

1 **Impact of natural NOM and AOM in Vacuum UV treatment**

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9 **Supplementary Information**

10 SUVA values before and after VUV treatment

11 SUVA, defined as UVA_{254} normalized to DOC, is a good predictor of the aromaticity of NOM. In 1999,
12 Edzwald and Tobiason proposed that a SUVA value greater than 4 indicates the predominance of aquatic
13 humic matter, highly aromatic and hydrophobic character, and high molecular weight. While SUVA values
14 between 2 and 4, are composed by a mixture of aquatic humic and non-humic matter, with a mix of aromatic
15 and aliphatic character and low to high molecular weight. When SUVA lower than 2, then a high fraction of
16 non-humic matter is present, aliphatic and hydrophobic character and low molecular weight [1].

17 *Table SI. 1. SUVA, $L m^{-1} mg C^{-1}$, values before and after VUV treatment*

		Lake			
		A	B	C	C*
Raw Water	Untreated (retention time = 0 sec)	2.70	4.27	4.01	3.89
	VUV treated (retention time = 9.4 sec)	2.49	3.97	3.84	3.48
Filtered water	Untreated (retention time = 0 sec)	3.06	-	2.71	2.73
	VUV treated (retention time = 9.4 sec)	2.91	-	2.55	2.48

19 LC-OCD-OND chromatograph

20 Figure SI 1 (a, d, g and j) shows the ultraviolet detector (UVD) signal from the LC-OCD-OND for the four
21 lakes. Without VUV treatment, in Lake A (where a bank filtration water system is responsible for NOM
22 removal), the UVD signal in filtered water is 25% lower than in raw water. Among the four lakes,
23 Lake B shows the higher UVD signal due to the high cell concentration (almost 4 times higher than
24 Lake A and two times Lake C and C*). DOC measurement was stable and UVD declined, thus modest
25 reductions of SUVA were obtained (about 0.2-0.4 units) (Table SI.1). In terms of TOC in Lake C and C*, it
26 also remained stable (reductions: -1% to -6%). This was expected given the low fluence applied.

27 DOC characterization by LC-OCD-OND are depicted in Figure SI 1 (b, e, h, and k). The figure revealed a
28 change in the molecular weight of NOM components after VUV treatment. While the overall reduction in
29 DOC was small, humics were strongly bleached (strong reduction in UV_{254} response). An increase in
30 biodegradability is observed through the increase in BDOC measurements (from +20% to +35%. Figure
31 SI.2) as the HMW fraction decreases.

