

## Electronic supplementary information

# Molecular level seasonality of dissolved organic matter in freshwater and its impact on drinking water treatment

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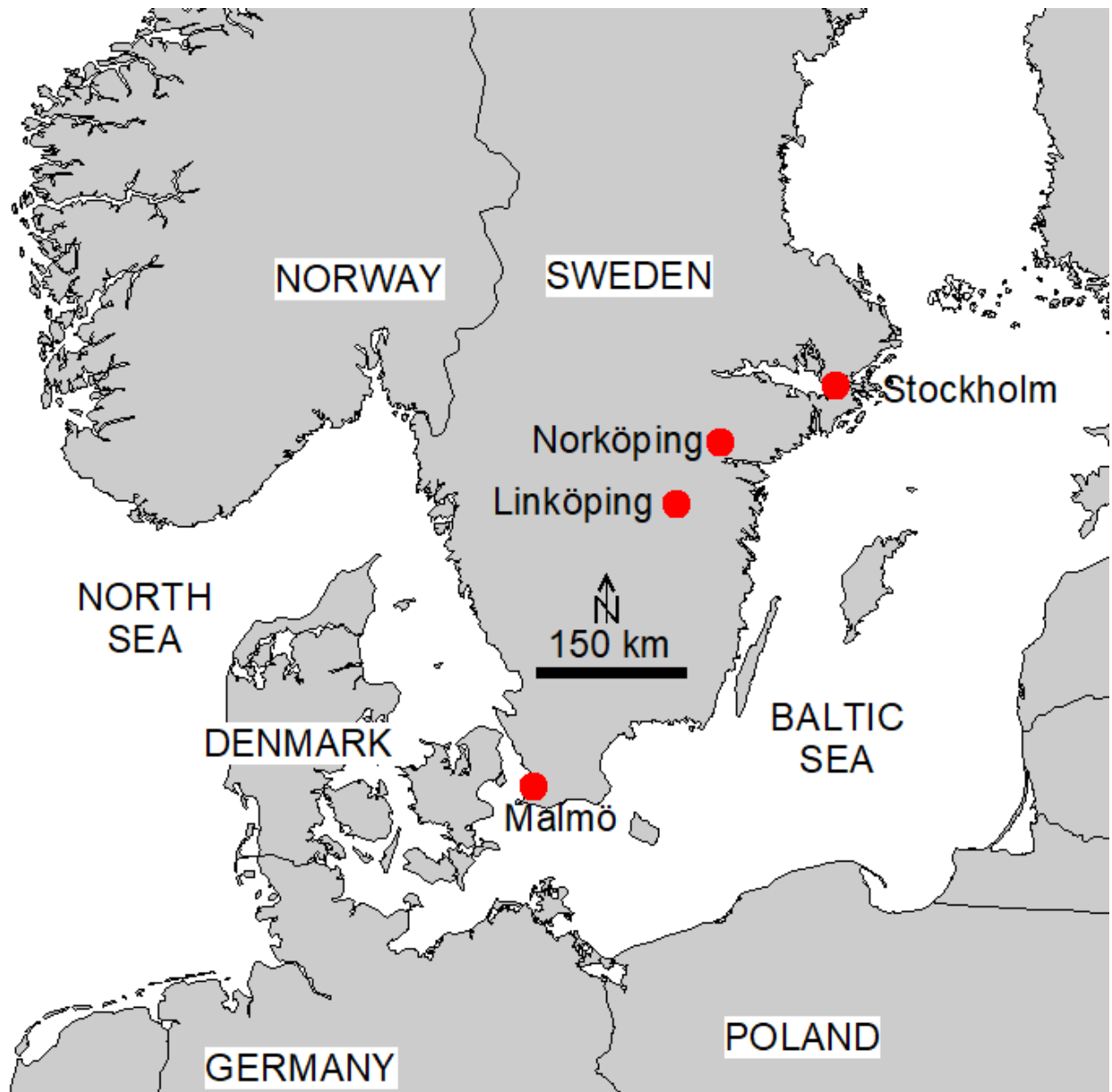
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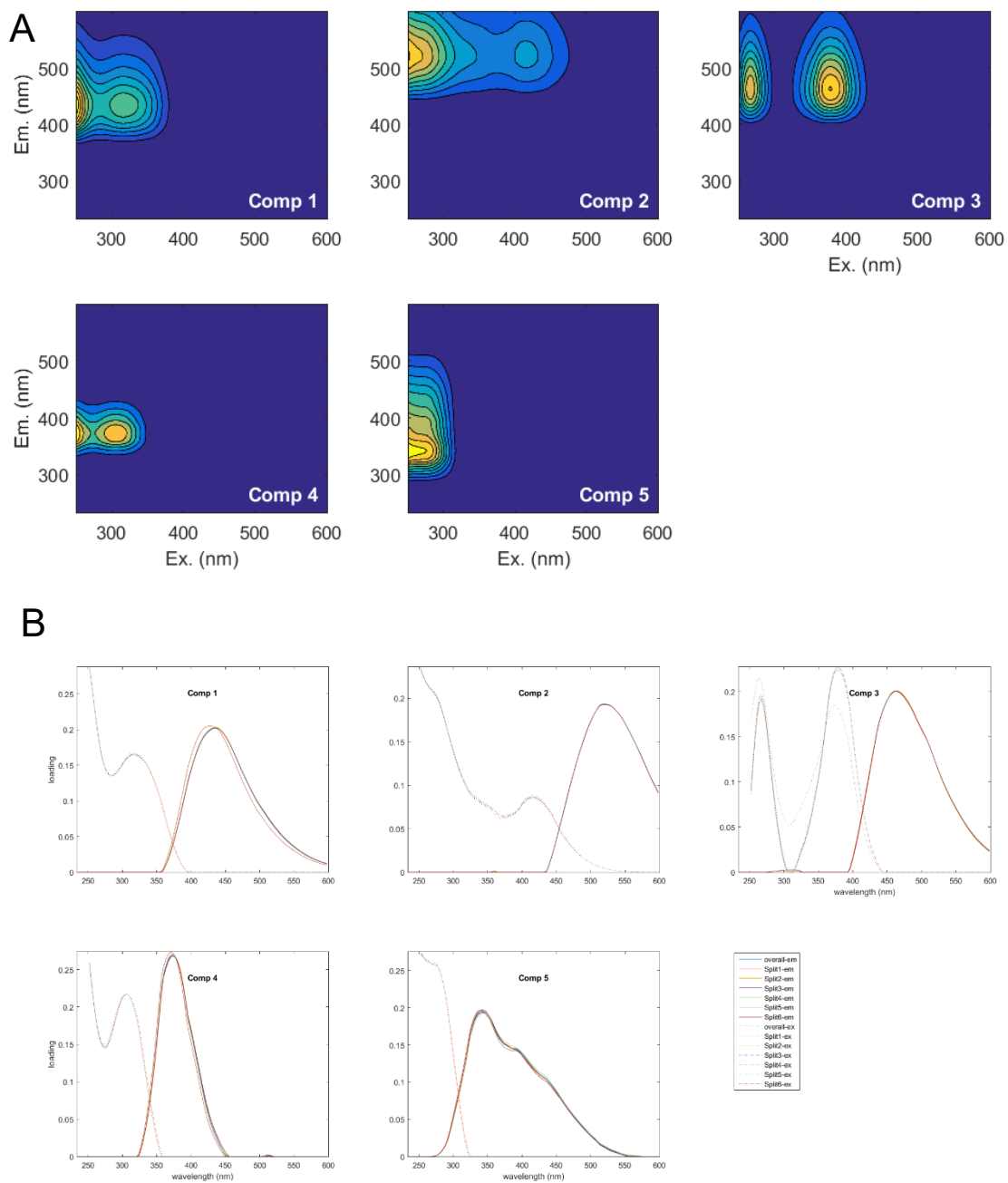


**Figure S1.** Map showing the locations of the cities Linköping (LIN), Norköping (NOR), Stockholm (STO) and Malmö (MAL), where samples were collected at DWTPs for the study.

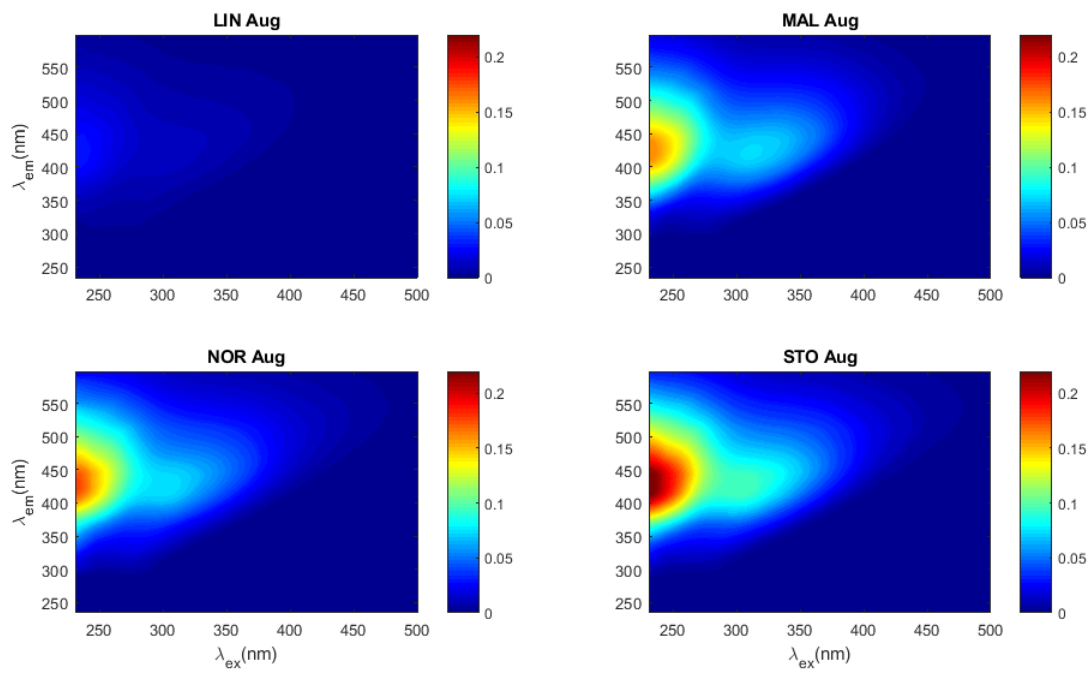
## **EEM PARAFAC analysis**

PARAFAC modelling of the about 300 EEM spectra resulted in a model explained by five components (Figure S2), which were matched with model components of previous studies (Table 2). In general, fluorescence peaks that emits at high wavelengths, contain many conjugated molecules and components 1, 2, 3 and 4 (C1, C2, C3, C4) have been described as humic-like components and component 5 (C5) as a protein-like component (Fellman et al., 2010). C1 and C2 were the EEM profiles that had the highest count of matches with prior model components (Table 2). C1 has been attributed as terrestrial humic-like material (Fellman et al., 2010, Coble, 1996, Guéguen et al., 2014), and this fluorescent signal has been observed for waters influenced by agriculture (Osburn et al., 2012). C2 has been described as a broader group of both humic-like and fulvic-like components (fulvic acids are typically smaller than humic acids and have a higher oxygen content) (Osburn et al., 2012). Due to the high emission maxima of C3 ( $>440$  nm, Table 2), this component have been associated with a humic-like component of terrestrial origin (Yang et al., 2015). The fluorescent signal of C4 has shorter emission wavelengths (Table 2), indicating less aromatic character, compared to C1-C3, and has been associated with low molecular weight molecules from other types of sources, e.g., from biological activity (e.g., plankton), in marine environments, and *in situ* production, e.g., primary production of DOM by algae (Fellman et al., 2010, Guéguen et al., 2014, Kellerman et al., 2015).

