra, and (bottom) DEPT-90 <sup>13</sup>C NMR spectra.