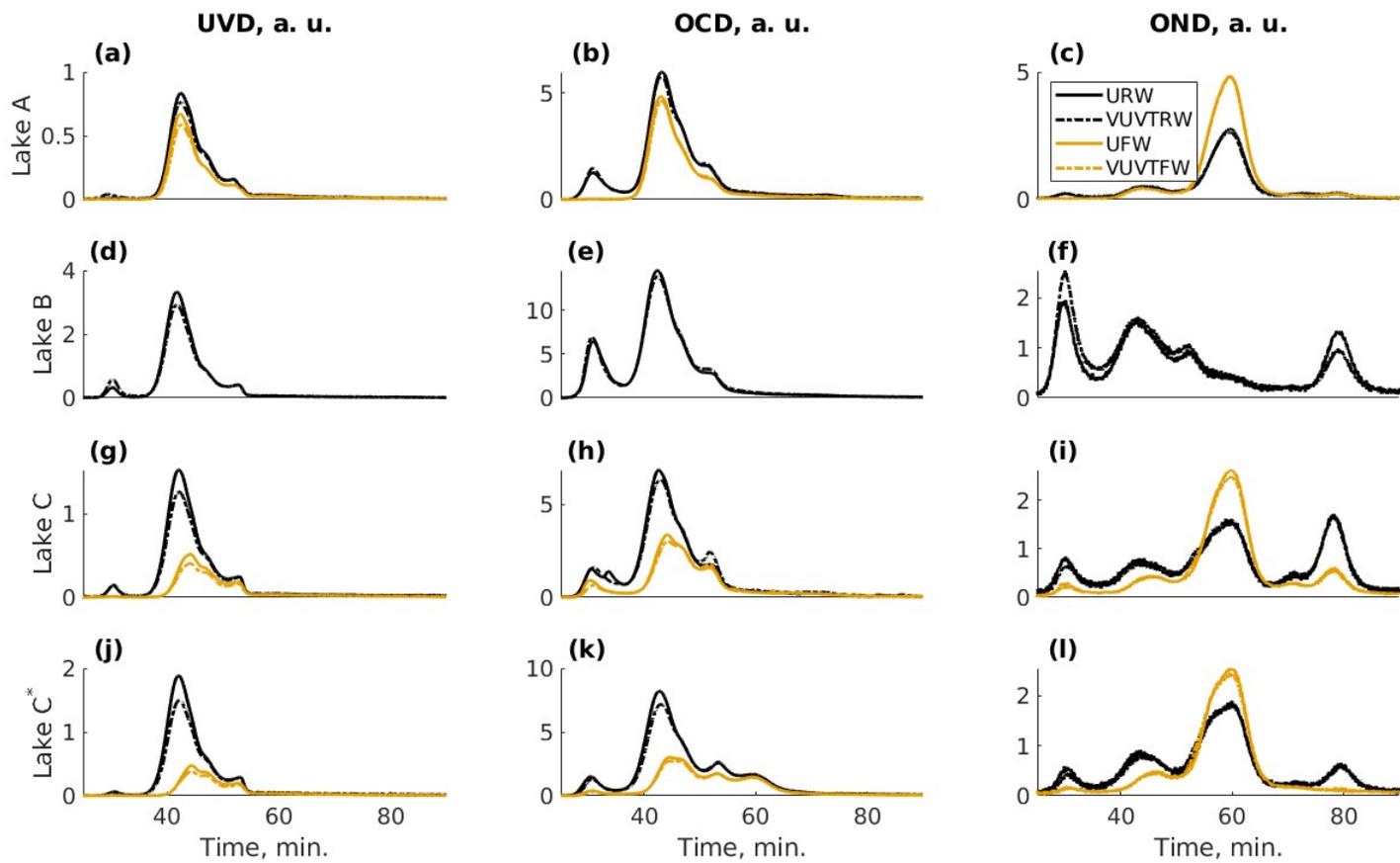


Figure SI. 1. LC-OCD-OND spectra (URW = Untreated Raw Water, VUVTRW = VUV Treated Raw Water, UFW = Untreated Filtered Water, UVTFW = VUV Treated Filtered Water, a.u. = arbitrary units)

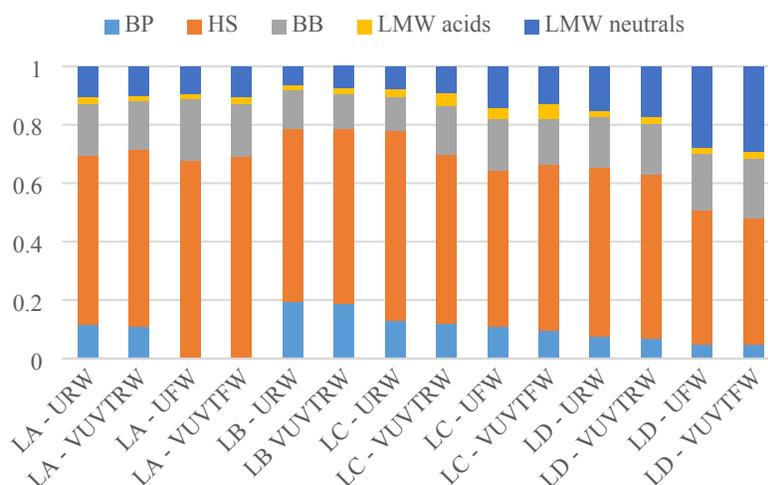


Figure SI. 2. Carbon fraction from LC-OCD-OND
 Lake A, LB = Lake B, LC = Lake C, LD = Lake C*, URW = untreated raw water, VUVRW = VUV treated raw water