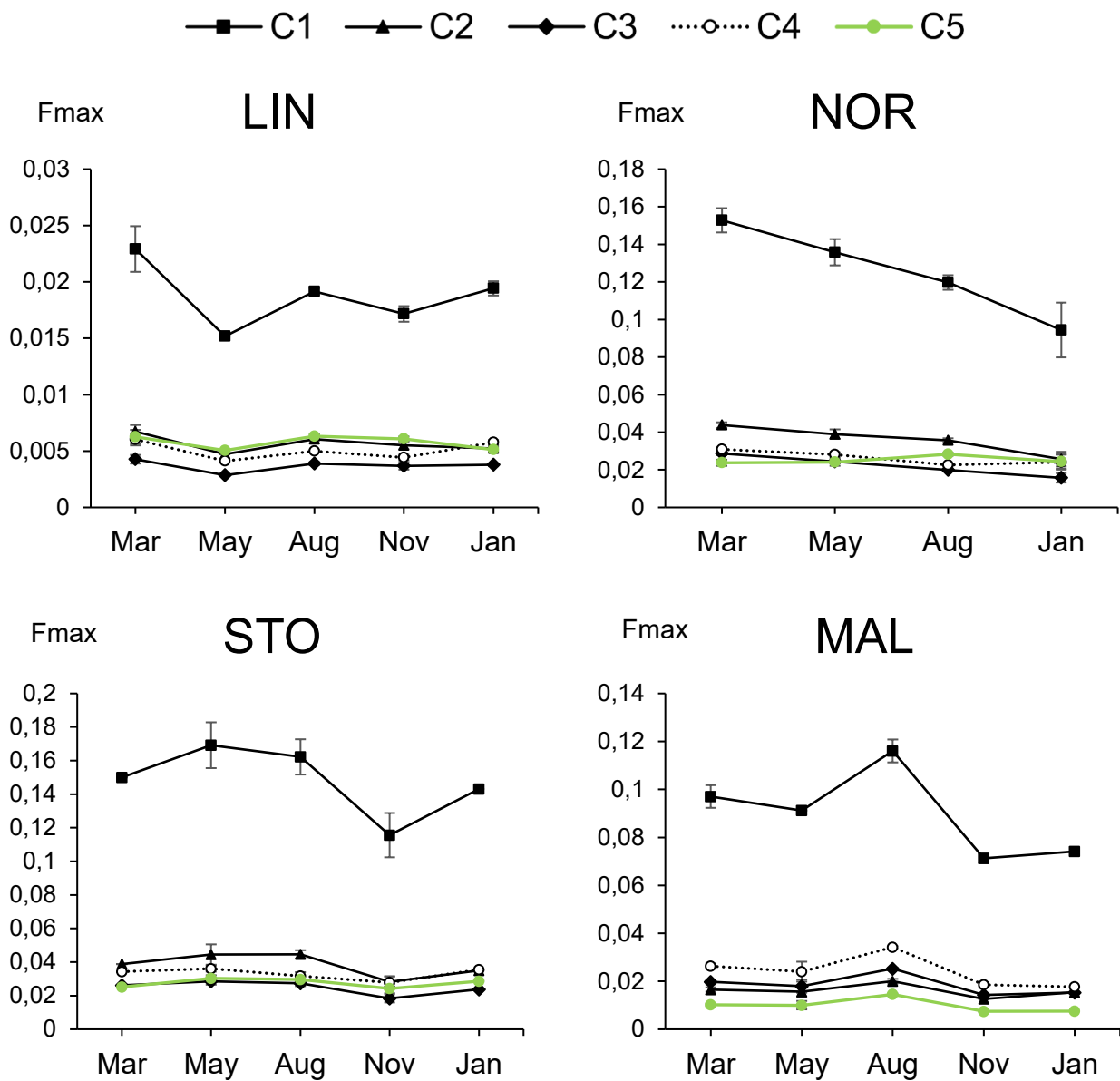
The fluorescent signal of C5 has been associated with protein-like or tryptophan-like compounds (Coble, 1996, Fellman et al., 2010, Osburn et al., 2012). Tryptophan is an amino acid that carries intrinsic fluorescent properties due to its fused aromatic ring side group; fluorescence from proteins mainly arises from the excitation and emission of the tryptophan side group. The origin of protein-like fluorescence has been linked to microbial activity, e.g., when certain microorganisms release enzymes to break down dead plant material (Wong and Williams, 2010). Protein-like fluorescence has also been found to increase by influences of waste water effluents or during algal blooms (Yang et al., 2015). Through combined fluorescence measurements and liquid chromatography with organic carbon detection (LC-OCD), protein-like fluorescence has been linked to a DOM fraction categorized as the biopolymer fraction (Baghoth et al., 2011).



**Figure S2.** EEM spectra of the five components (comp) identified using Parallel Factor Analysis (PARAFAC) (A) and excitation/emission spectra of each component during split-validation (B).

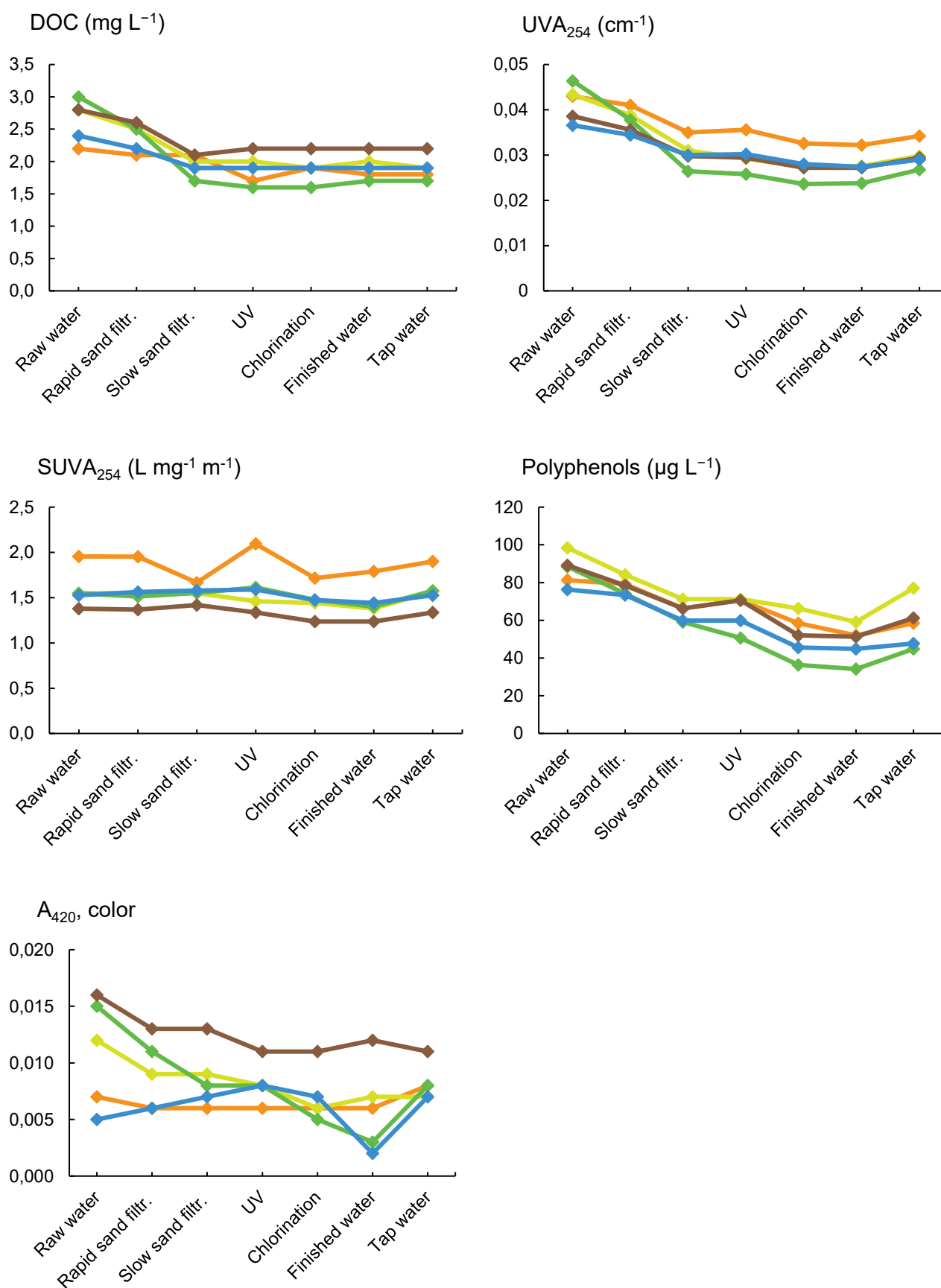


**Figure S3.** Variations in raw water EEM fluorescence spectra compared at the highest fluorescence scale (in Raman units) shown for samples from the four DWTPs collected in August.



**Figure S4.** Variability in raw water characteristics defined by the five PARAFAC components, C1-C5 (Fig S1). Average values of two replicates and standard deviations are presented. Data for NOR in November is missing. Note the different y-axis scales.

LIN      Mar ■      May ■      Aug ■      Nov ■      Jan ■



**Figure S5.** Changes in DOC, UVA<sub>254</sub>, SUVA<sub>254</sub>, polyphenols and A<sub>420</sub> upon treatment at LIN.

NOR

Mar ■ May ■ Aug ■ Nov ■ Jan ■

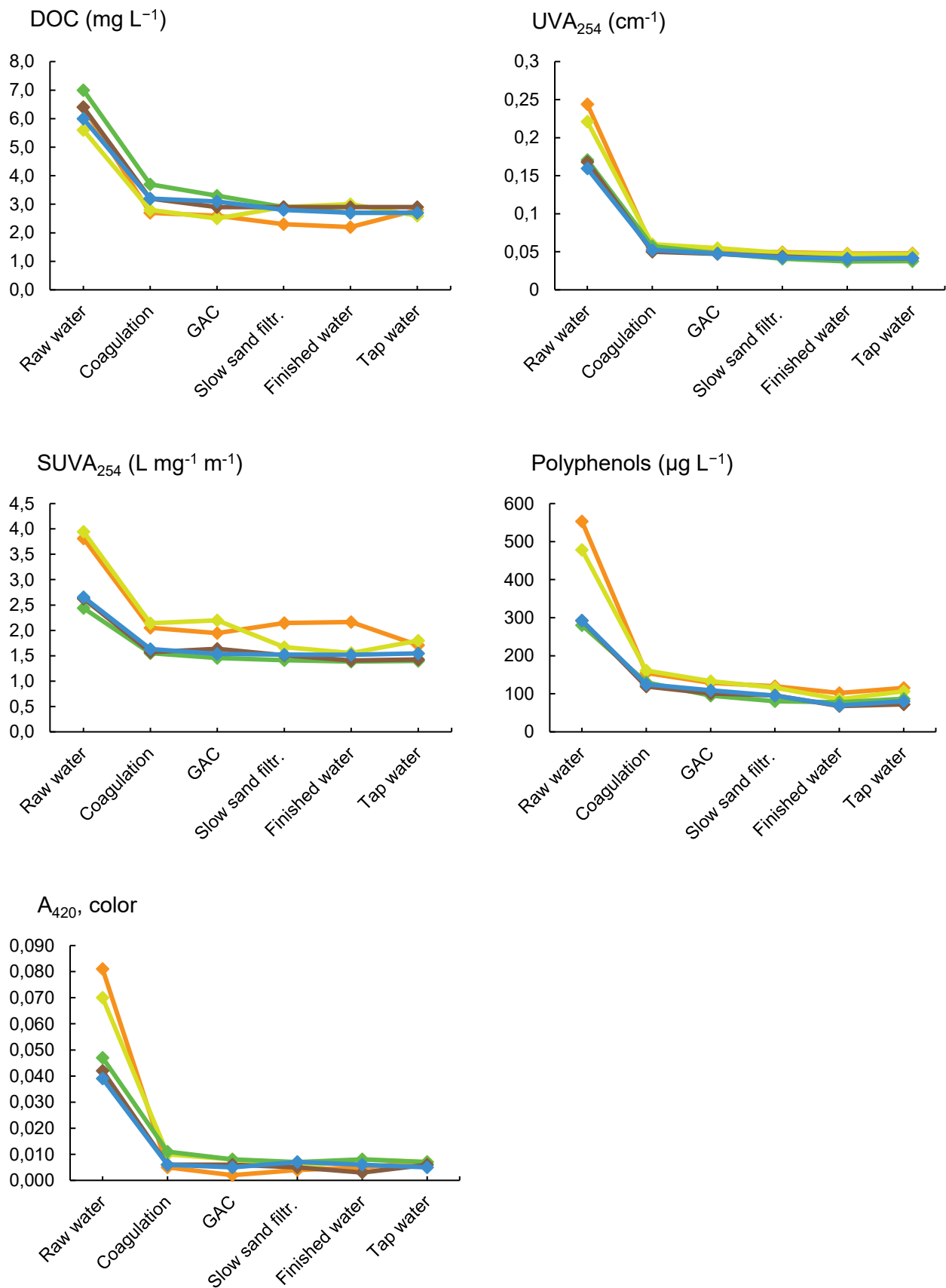
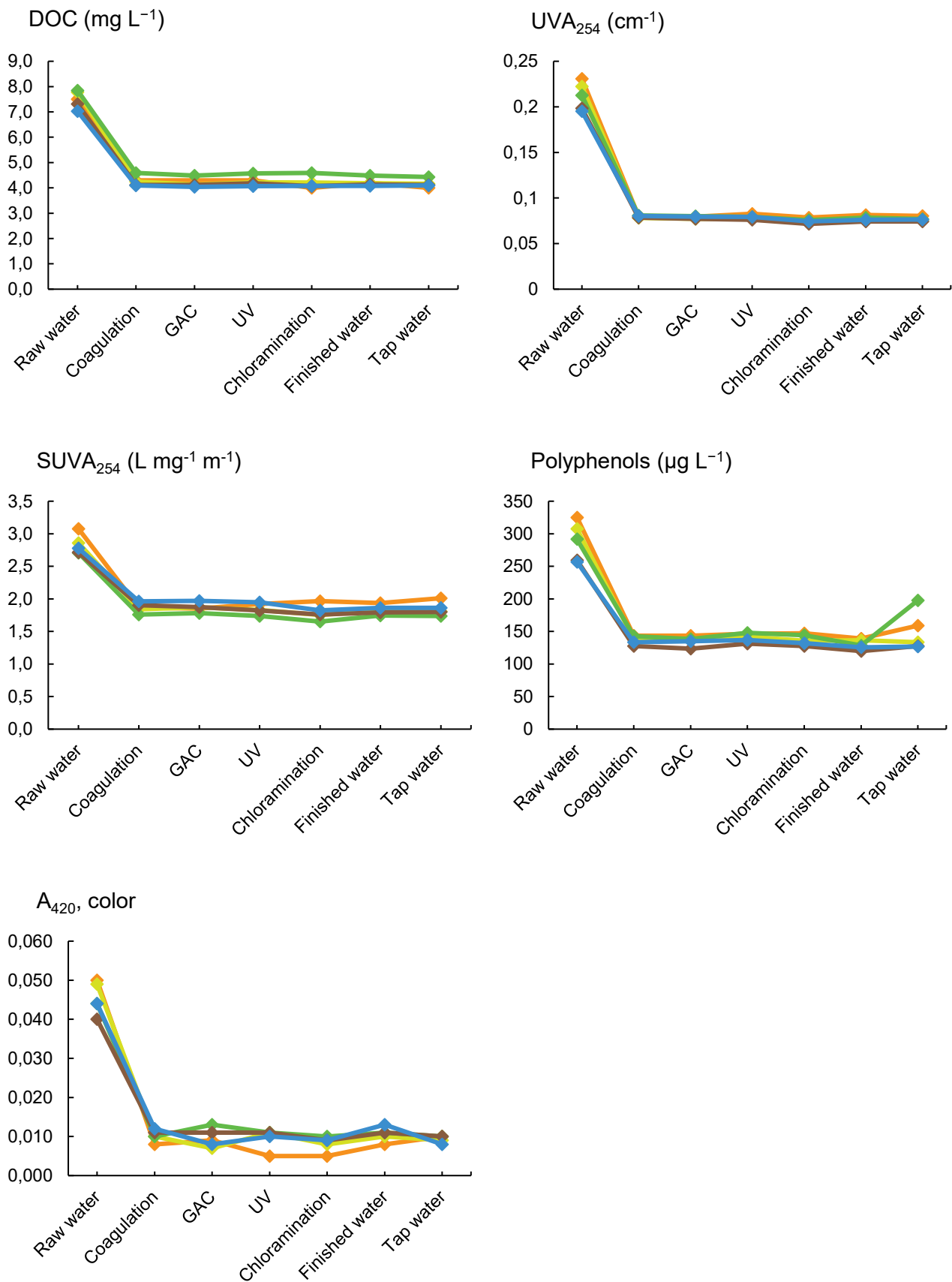


Figure S6. Changes in DOC, UVA<sub>254</sub>, SUVA<sub>254</sub>, polyphenols and A<sub>420</sub> upon treatment at NOR.



# STO

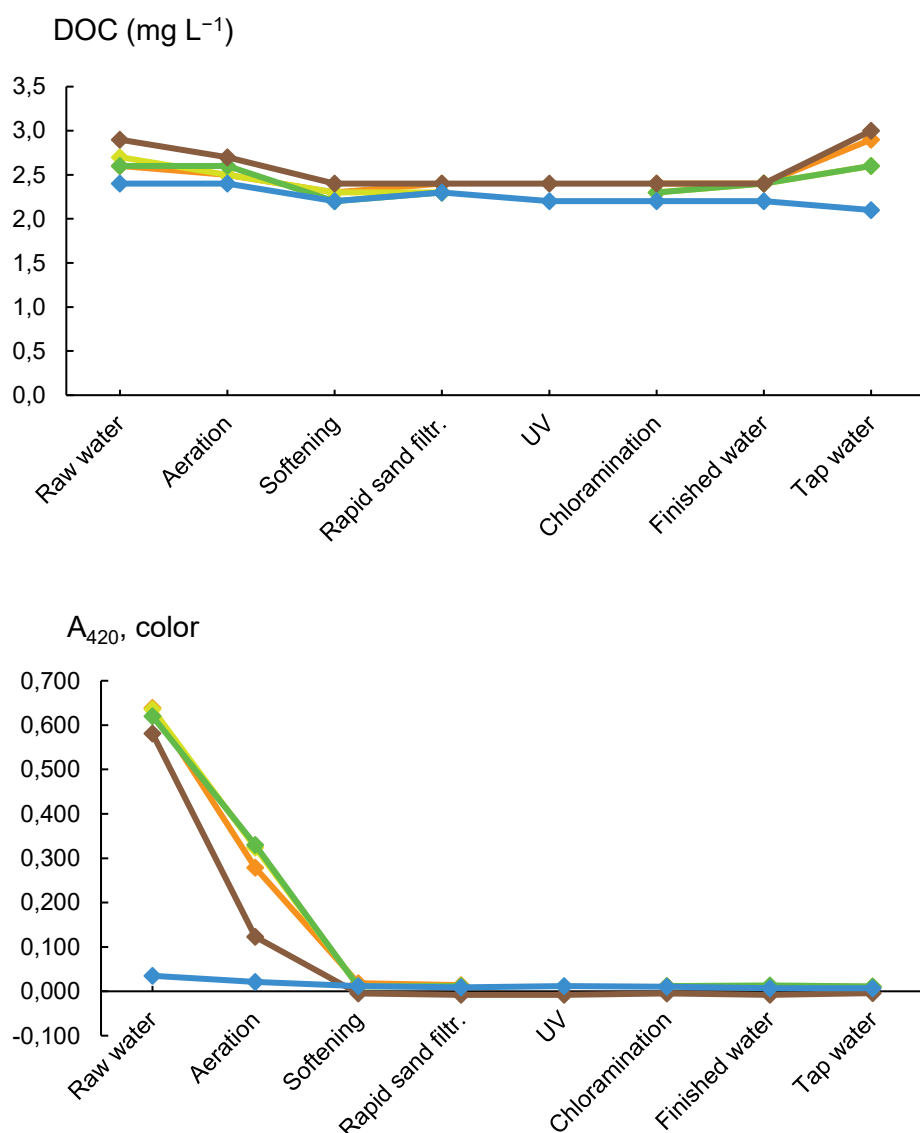
Mar ■ May ■ Aug ■ Nov ■ Jan ■



**Figure S7.** Changes in DOC, UVA<sub>254</sub>, SUVA<sub>254</sub>, polyphenols and A<sub>420</sub> upon treatment at STO.

MAL

Mar ■ May ■ Aug ■ Nov ■ Jan ■

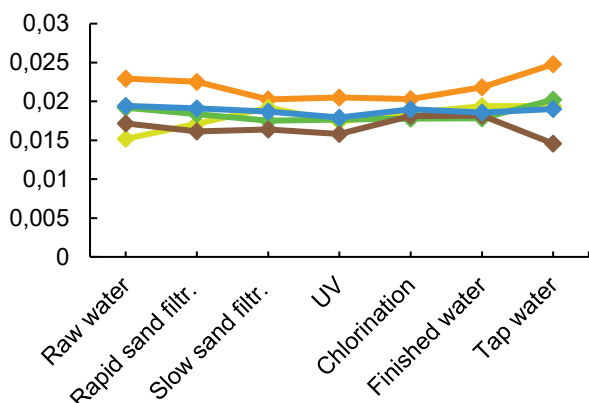


**Figure S8.** Changes in DOC and A<sub>420</sub> upon treatment at MAL. Data for UVA<sub>254</sub>, SUVA<sub>254</sub>, and polyphenols are not shown due to interferences of iron on absorbance measurements for DOM analysis at this water source. UV was installed during the sample campaign, why data at this sampling point only are available for two occasions.

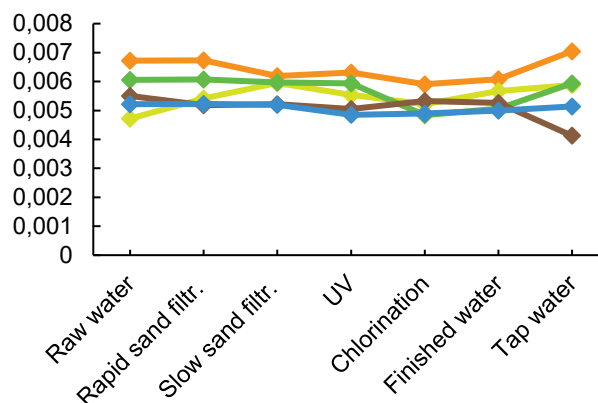
LIN

Mar ■ May ■ Aug ■ Nov ■ Jan ■

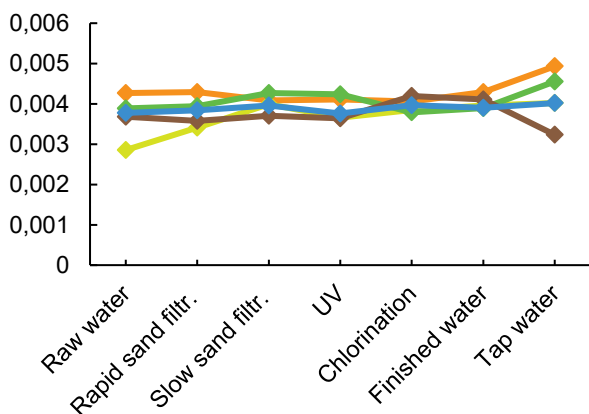
Fmax, C1



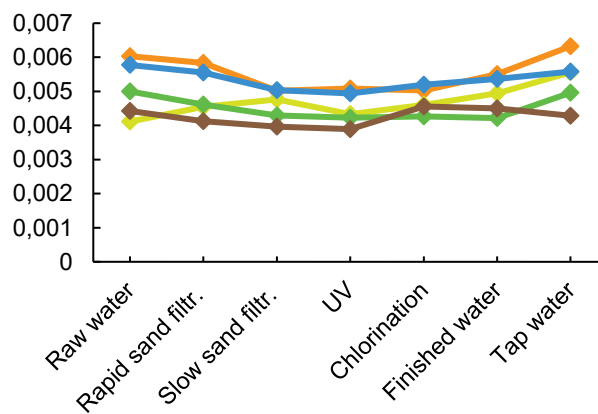
Fmax, C2



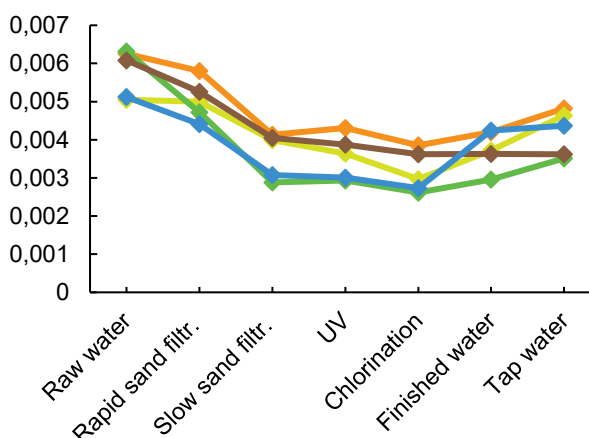
Fmax, C3



Fmax, C4



Fmax, C5

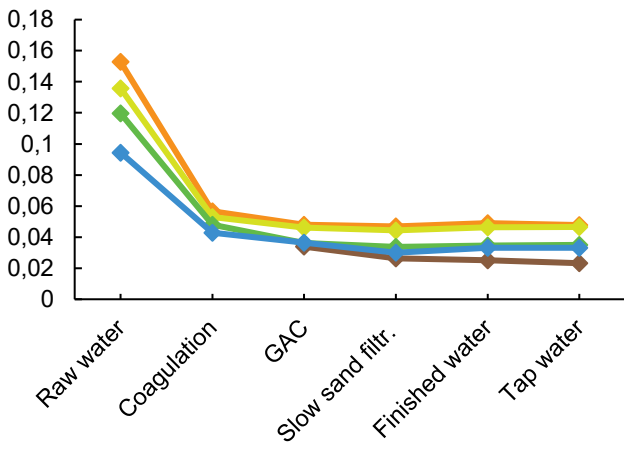


**Figure S9.** Average Fmax values of the five PARAFAC components and the effect of treatment in LIN. The average of two replicates are shown, standard deviations are reported in Table S5.