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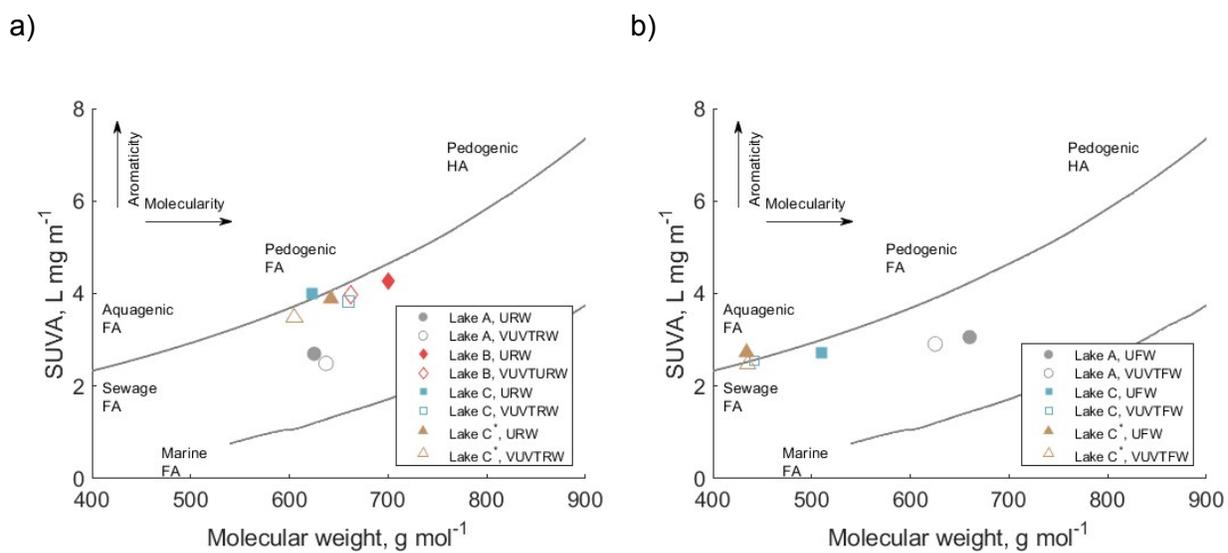


Figure SI. 3. Humic substances diagram (HS-diagram) before and after (retention time=9.4 sec) VUV treatment. a) Raw water: URW = untreated raw water, VUVTRW VUV treated raw water. b) Filtered water: UFW = untreated filtered water, VUVTFW VUV treated filtered water. Based on [2]

34 Biodegradability though biological dissolved organic carbon (BDOC) measurements

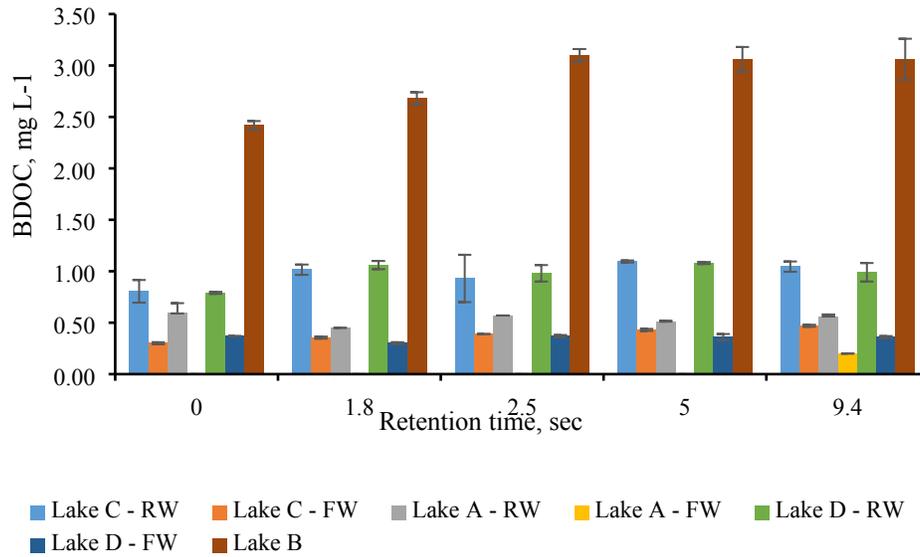


Figure SI. 4. BDCO before and after VUV treatment. RW = Raw Water, FW = Filtered Water

35 PARAFAC modeling

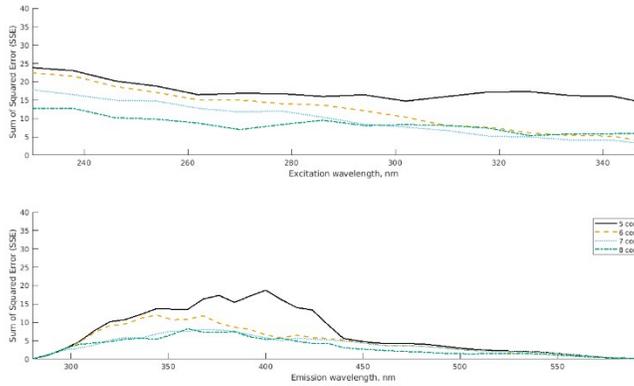


Figure SI. 5. SSE of PARAFAC models with different number of component (for supplementary information)