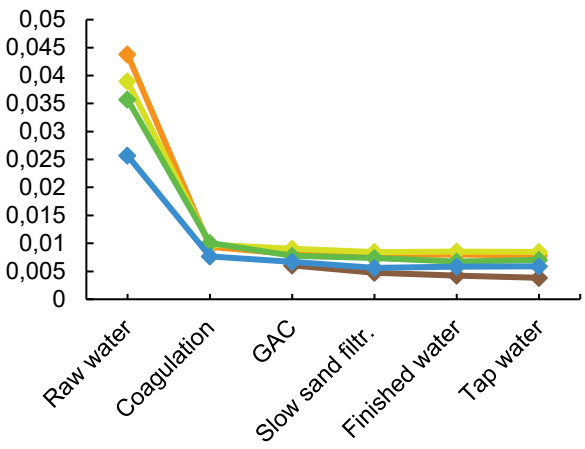
# NOR

Mar ■ May ■ Aug ■ Nov ■ Jan ■

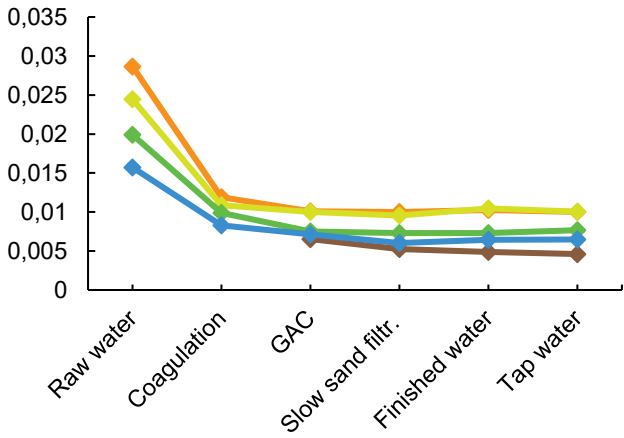
Fmax, C1



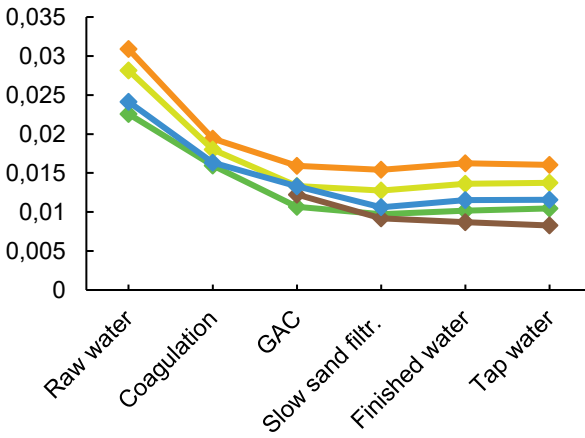
Fmax, C2



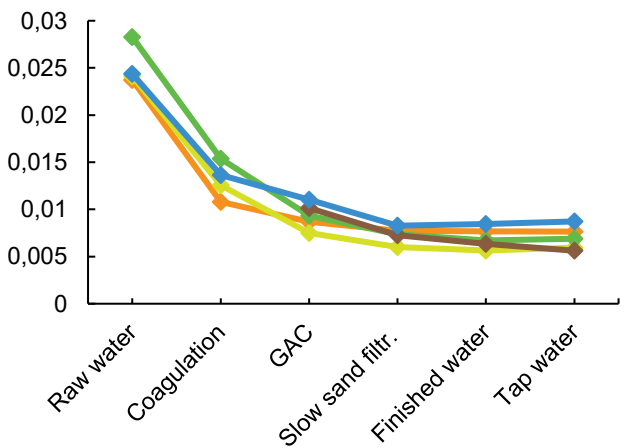
Fmax, C3



Fmax, C4



Fmax, C5

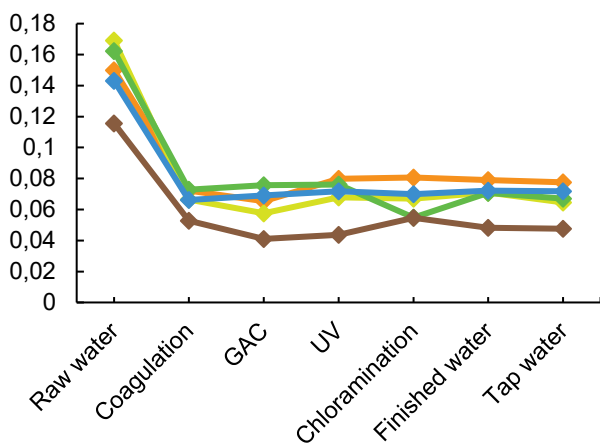


**Figure S10.** Average Fmax values of the five PARAFAC components and the effect of treatment in NOR. The average of two replicates are shown, standard deviations are reported in Table S5. Data for Raw water and coagulation are missing for November.

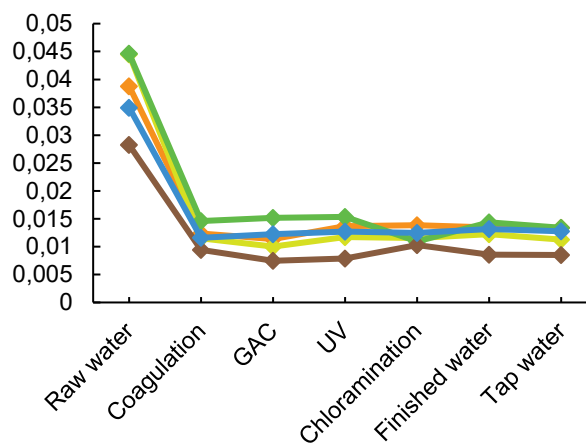
# STO

Mar ■ May ■ Aug ■ Nov ■ Jan ■

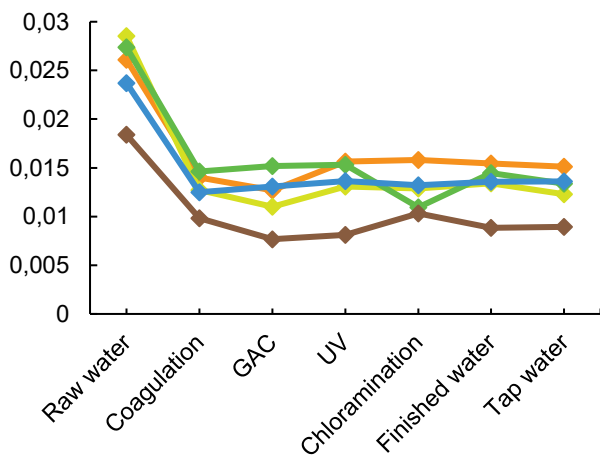
Fmax, C1



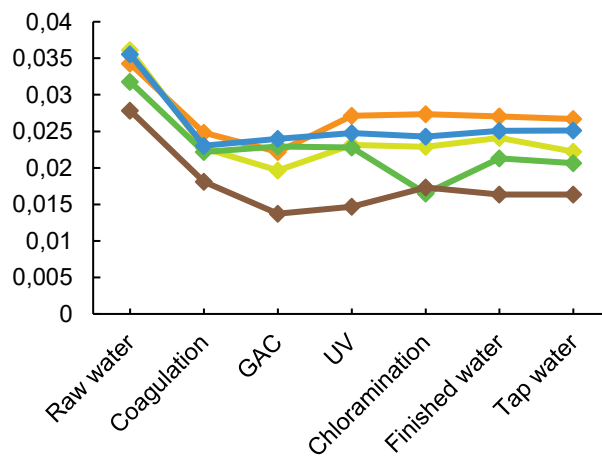
Fmax, C2



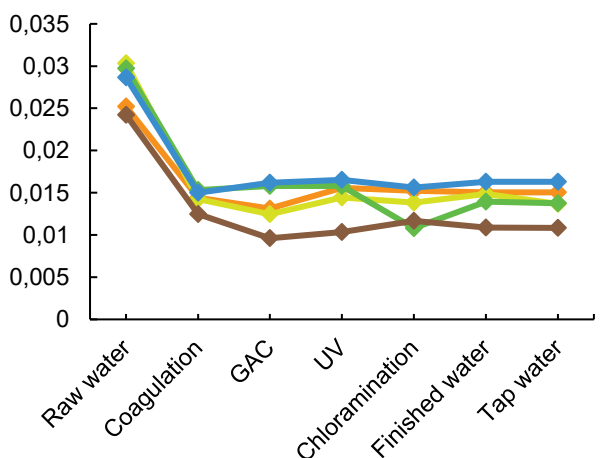
Fmax, C3



Fmax, C4



Fmax, C5

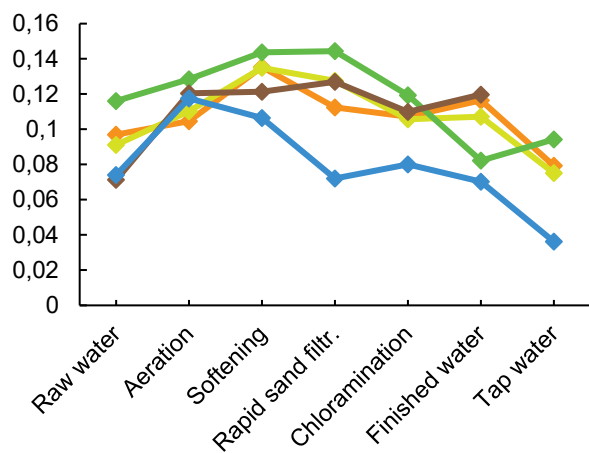


**Figure S11.** Average Fmax values of the five PARAFAC components and the effect of treatment in STO. The average of two replicates are shown, standard deviations are reported in Table S5.