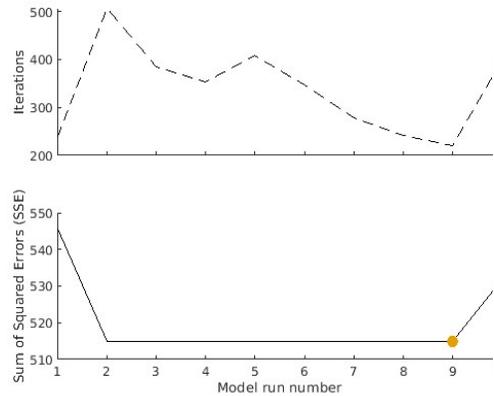


Figure SI. 6. Robustness of the 7-components PARAFAC model. Ten runs with random starting points (for supplementary information)

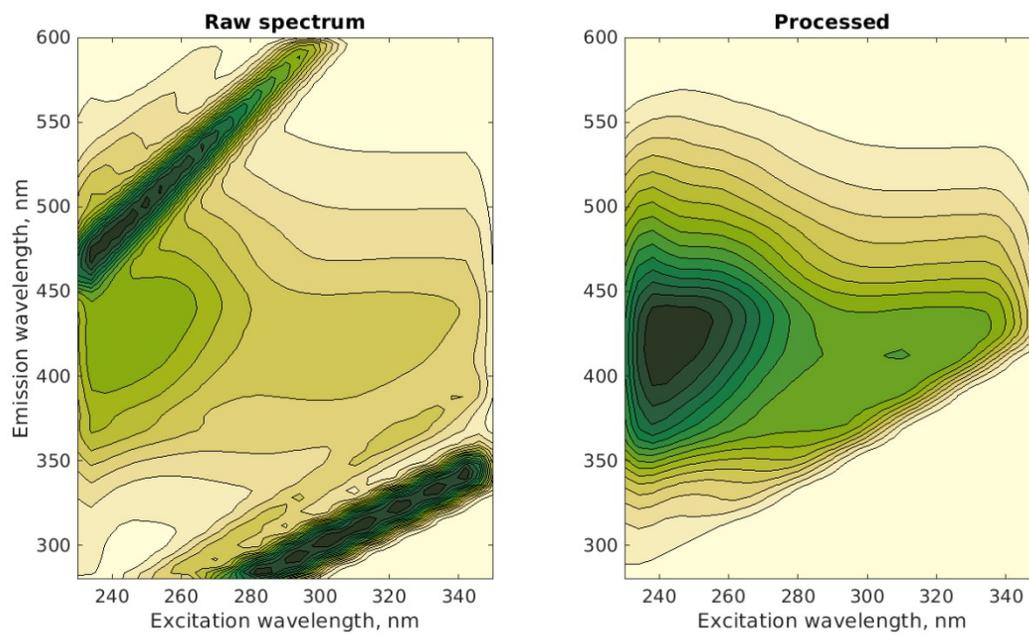


Figure SI. 7. Dataset pre-processing (for supplementary information)

36 PARAFAC Modeling components

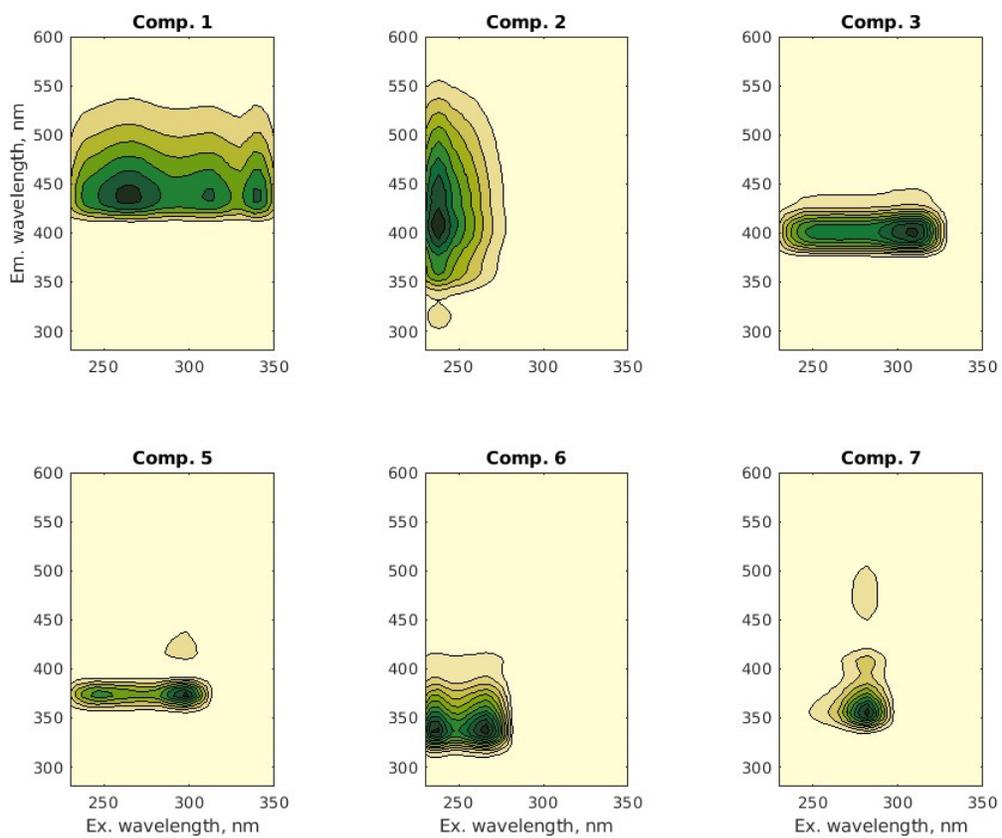


Figure SI. 8. Components obtained from PARAFAC modelling

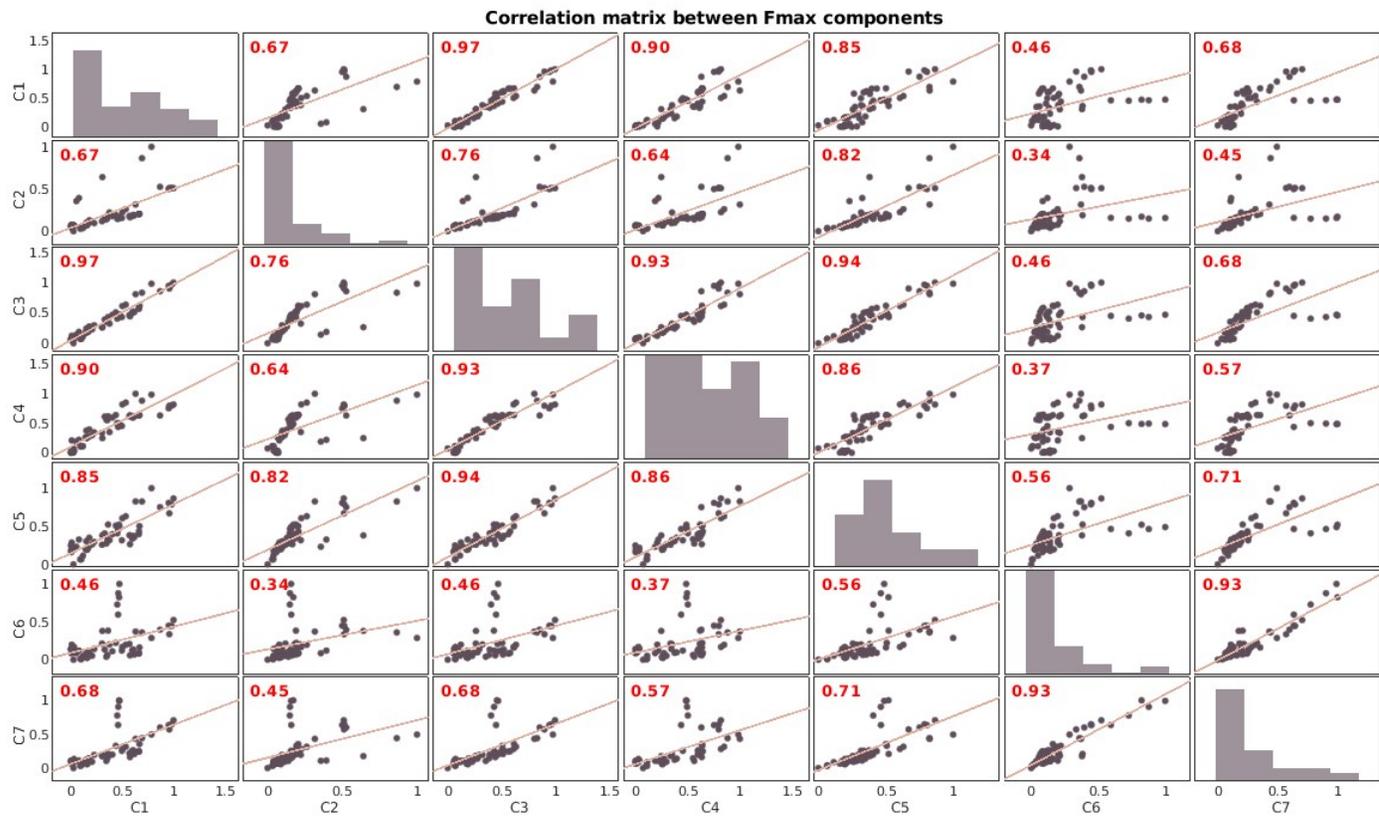


Figure SI. 9. F_{max} correlation between PARAFAC components

39 SOM modeling

40 SOM is a pattern recognition method. It clusterizes and reduces the dimensionality of input FEEM without
41 make assumption about the data structure [3]. A dataset of 64 spectra were analyzed with the PARAllel
42 FACtor analysis (PARAFAC) decomposition routines for Excitation Emission Matrices (drEEM, version
43 0.2.0 toolbox [4]) and self-organizing maps (SOM) with the SOM-Toolbox version 2.1 [5] both running on
44 Matlab® R2018B software).

45 Figure SI 8 shows the U-matrix from the best SOM map selected along with the component planes of each
46 of the components retrieved by PARAFAC. The U-matrix shows the distance between neighbouring
47 neurons in the map, where large values (darker color) indicate highly dissimilar neurons. The component
48 planes are useful to visualize the importance of different excitation-emission pairs or variables for each
49 sample (neurons, from the SOM solution). We selected the pairs corresponding to the components
50 proportions from PARAFAC (Figure SI.7).

51 The distribution of the samples obtained from the solution of SOM is shown through the best matching unit
52 (BMU) graphics. In Figure SI 8 we show a combined image of the BMU overlapped with multiple hits