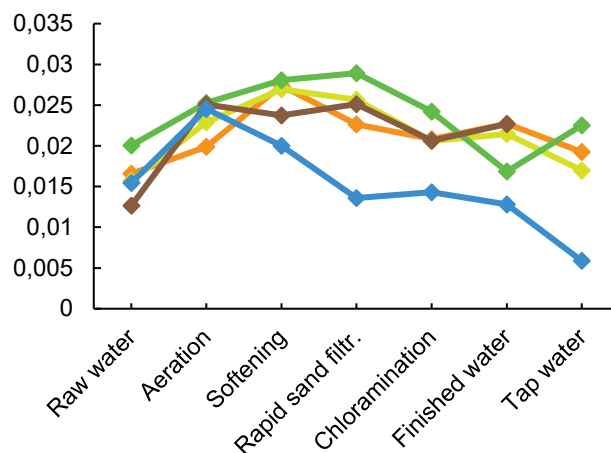
MAL

Mar ■ May ■ Aug ■ Nov ■ Jan ■

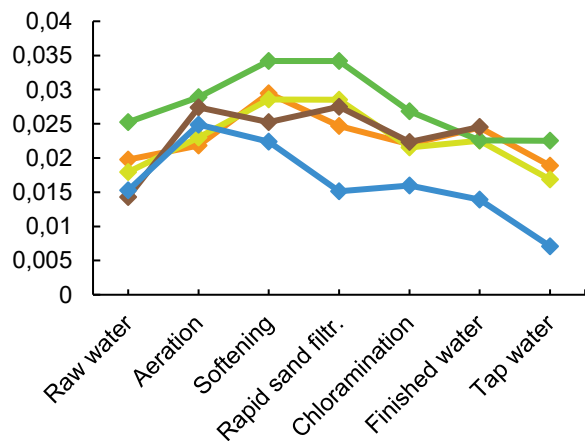
Fmax, C1



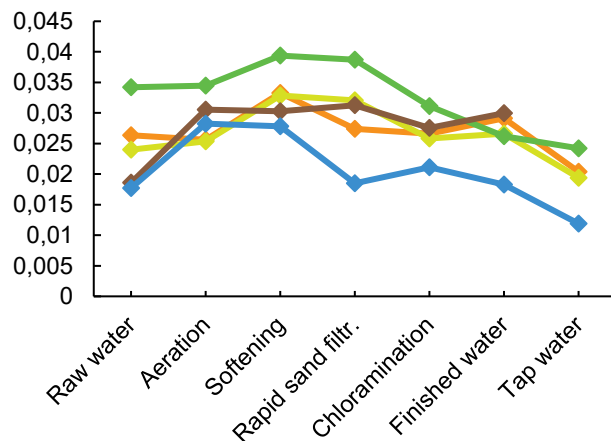
Fmax, C2



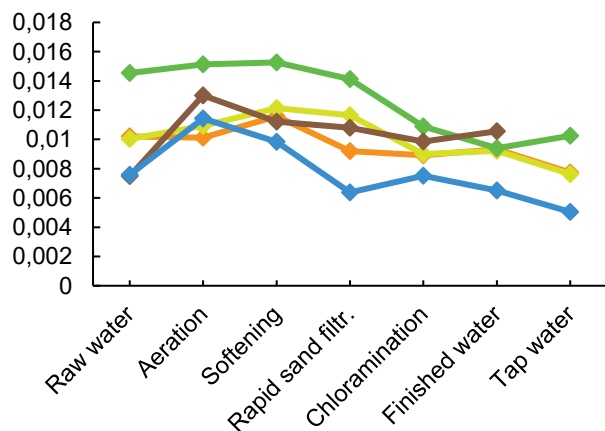
Fmax, C3



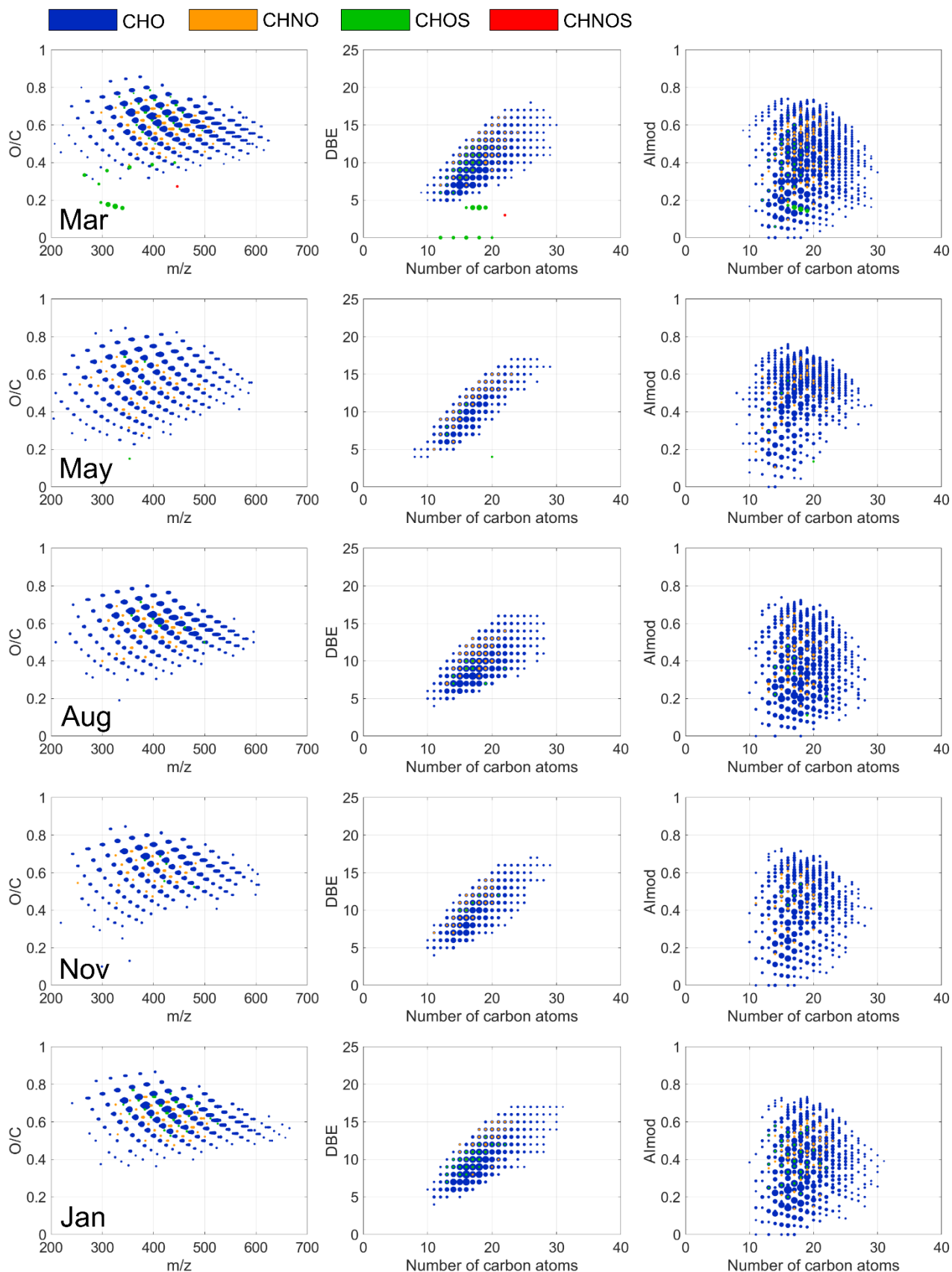
Fmax, C4



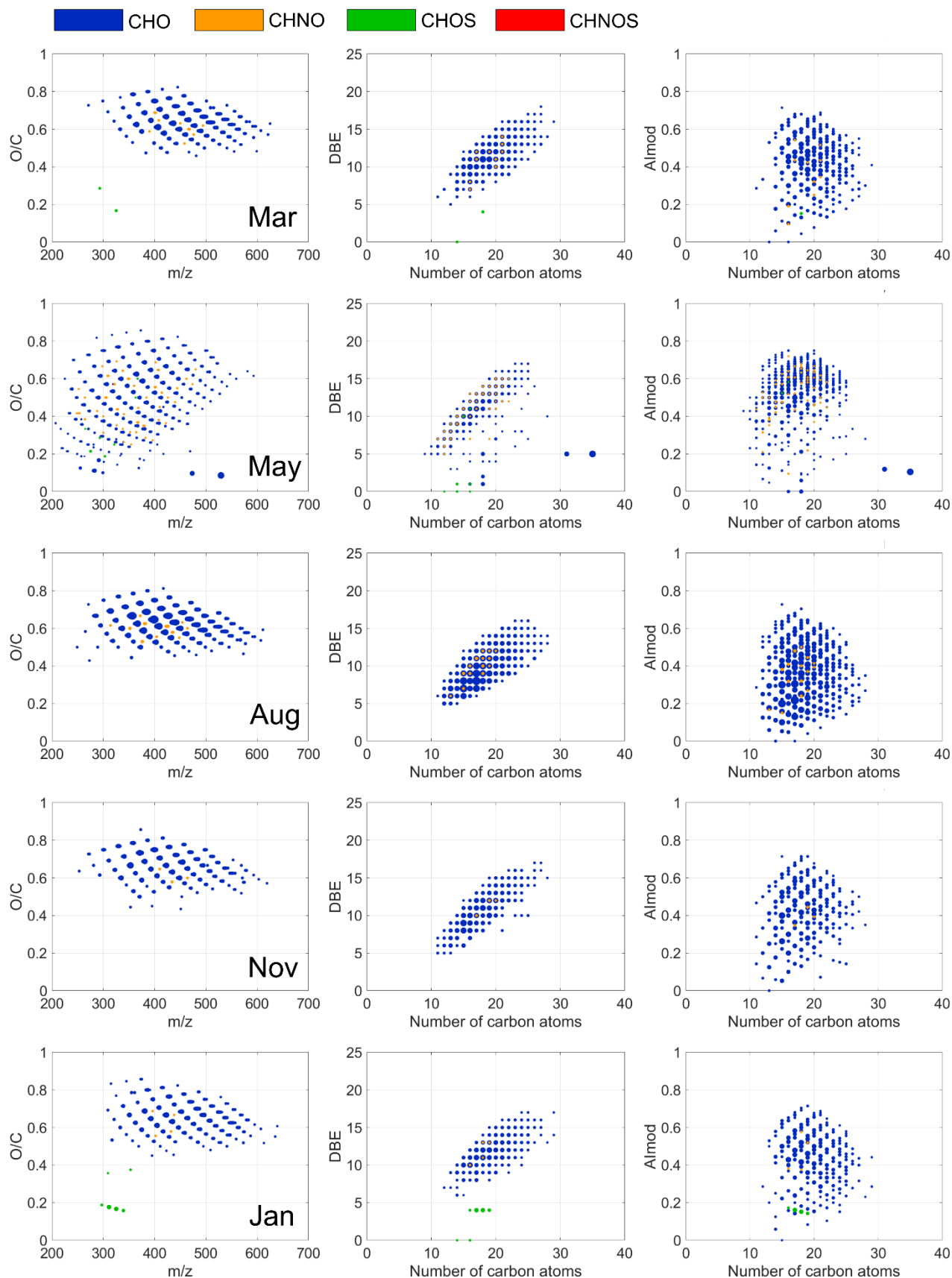
Fmax, C5



**Figure S12.** Average Fmax values of the five PARAFAC components and the effect of treatment in MAL. The average of two replicates are shown, standard deviations are reported in Table S5.

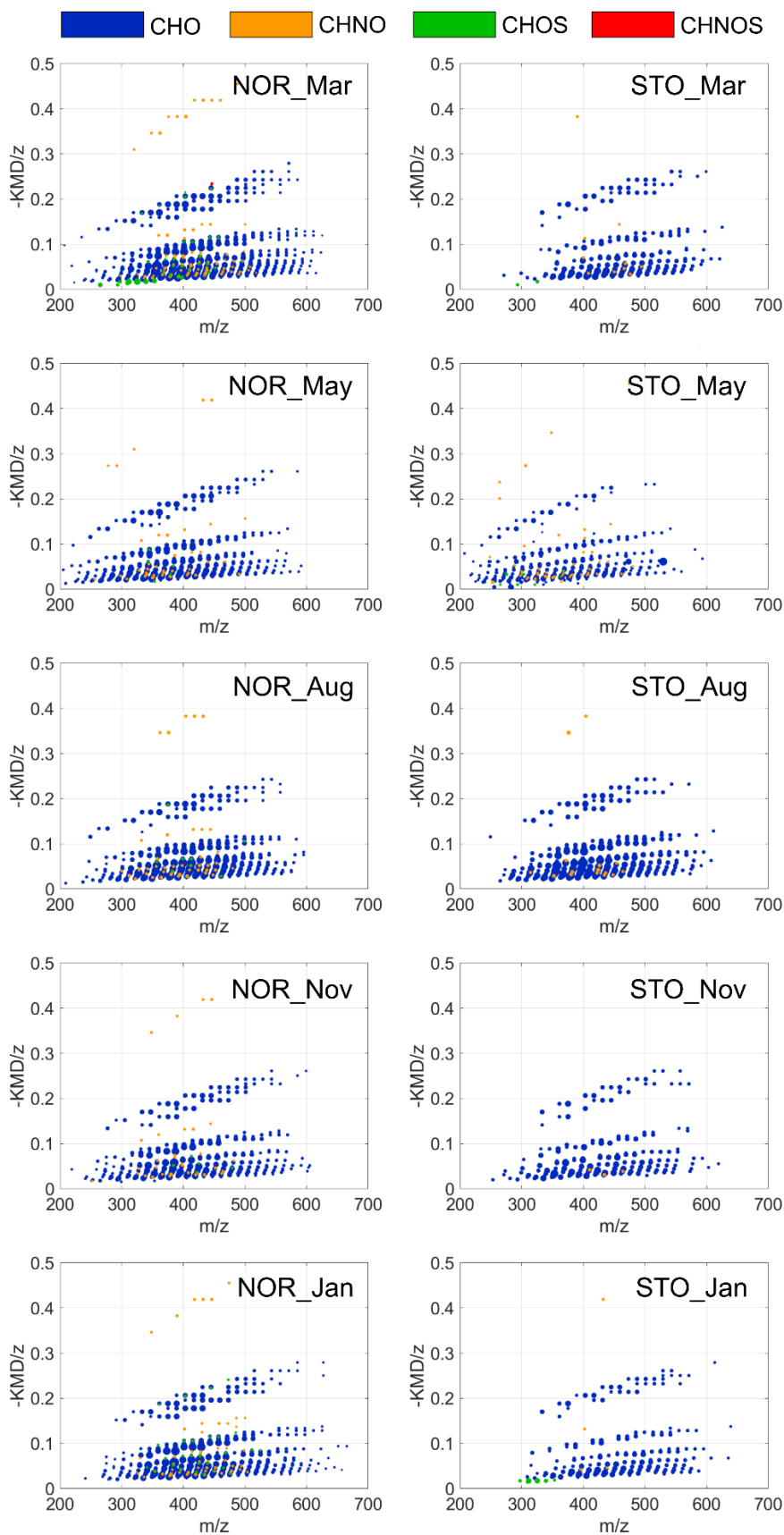


**Figure S13.** Molecular compositions that decreased in relative abundance by more than 50% after coagulation at NOR, presented with mass-edited O/C ratios (left panel), double bond equivalents (DBE) (middle panel) and a modified aromaticity index ( $Al_{mod}$ ) (right panel). DOM changes for five months shown separately. Point size reflect relative mass peak intensity in a sample.

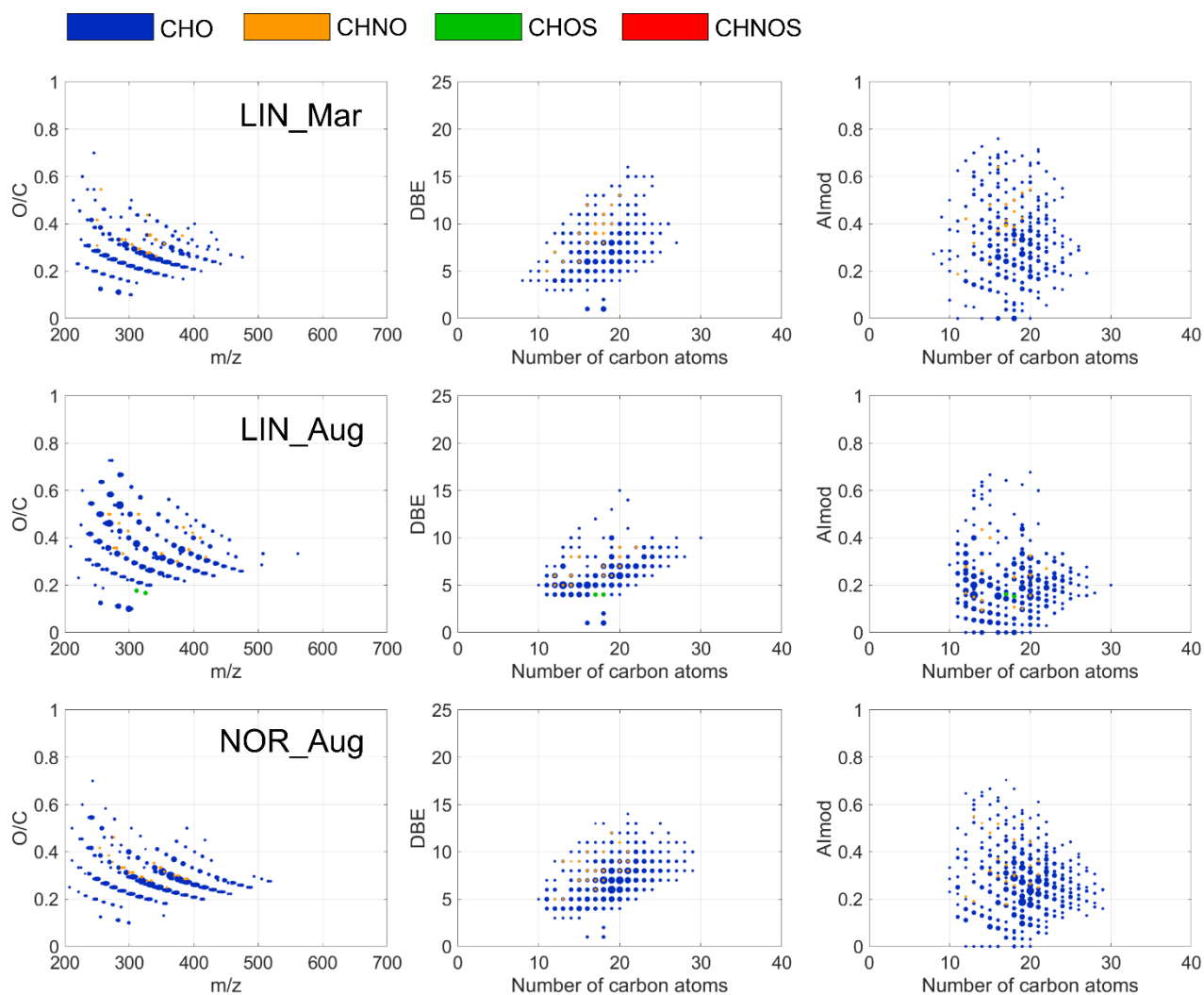


**Figure S14.** Molecular compositions that decreased in relative abundance by more than 50% after coagulation at STO, presented with mass-edited O/C ratios (left panel), double bond equivalents (DBE) (middle panel) and a modified aromaticity index ( $AI_{mod}$ ) (right panel). DOM changes for five months shown separately. Point size reflect relative mass peak intensity in a sample.