53 histograms (colored hexagons) for each lake showing the importance of each neuron representing the lakes
54 and the type of water (the larger the colored hexagon the higher number of hits in that neuron). The BMU
55 is the best solution found by SOM and shows the distribution of the four lakes across the map for raw (R)
56 and filtered (F) water. For example, samples associated to Lake B are located at different neurons in the
57 map (the top-left and bottom-left in Figure SI 7), indicating the differences in NOM composition from raw
58 and filtered. Basically, raw water contains NOM and AOM (particulate + dissolved) and filtered water only
59 contains the dissolved fraction of NOM and AOM. Thus, SOM maps locate raw and filtered water in different
60 region (Figure SI 7). Therefore, SOM allows discriminating differences in cell concentration between
61 samples. We identify two clusters for Lake A: one (bottom-right) belongs to raw water from the lake while
62 the other (middle) corresponds to filtered water through the well (Lake A supplies a bank filtration water
63 system). Lakes C and C* show share similar neurons in the map, because they represent the same lake
64 without (Lake C) and with (Lake C*) bloom conditions. For a given lake is seen that the SOM solution

65 properly discriminate between raw and filtered water, particularly those with bloom condition (Lakes B and
66 C*).

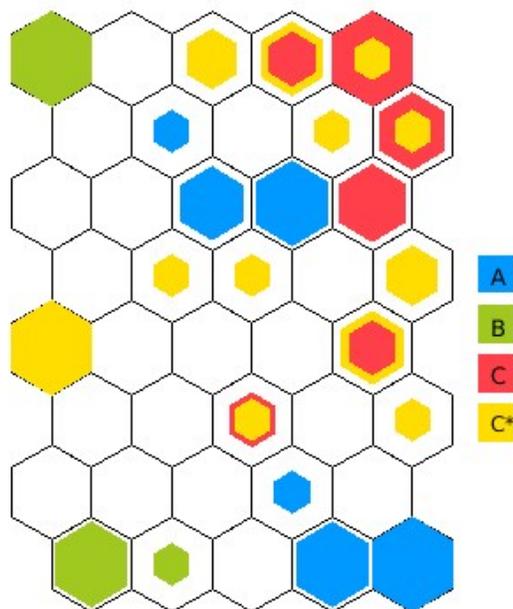


Figure SI. 10. SOM map with multiple hits overlapped showing the sites distribution for raw and filtered water.

67 By comparing Figure SI 7 and each component plane from Figure SI 8 is possible to correlate that
68 PARAFAC components C1(1), C6(1), C6(2) and C7(1) characterize Lake B. Furthermore, C6(1), C6(2) and
69 C7(1) are essentially due to samples from Lake B. This allows us to conclude that Lake B is mostly
70 characterized by humic acids-like and (Region V, Figure 3) and soluble microbial by-product-like (Region
71 IV, Figure 3). Component planes show that raw water in Lake A is associated to C1(1), C5(1), C5(2) and
72 C7(2), while filtered water correlates to C1(2), C3(1), C4(1) and C7(2). Here also the regions that
73 characterize the lake are V and IV (Figure 3). Lakes C and C* are characterized by C1(2), C1(3), C3(2),
74 C4(1) and C7(2), lying in regions V and IV.

75 By overlapping PARAFAC and SOM results, it is possible to distinguish components that characterise each
76 lake. All four lake could be characterize by a reduced number of regions (Figure 3): region IV(humic acid-

77 like) and V (soluble microbial by-product-like). By focusing on this FEEM region, it could be possible to
78 make a relationship with NOM / AOM and DBP's formation and reactivity.

79

Table SI. 2. SOM and PARAFAC. Characteristic component of each lake

Lake	PARAFAC Component
A	C1(1), C5(1), C5(2) and C7(2) (Raw water)
	C1(2), C3(1), C4(1) and C7(2) (filtered water)
B	C1(1), C6(1), C6(2) and C7(1)
C and C*	C1(2), C1(3), C3(2), C4(1) and C7(2)

80 Combination of PARAFAC components and SOM allowed to characterize each lake with fewer
81 components:

- 82 ▪ Components C1(1), C6(1), C6(2) and C7(1) characterize Lake B
- 83 ▪ Raw water in Lake A is associated to C1(1), C5(1), C5(2) and C7(2), while filtered water
84 correlates to C1(2), C3(1), C4(1) and C7(2)

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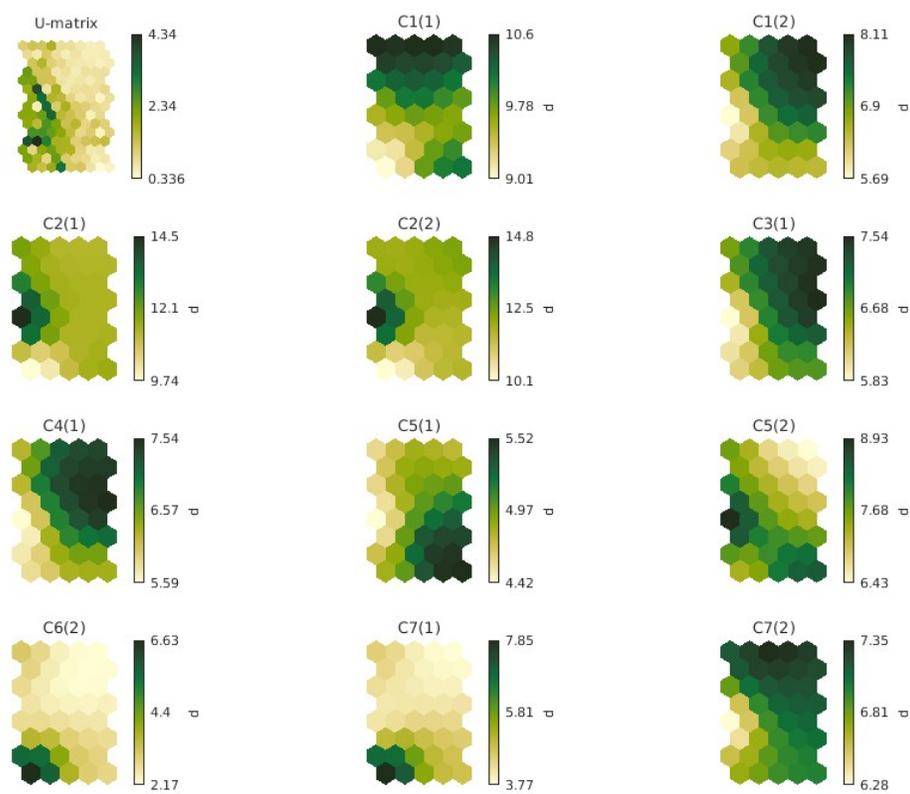