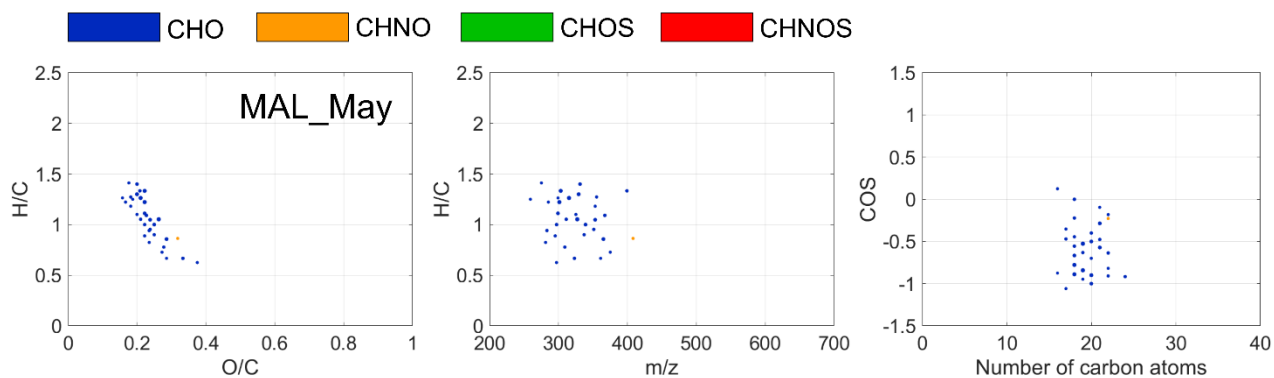




**Figure S15.** Molecular compositions that decreased in relative abundance by more than 50% after coagulation, presented with modified Kendrick mass plots for NOR (left panel) and STO (right panel). DOM changes for five months shown separately. Point size reflect relative mass peak intensity in a sample.



**Figure S16.** Molecular compositions that decreased in relative abundance by more than 50% after slow sand filtration at LIN (March and August) and NOR (August), presented with mass-edited O/C ratios (left panel), double bond equivalents (DBE) (middle panel) and a modified aromaticity index (AI<sub>mod</sub>) (right panel). Point size reflect relative mass peak intensity in a sample.



**Figure S17.** Van Krevelen diagrams (left panel), mass edited H/C ratios (middle panel) and average oxidation state of carbon plots (right panel) of the molecular compositions that decreased in relative abundance by more than 50% after softening in MAL in May. Point size reflect relative mass peak intensity in a sample.

**Table S1.** Counts of  $m/z$  ions and computed indices of molecular formulae detected in the source water of LIN throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). RSD = relative standard deviation. DBE = Double bond equivalences,  $AI_{\text{mod}}$  = a modified aromaticity index.

Characterization of LIN DOM	Mar	May	Aug	Nov	Jan	RSD
number of DOM compositions	2173	2243	2076	2080	1826	8%
average H [%]	44.9	44.6	45.2	44.8	46.0	1%
average C [%]	37.1	36.9	36.7	36.7	36.5	1%
average O [%]	17.8	18.3	17.9	18.3	17.3	2%
average N [%]	0.15	0.14	0.15	0.15	0.13	6%
average S [%]	0.04	0.05	0.04	0.05	0.04	12%
computed average H/C ratio	1.21	1.21	1.23	1.22	1.26	2%
computed average O/C ratio	0.48	0.50	0.49	0.50	0.47	3%
computed average N/C ratio	0.004	0.004	0.004	0.004	0.004	0%
computed average S/C ratio	0.001	0.001	0.001	0.001	0.001	0%
average carbon oxidation state (Cos)	-0.238	-0.204	-0.241	-0.211	-0.295	15%
average DBE	8.2	8.3	8.0	8.1	7.9	2%
average DBE/C	0.45	0.45	0.44	0.45	0.43	2%
average $AI_{\text{mod}}$	0.28	0.27	0.26	0.26	0.24	6%
mass weighted average [Da]	382.3	389.6	386.2	383.9	390.4	1%

**Table S2.** Counts of  $m/z$  ions and computed indices of molecular formulae detected in the source water of NOR throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). RSD = relative standard deviation. DBE = Double bond equivalences,  $AI_{\text{mod}}$  = a modified aromaticity index.

Characterization of NOR DOM	Mar	May	Aug	Nov	Jan	RSD
number of DOM compositions	2846	2388	2187	2362	2859	12%
average H [%]	42.9	43.8	44.4	43.5	42.7	2%
average C [%]	37.8	38.1	37.7	37.4	37.4	1%
average O [%]	19.0	18.0	17.7	18.9	19.6	4%
average N [%]	0.14	0.14	0.13	0.14	0.14	3%
average S [%]	0.09	0.05	0.05	0.07	0.11	35%
computed average H/C ratio	1.14	1.15	1.18	1.16	1.14	1%
computed average O/C ratio	0.50	0.47	0.47	0.50	0.52	4%
computed average N/C ratio	0.004	0.004	0.004	0.004	0.004	0%
computed average S/C ratio	0.002	0.001	0.001	0.002	0.003	46%
average carbon oxidation state (Cos)	-0.117	-0.189	-0.222	-0.139	-0.083	37%
average DBE	9.1	8.9	8.7	8.8	9.1	2%
average DBE/C	0.49	0.48	0.47	0.47	0.48	2%
average $AI_{\text{mod}}$	0.31	0.32	0.30	0.29	0.29	4%
mass weighted average [Da]	399.8	385.9	388.3	396.4	407.2	2%

**Table S3.** Counts of  $m/z$  ions and computed indices of molecular formulae detected in the source water of STO throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). RSD = relative standard deviation. DBE = Double bond equivalences,  $AI_{\text{mod}}$  = a modified aromaticity index.

Characterization of STO DOM	Mar	May	Aug	Nov	Jan	RSD
number of DOM compositions	2185	2719	1688	1589	2198	22%
average H [%]	42.9	44.7	43.5	43.1	43.0	2%
average C [%]	37.4	37.7	37.3	37.3	37.3	0%
average O [%]	19.5	17.4	19.0	19.4	19.5	5%
average N [%]	0.11	0.15	0.10	0.09	0.10	21%
average S [%]	0.08	0.05	0.05	0.05	0.09	30%
computed average H/C ratio	1.15	1.18	1.16	1.16	1.15	1%
computed average O/C ratio	0.52	0.46	0.51	0.52	0.52	5%
computed average N/C ratio	0.003	0.004	0.003	0.003	0.003	14%
computed average S/C ratio	0.002	0.001	0.001	0.001	0.003	56%
average carbon oxidation state (Cos)	-0.095	-0.245	-0.133	-0.106	-0.098	47%
average DBE	9.3	8.8	9.0	8.9	9.1	2%
average DBE/C	0.48	0.46	0.47	0.48	0.48	2%
average $AI_{\text{mod}}$	0.29	0.30	0.29	0.29	0.29	2%
mass weighted average [Da]	417.3	392.5	407.8	404.2	414.1	2%

**Table S4.** Counts of  $m/z$  ions and computed indices of molecular formulae detected in the source water of MAL throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). RSD = relative standard deviation. DBE = Double bond equivalences,  $AI_{\text{mod}}$  = a modified aromaticity index.

Characterization of MAL DOM	Mar	May	Aug	Nov	Jan	RSD
number of DOM compositions	2062	2106	2108	1927	2318	7%
average H [%]	44.4	44.7	44.7	44.3	44.0	1%
average C [%]	37.7	37.8	37.9	37.7	38.0	0%
average O [%]	17.6	17.3	17.2	17.8	17.8	2%
average N [%]	0.14	0.16	0.16	0.14	0.18	11%
average S [%]	0.05	0.05	0.05	0.05	0.06	9%
computed average H/C ratio	1.18	1.18	1.18	1.17	1.16	1%
computed average O/C ratio	0.47	0.46	0.45	0.47	0.47	2%
computed average N/C ratio	0.004	0.004	0.004	0.004	0.005	11%
computed average S/C ratio	0.001	0.001	0.001	0.001	0.002	37%
average carbon oxidation state (Cos)	-0.233	-0.258	-0.259	-0.222	-0.212	9%
average DBE	9.1	8.9	8.9	9.1	9.3	2%
average DBE/C	0.46	0.46	0.47	0.47	0.47	1%
average $AI_{\text{mod}}$	0.30	0.30	0.30	0.30	0.31	1%
mass weighted average [Da]	408.3	396.3	395.2	407.0	406.3	2%

**Table S5.** Standard deviations of component C1 and C2-C5 for LIN, NOR, STO and MAL, acquired from EEM fluorescence measurements based on two replicates. The highest standard deviations for each sampling point (combining the five months of sampling) are reported.

<b>DWTP</b>	<b>Treatment stage</b>	<b>Max std C1</b>	<b>Max std C2-C5</b>
LIN	Raw water	0.002027	0.000619
	Rapid sand filtr.	0.003386	0.001001
	Slow sand filtr.	0.003630	0.005949
	UV	0.005404	0.001505
	Chlorination	0.006792	0.001948
	Finished water	0.003018	0.000867
	Tap water	0.008600	0.002771
NOR	Raw water	0.014572	0.043782
	Coagulation	0.001929	0.019418
	GAC	0.003464	0.015902
	Slow sand filtr.	0.006276	0.015408
	Finished water	0.000910	0.016249
	Tap water	0.002853	0.016026
STO	Raw water	0.013644	0.006051
	Coagulation	0.007772	0.002775
	GAC	0.013221	0.019643
	UV	0.004475	0.001372
	Chloramination	0.020626	0.005732
	Finished water	0.010721	0.003861
	Tap water	0.004174	0.001576
MAL	Raw water	0.004779	0.001105
	Aeration	0.032102	0.009941
	Softening	0.010560	0.002524
	Rapid sand filtr.	0.004191	0.001194
	Chloramination	0.023233	0.006211
	Finished water	0.012459	0.007084
	Tap water	0.009970	0.002664

**Table S6.** Counts of  $m/z$  ions and computed indices of molecular formulae detected at the different stages of water treatment in LIN throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). The sample points (1-7) are: 1 - Raw water, 2 - Rapid sand filtr., 3 - Slow sand filtr., 4 - UV, 5 - Chlorination, 6 - Finished water, 7 - Tap water. DBE = Double bond equivalences,  $AI_{mod}$  = a modified aromaticity index.

Characterization of DOM Mar	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2173	2203	2202	2162	2267	2216	2255
average H [%]	44.9	45.3	44.5	44.5	44.1	44.4	44.3
average C [%]	37.1	37.4	37.1	37.1	37.1	37.2	37.1
average O [%]	17.8	17.1	18.3	18.2	18.6	18.2	18.4
average N [%]	0.15	0.16	0.14	0.15	0.15	0.16	0.15
average S [%]	0.04	0.04	0.05	0.04	0.06	0.05	0.05
computed average H/C ratio	1.21	1.21	1.20	1.20	1.19	1.19	1.20
computed average O/C ratio	0.48	0.46	0.49	0.49	0.50	0.49	0.50
computed average N/C ratio	0.0041	0.0042	0.0039	0.0040	0.0041	0.0042	0.0040
computed average S/C ratio	0.0010	0.0010	0.0013	0.0012	0.0017	0.0012	0.0015
average carbon oxidation state (Cos)	-0.238	-0.279	-0.202	-0.207	-0.173	-0.198	-0.190
average DBE	8.2	8.2	8.4	8.4	8.3	8.3	8.3
average DBE/C	0.45	0.45	0.46	0.46	0.46	0.46	0.46
average $AI_{mod}$	0.27	0.28	0.27	0.27	0.28	0.28	0.27
mass weighted average [Da]	382.3	373.3	388.0	387.9	385.3	380.6	384.3
Characterization of DOM May	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2243	2137	1967	1917	1935	1847	2024
average H [%]	44.6	44.9	45.0	44.5	44.4	44.5	44.7
average C [%]	36.9	37.0	37.1	36.9	36.9	36.8	37.0
average O [%]	18.3	17.9	17.7	18.4	18.5	18.5	18.1
average N [%]	0.14	0.15	0.14	0.14	0.14	0.14	0.15
average S [%]	0.05	0.05	0.04	0.04	0.04	0.04	0.04
computed average H/C ratio	1.21	1.21	1.21	1.21	1.20	1.21	1.21
computed average O/C ratio	0.50	0.48	0.48	0.50	0.50	0.50	0.49
computed average N/C ratio	0.0039	0.0040	0.0039	0.0037	0.0039	0.0037	0.0039
computed average S/C ratio	0.0013	0.0012	0.0011	0.0012	0.0012	0.0012	0.0011
average carbon oxidation state (Cos)	-0.204	-0.230	-0.244	-0.200	-0.190	-0.192	-0.219
average DBE	8.3	8.2	8.3	8.4	8.3	8.3	8.2
average DBE/C	0.45	0.45	0.45	0.45	0.45	0.45	0.45
average $AI_{mod}$	0.27	0.27	0.27	0.27	0.27	0.26	0.27
mass weighted average [Da]	389.6	383.4	385.0	395.7	390.6	393.3	384.1
Characterization of DOM Aug	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2076	1700	1876	1898	1727	1783	1813
average H [%]	45.2	45.9	44.6	44.5	44.4	44.5	45.0
average C [%]	36.7	36.8	36.8	36.8	36.6	36.8	36.9
average O [%]	17.9	17.2	18.4	18.5	18.8	18.6	18.0
average N [%]	0.15	0.14	0.14	0.14	0.15	0.15	0.15
average S [%]	0.04	0.03	0.05	0.05	0.04	0.04	0.04

computed average H/C ratio	1.23	1.25	1.21	1.21	1.21	1.21	1.22
computed average O/C ratio	0.49	0.47	0.50	0.50	0.51	0.50	0.49
computed average N/C ratio	0.0040	0.0038	0.0039	0.0038	0.0040	0.0040	0.0040
computed average S/C ratio	0.0012	0.0009	0.0013	0.0014	0.0011	0.0011	0.0010
average carbon oxidation state (Cos)	-0.241	-0.298	-0.202	-0.192	-0.174	-0.189	-0.230
average DBE	8.0	8.0	8.4	8.4	8.3	8.3	8.2
average DBE/C	0.44	0.43	0.45	0.45	0.45	0.45	0.45
average AI <sub>mod</sub>	0.26	0.26	0.26	0.26	0.26	0.26	0.26
mass weighted average [Da]	386.2	382.5	395.8	396.6	398.0	393.9	388.3
Characterization of DOM Nov	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2080	2033	2030	1818	1731	1900	1893
average H [%]	44.8	44.7	44.5	44.6	44.6	44.4	44.2
average C [%]	36.7	36.7	36.7	36.6	36.6	36.7	36.5
average O [%]	18.3	18.4	18.6	18.6	18.5	18.8	19.1
average N [%]	0.15	0.15	0.14	0.13	0.14	0.14	0.13
average S [%]	0.05	0.05	0.06	0.05	0.04	0.05	0.06
computed average H/C ratio	1.22	1.22	1.21	1.22	1.22	1.21	1.21
computed average O/C ratio	0.50	0.50	0.51	0.51	0.51	0.51	0.52
computed average N/C ratio	0.0040	0.0040	0.0039	0.0036	0.0039	0.0039	0.0037
computed average S/C ratio	0.0014	0.0014	0.0017	0.0015	0.0012	0.0014	0.0018
average carbon oxidation state (Cos)	-0.211	-0.202	-0.188	-0.191	-0.192	-0.172	-0.152
average DBE	8.1	8.1	8.2	8.2	8.1	8.1	8.2
average DBE/C	0.45	0.45	0.45	0.45	0.45	0.45	0.45
average AI <sub>mod</sub>	0.26	0.26	0.26	0.26	0.26	0.26	0.25
mass weighted average [Da]	383.9	385.6	388.4	392.3	388.1	386.4	393.9
Characterization of DOM Jan	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	1826	1954	2179	2221	2146	2127	2112
average H [%]	46.0	45.4	44.6	44.3	44.3	44.4	44.7
average C [%]	36.5	36.6	36.7	36.7	36.6	36.7	36.8
average O [%]	17.3	17.8	18.5	18.8	18.9	18.7	18.3
average N [%]	0.13	0.14	0.15	0.15	0.15	0.15	0.16
average S [%]	0.04	0.04	0.06	0.06	0.06	0.05	0.04
computed average H/C ratio	1.26	1.24	1.22	1.21	1.21	1.21	1.22
computed average O/C ratio	0.47	0.49	0.51	0.51	0.52	0.51	0.50
computed average N/C ratio	0.0036	0.0039	0.0040	0.0040	0.0040	0.0042	0.0043
computed average S/C ratio	0.0010	0.0011	0.0016	0.0016	0.0017	0.0013	0.0012
average carbon oxidation state (Cos)	-0.295	-0.256	-0.193	-0.173	-0.164	-0.178	-0.202
average DBE	7.9	8.0	8.2	8.3	8.2	8.2	8.1
average DBE/C	0.43	0.44	0.45	0.45	0.45	0.45	0.45
average AI <sub>mod</sub>	0.24	0.25	0.26	0.26	0.26	0.26	0.26
mass weighted average [Da]	390.4	388.6	393.5	393.0	391.8	389.9	384.5



**Table S7.** Counts of  $m/z$  ions and computed indices of molecular formulae detected at the different stages of water treatment in NOR throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). The sample points (1-6) are: 1 - Raw water. 2 - Coagulation.. 3 - GAC. 4 - Slow sand filtr.. 5 - Finished water. 6 - Tap water. DBE = Double bond equivalences.  $AI_{mod}$  = a modified aromaticity index.

Characterization of DOM Mar	P1	P2	P3	P4	P5	P6
number of DOM compositions	2846	2817	2347	2447	2497	2515
average H [%]	42.9	45.5	46.3	45.9	45.8	45.4
average C [%]	37.8	37.3	37.3	37.2	37.2	37.0
average O [%]	19.0	16.9	16.2	16.7	16.8	17.4
average N [%]	0.14	0.15	0.13	0.14	0.14	0.14
average S [%]	0.09	0.10	0.07	0.08	0.07	0.09
computed average H/C ratio	1.14	1.22	1.24	1.23	1.23	1.22
computed average O/C ratio	0.50	0.45	0.43	0.45	0.45	0.47
computed average N/C ratio	0.0038	0.0039	0.0036	0.0037	0.0039	0.0038
computed average S/C ratio	0.0023	0.0026	0.0019	0.0021	0.0020	0.0024
average carbon oxidation state (Cos)	-0.117	-0.298	-0.355	-0.318	-0.311	-0.272
average DBE	9.1	8.3	8.2	8.3	8.3	8.3
average DBE/C	0.49	0.45	0.43	0.44	0.44	0.44
average $AI_{mod}$	0.31	0.28	0.27	0.27	0.27	0.27
mass weighted average [Da]	399.8	383.8	384.1	389.0	387.8	390.9
Characterization of DOM May	P1	P2	P3	P4	P5	P6
number of DOM compositions	2388	2601	2776	2523	2528	2235
average H [%]	43.8	45.7	45.2	44.9	45.3	46.2
average C [%]	38.1	37.4	37.3	37.2	37.2	37.4
average O [%]	18.0	16.7	17.2	17.8	17.2	16.3
average N [%]	0.14	0.14	0.14	0.14	0.15	0.14
average S [%]	0.05	0.08	0.10	0.09	0.08	0.05
computed average H/C ratio	1.15	1.22	1.21	1.21	1.22	1.24
computed average O/C ratio	0.47	0.45	0.46	0.48	0.46	0.44
computed average N/C ratio	0.0036	0.0037	0.0038	0.0037	0.0039	0.0038
computed average S/C ratio	0.0013	0.0021	0.0026	0.0023	0.0021	0.0014
average carbon oxidation state (Cos)	-0.189	-0.311	-0.275	-0.237	-0.274	-0.348
average DBE	8.9	8.4	8.4	8.5	8.3	8.2
average DBE/C	0.48	0.44	0.45	0.45	0.45	0.44
average $AI_{mod}$	0.32	0.28	0.28	0.27	0.28	0.28
mass weighted average [Da]	385.9	386.9	385.5	396.9	385.0	380.4
Characterization of DOM Aug	P1	P2	P3	P4	P5	P6
number of DOM compositions	2187	2231	2297	2354	2056	2234
average H [%]	44.4	46.8	46.5	45.3	45.4	45.7
average C [%]	37.7	37.2	37.2	37.0	36.9	37.0
average O [%]	17.7	15.8	16.1	17.5	17.5	17.2
average N [%]	0.13	0.14	0.14	0.14	0.13	0.14
average S [%]	0.05	0.06	0.06	0.08	0.07	0.06

computed average H/C ratio	1.18	1.26	1.25	1.22	1.23	1.24
computed average O/C ratio	0.47	0.42	0.43	0.47	0.48	0.46
computed average N/C ratio	0.0035	0.0037	0.0038	0.0037	0.0036	0.0038
computed average S/C ratio	0.0013	0.0016	0.0016	0.0022	0.0019	0.0017
average carbon oxidation state (Cos)	-0.222	-0.393	-0.364	-0.265	-0.265	-0.293
average DBE	8.7	8.0	8.1	8.4	8.2	8.2
average DBE/C	0.47	0.43	0.43	0.44	0.44	0.44
average AI <sub>mod</sub>	0.30	0.27	0.27	0.27	0.26	0.26
mass weighted average [Da]	388.3	378.9	380.8	395.6	389.7	388.7
Characterization of DOM Nov	P1	P2	P3	P4	P5	P6
number of DOM compositions	2362	2414	2553	2557	2334	2252
average H [%]	43.5	45.2	45.2	45.4	45.4	45.4
average C [%]	37.4	36.9	37.0	37.0	36.9	36.8
average O [%]	18.9	17.6	17.6	17.3	17.4	17.5
average N [%]	0.14	0.14	0.14	0.15	0.15	0.14
average S [%]	0.07	0.11	0.11	0.10	0.09	0.09
computed average H/C ratio	1.16	1.22	1.22	1.23	1.23	1.23
computed average O/C ratio	0.50	0.48	0.47	0.47	0.47	0.48
computed average N/C ratio	0.0037	0.0037	0.0039	0.0040	0.0040	0.0039
computed average S/C ratio	0.0020	0.0031	0.0030	0.0028	0.0024	0.0023
average carbon oxidation state (Cos)	-0.139	-0.256	-0.258	-0.275	-0.271	-0.265
average DBE	8.8	8.3	8.3	8.2	8.2	8.2
average DBE/C	0.47	0.44	0.44	0.44	0.44	0.44
average AI <sub>mod</sub>	0.29	0.27	0.27	0.27	0.26	0.26
mass weighted average [Da]	396.4	393.4	391.2	388.1	387.4	391.4
Characterization of DOM Jan	P1	P2	P3	P4	P5	P6
number of DOM compositions	2859	2901	2665	2771	2546	2577
average H [%]	42.7	45.1	46.0	45.4	45.5	45.3
average C [%]	37.4	37.0	37.0	36.9	36.9	36.9
average O [%]	19.6	17.7	16.8	17.4	17.3	17.6
average N [%]	0.14	0.15	0.15	0.15	0.15	0.15
average S [%]	0.11	0.14	0.11	0.12	0.10	0.10
computed average H/C ratio	1.14	1.22	1.24	1.23	1.23	1.23
computed average O/C ratio	0.52	0.48	0.45	0.47	0.47	0.48
computed average N/C ratio	0.0038	0.0039	0.0039	0.0041	0.0041	0.0040
computed average S/C ratio	0.0030	0.0037	0.0029	0.0031	0.0028	0.0028
average carbon oxidation state (Cos)	-0.0825	-0.2470	-0.3171	-0.2731	-0.2758	-0.2544
average DBE	9.1	8.4	8.1	8.2	8.2	8.2
average DBE/C	0.48	0.45	0.43	0.44	0.44	0.44
average AI <sub>mod</sub>	0.29	0.27	0.26	0.26	0.26	0.26
mass weighted average [Da]	407.2	395.1	384.5	387.0	387.9	392.8

**Table S8.** Counts of  $m/z$  ions and computed indices of molecular formulae detected at the different stages of water treatment in STO throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). The sample points (1-7) are: 1 - Raw water. 2 - Coagulation. 3 - GAC. 4 - UV. 5 - Chloramination. 6 - Finished water. 7 - Tap water. DBE = Double bond equivalences.  $AI_{mod}$  = a modified aromaticity index.

Characterization of DOM Mar	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2185	2032	2309	2835	2289	2406	2630
average H [%]	42.9	44.7	44.6	45.5	44.7	44.9	45.3
average C [%]	37.4	37.0	37.0	37.2	36.9	36.9	37.1
average O [%]	19.5	18.1	18.2	17.1	18.1	18.0	17.4
average N [%]	0.11	0.09	0.10	0.13	0.10	0.14	0.14
average S [%]	0.08	0.11	0.13	0.08	0.12	0.09	0.08
computed average H/C ratio	1.15	1.21	1.21	1.23	1.21	1.22	1.22
computed average O/C ratio	0.52	0.49	0.49	0.46	0.49	0.49	0.47
computed average N/C ratio	0.0029	0.0024	0.0026	0.0036	0.0028	0.0037	0.0037
computed average S/C ratio	0.0022	0.0031	0.0035	0.0022	0.0032	0.0026	0.0022
average carbon oxidation state (Cos)	-0.095	-0.213	-0.210	-0.291	-0.215	-0.228	-0.266
average DBE	9.3	8.6	8.7	8.5	8.7	8.7	8.6
average DBE/C	0.48	0.45	0.45	0.44	0.45	0.45	0.44
average $AI_{mod}$	0.29	0.27	0.27	0.27	0.26	0.26	0.27
mass weighted average [Da]	417.3	407.8	410.8	398.7	411.2	411.3	403.2
Characterization of DOM May	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2719	2762	2640	1534	2774	1648	1678
average H [%]	44.7	45.5	46.0	46.1	46.2	46.3	46.4
average C [%]	37.7	37.2	37.1	36.9	37.2	37.0	37.0
average O [%]	17.4	17.1	16.8	16.9	16.4	16.6	16.4
average N [%]	0.15	0.13	0.14	0.12	0.15	0.11	0.11
average S [%]	0.05	0.08	0.06	0.04	0.06	0.04	0.04
computed average H/C ratio	1.18	1.22	1.24	1.25	1.24	1.25	1.26
computed average O/C ratio	0.46	0.46	0.45	0.46	0.44	0.45	0.44
computed average N/C ratio	0.0039	0.0035	0.0039	0.0033	0.0039	0.0029	0.0030
computed average S/C ratio	0.0013	0.0022	0.0017	0.0011	0.0016	0.0012	0.0012
average carbon oxidation state (Cos)	-0.245	-0.288	-0.320	-0.321	-0.341	-0.339	-0.351
average DBE	8.8	8.5	8.4	8.3	8.3	8.3	8.2
average DBE/C	0.46	0.44	0.43	0.43	0.43	0.43	0.43
average $AI_{mod}$	0.30	0.27	0.26	0.26	0.27	0.26	0.26
mass weighted average [Da]	392.5	398.0	400.1	400.3	391.5	396.1	395.7
Characterization of DOM Aug	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	1688	1687	1412	1187	1119	1269	1523
average H [%]	43.5	46.1	45.9	45.3	45.2	45.3	45.2
average C [%]	37.3	37.0	37.0	36.9	36.8	36.9	36.9
average O [%]	19.0	16.8	17.0	17.6	17.8	17.7	17.8
average N [%]	0.10	0.10	0.09	0.07	0.07	0.08	0.09
average S [%]	0.05	0.05	0.04	0.05	0.05	0.06	0.06

computed average H/C ratio	1.16	1.24	1.24	1.23	1.23	1.23	1.22
computed average O/C ratio	0.51	0.45	0.46	0.48	0.48	0.48	0.48
computed average N/C ratio	0.0026	0.0027	0.0025	0.0020	0.0019	0.0021	0.0023
computed average S/C ratio	0.0013	0.0013	0.0011	0.0013	0.0013	0.0015	0.0017
average carbon oxidation state (Cos)	-0.133	-0.323	-0.305	-0.262	-0.249	-0.255	-0.248
average DBE	9.0	8.3	8.2	8.2	8.2	8.2	8.4
average DBE/C	0.47	0.43	0.43	0.44	0.44	0.44	0.44
average AI <sub>mod</sub>	0.29	0.26	0.26	0.26	0.26	0.26	0.26
mass weighted average [Da]	407.8	397.1	393.0	390.8	390.0	392.4	399.1
Characterization of DOM Nov	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	1589	1696	1581	2139	1406	2902	2717
average H [%]	43.1	44.7	45.1	45.3	45.6	45.2	45.8
average C [%]	37.3	37.0	37.0	37.1	36.9	37.0	37.1
average O [%]	19.4	18.1	17.8	17.5	17.4	17.6	16.9
average N [%]	0.09	0.09	0.09	0.11	0.08	0.13	0.14
average S [%]	0.05	0.07	0.06	0.07	0.05	0.10	0.07
computed average H/C ratio	1.16	1.21	1.22	1.22	1.24	1.22	1.24
computed average O/C ratio	0.52	0.49	0.48	0.47	0.47	0.48	0.45
computed average N/C ratio	0.0025	0.0025	0.0024	0.0030	0.0022	0.0036	0.0037
computed average S/C ratio	0.0014	0.0019	0.0017	0.0018	0.0014	0.0026	0.0019
average carbon oxidation state (Cos)	-0.106	-0.218	-0.243	-0.261	-0.275	-0.258	-0.310
average DBE	8.9	8.5	8.4	8.6	8.2	8.4	8.3
average DBE/C	0.48	0.45	0.44	0.44	0.44	0.44	0.44
average AI <sub>mod</sub>	0.29	0.27	0.27	0.27	0.26	0.27	0.27
mass weighted average [Da]	404.2	400.2	398.0	404.7	394.1	395.2	392.2
Characterization of DOM Jan	P1	P2	P3	P4	P5	P6	P7
number of DOM compositions	2198	2214	2237	2194	2206	2048	2021
average H [%]	43.0	44.7	44.8	44.8	44.7	44.8	44.8
average C [%]	37.3	36.9	36.9	36.9	36.8	36.8	36.8
average O [%]	19.5	18.2	18.1	18.1	18.3	18.2	18.2
average N [%]	0.10	0.10	0.10	0.09	0.10	0.09	0.09
average S [%]	0.09	0.12	0.11	0.11	0.12	0.12	0.11
computed average H/C ratio	1.15	1.21	1.21	1.21	1.21	1.22	1.22
computed average O/C ratio	0.52	0.49	0.49	0.49	0.50	0.49	0.49
computed average N/C ratio	0.0026	0.0026	0.0026	0.0025	0.0026	0.0024	0.0025
computed average S/C ratio	0.0025	0.0033	0.0030	0.0030	0.0033	0.0032	0.0030
average carbon oxidation state (Cos)	-0.098	-0.210	-0.220	-0.217	-0.205	-0.213	-0.214
average DBE	9.1	8.5	8.6	8.6	8.6	8.5	8.5
average DBE/C	0.48	0.45	0.45	0.45	0.45	0.45	0.45
average AI <sub>mod</sub>	0.29	0.26	0.27	0.27	0.26	0.26	0.26
mass weighted average [Da]	414.1	406.1	407.8	407.8	408.9	406.7	406.0

**Table S9.** Counts of  $m/z$  ions and computed indices of molecular formulae detected at the different stages of water treatment in MAL throughout five sample occasions as computed from negative electrospray 12T FT-ICR mass spectra for singly charged ions reported in neutral form (the mass of a proton added). The sample points (1-8) are: 1 - Raw water. 2 - Aeration. 3 - Softening. 4 - Rapid sand filtr.. 5 - UV. 6 - Chloramination 7 - Finished water. 8 - Tap water. DBE = Double bond equivalences.  $AI_{mod}$  = a modified aromaticity index.

Characterization of DOM Mar	P1	P2	P3	P4	P5	P6	P7	P8
number of DOM compositions	2062	2157	2276	2278	NA	2164	2036	2029
average H [%]	44.4	44.3	43.9	44.1	NA	44.4	44.7	44.3
average C [%]	37.7	37.8	37.9	38.0	NA	37.9	37.9	37.3
average O [%]	17.6	17.7	17.9	17.6	NA	17.5	17.3	18.0
average N [%]	0.14	0.16	0.19	0.18	NA	0.17	0.17	0.22
average S [%]	0.05	0.05	0.06	0.06	NA	0.05	0.04	0.11
computed average H/C ratio	1.18	1.17	1.16	1.16	NA	1.17	1.18	1.19
computed average O/C ratio	0.47	0.47	0.47	0.46	NA	0.46	0.46	0.48
computed average N/C ratio	0.0038	0.0043	0.0049	0.0048	NA	0.0046	0.0045	0.0058
computed average S/C ratio	0.0014	0.0014	0.0015	0.0015	NA	0.0013	0.0011	0.0029
average carbon oxidation state (Cos)	-0.233	-0.228	-0.205	-0.224	NA	-0.237	-0.256	-0.212
average DBE	9.1	9.1	9.2	9.1	NA	9.0	8.9	8.7
average DBE/C	0.46	0.47	0.48	0.47	NA	0.47	0.46	0.46
average $AI_{mod}$	0.30	0.30	0.31	0.31	NA	0.30	0.30	0.29
mass weighted average [Da]	408.3	404.8	402.2	400.2	NA	398.3	396.3	398.1
Characterization of DOM May	P1	P2	P3	P4	P5	P6	P7	P8
number of DOM compositions	2106	2209	2254	2117	NA	1999	1914	1921
average H [%]	44.7	44.2	43.8	44.0	NA	44.2	44.5	44.5
average C [%]	37.8	38.1	37.8	37.8	NA	37.7	37.6	37.4
average O [%]	17.3	17.5	18.1	18.0	NA	17.9	17.7	17.9
average N [%]	0.16	0.17	0.18	0.18	NA	0.17	0.17	0.20
average S [%]	0.05	0.06	0.07	0.06	NA	0.05	0.05	0.08
computed average H/C ratio	1.18	1.16	1.16	1.16	NA	1.17	1.18	1.19
computed average O/C ratio	0.46	0.46	0.48	0.47	NA	0.47	0.47	0.48
computed average N/C ratio	0.0041	0.0044	0.0046	0.0047	NA	0.0046	0.0046	0.0054
computed average S/C ratio	0.0013	0.0015	0.0018	0.0015	NA	0.0014	0.0014	0.0022
average carbon oxidation state (Cos)	-0.258	-0.234	-0.193	-0.204	NA	-0.213	-0.229	-0.225
average DBE	8.9	9.1	9.1	9.2	NA	9.0	8.9	8.7
average DBE/C	0.46	0.47	0.47	0.47	NA	0.47	0.46	0.46
average $AI_{mod}$	0.30	0.31	0.30	0.30	NA	0.30	0.29	0.29
mass weighted average [Da]	396.3	396.6	403.3	404.3	NA	400.2	399.6	397.8
Characterization of DOM Aug	P1	P2	P3	P4	P5	P6	P7	P8
number of DOM compositions	2108	2047	2230	2135	NA	1930	1945	1901
average H [%]	44.7	44.2	44.0	43.8	NA	44.0	44.2	44.1
average C [%]	37.9	37.8	37.9	37.7	NA	37.5	37.6	37.4
average O [%]	17.2	17.8	17.9	18.3	NA	18.3	17.9	18.2
average N [%]	0.16	0.15	0.18	0.17	NA	0.17	0.16	0.19
average S [%]	0.05	0.05	0.06	0.07	NA	0.05	0.05	0.07

computed average H/C ratio	1.18	1.17	1.16	1.16	NA	1.17	1.18	1.18
computed average O/C ratio	0.45	0.47	0.47	0.48	NA	0.49	0.48	0.49
computed average N/C ratio	0.0043	0.0041	0.0048	0.0045	NA	0.0044	0.0043	0.0050
computed average S/C ratio	0.0013	0.0013	0.0016	0.0019	NA	0.0014	0.0012	0.0018
average carbon oxidation state (Cos)	-0.259	-0.216	-0.207	-0.183	NA	-0.189	-0.212	-0.197
average DBE	8.9	9.2	9.1	9.1	NA	9.0	9.0	8.9
average DBE/C	0.47	0.47	0.47	0.47	NA	0.47	0.47	0.46
average AI <sub>mod</sub>	0.30	0.30	0.31	0.30	NA	0.29	0.29	0.29
mass weighted average [Da]	395.2	407.5	400.9	406.2	NA	405.1	405.5	405.5
Characterization of DOM Nov	P1	P2	P3	P4	P5	P6	P7	P8
number of DOM compositions	1927	1872	2071	2070	2181	1881	2021	1902
average H [%]	44.3	43.8	43.6	43.6	43.6	43.7	44.1	44.5
average C [%]	37.7	37.8	37.8	37.7	37.7	37.5	37.6	37.4
average O [%]	17.8	18.2	18.5	18.4	18.4	18.6	18.1	17.8
average N [%]	0.14	0.15	0.17	0.17	0.16	0.16	0.17	0.23
average S [%]	0.05	0.06	0.07	0.07	0.08	0.06	0.05	0.07
computed average H/C ratio	1.17	1.16	1.15	1.16	1.16	1.16	1.17	1.19
computed average O/C ratio	0.47	0.48	0.49	0.49	0.49	0.50	0.48	0.48
computed average N/C ratio	0.0038	0.0039	0.0044	0.0044	0.0044	0.0042	0.0045	0.0062
computed average S/C ratio	0.0014	0.0015	0.0018	0.0018	0.0020	0.0016	0.0014	0.0019
average carbon oxidation state (Cos)	-0.222	-0.188	-0.168	-0.173	-0.171	-0.165	-0.199	-0.226
average DBE	9.1	9.3	9.3	9.2	9.2	9.1	9.0	8.7
average DBE/C	0.47	0.47	0.48	0.48	0.48	0.47	0.47	0.46
average AI <sub>mod</sub>	0.30	0.30	0.30	0.30	0.30	0.29	0.29	0.29
mass weighted average [Da]	407.0	411.8	410.8	409.9	409.1	411.3	404.1	395.4
Characterization of DOM Jan	P1	P2	P3	P4	P5	P6	P7	P8
number of DOM compositions	2318	2270	2123	2095	2171	2072	2136	1960
average H [%]	44.0	43.9	44.1	44.1	44.1	44.4	44.1	45.4
average C [%]	38.0	37.9	37.8	37.7	37.7	37.5	37.5	37.0
average O [%]	17.8	18.0	17.9	17.9	18.0	18.0	18.2	17.4
average N [%]	0.18	0.17	0.18	0.17	0.18	0.16	0.17	0.10
average S [%]	0.06	0.06	0.06	0.06	0.06	0.06	0.07	0.10
computed average H/C ratio	1.16	1.16	1.17	1.17	1.17	1.18	1.17	1.23
computed average O/C ratio	0.47	0.48	0.47	0.48	0.48	0.48	0.49	0.47
computed average N/C ratio	0.0047	0.0045	0.0047	0.0046	0.0047	0.0043	0.0044	0.0027
computed average S/C ratio	0.0015	0.0017	0.0015	0.0017	0.0017	0.0016	0.0018	0.0026
average carbon oxidation state (Cos)	-0.212	-0.199	-0.210	-0.208	-0.206	-0.215	-0.195	-0.273
average DBE	9.3	9.3	9.2	9.1	9.1	8.9	9.0	8.2
average DBE/C	0.47	0.47	0.47	0.47	0.47	0.46	0.47	0.44
average AI <sub>mod</sub>	0.31	0.30	0.30	0.30	0.30	0.29	0.29	0.27
mass weighted average [Da]	406.3	408.8	408.1	406.0	403.9	402.0	404.2	389.6

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