Figure SI. 11. U-matrix and component planes for the proportions of PARAFAC components resulted from SOM analysis

87 Disinfection by-products formation: correlation data

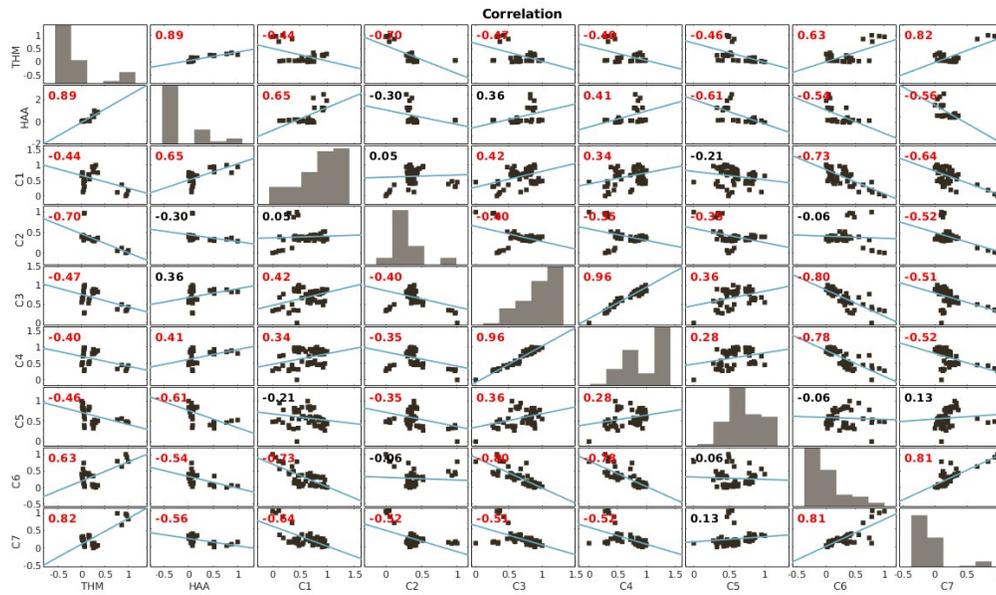


Figure SI. 12. THM and HAA correlation with PARAFAC components

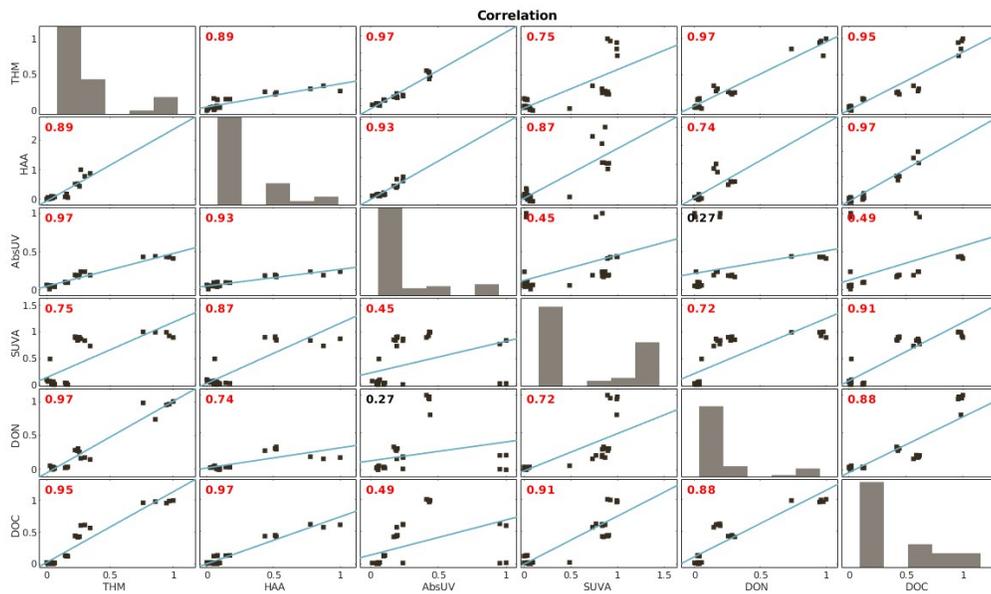


Figure SI. 13. THM and HAA correlation with UVA₂₅₄, SUVA, DON and DOC

Table SI. 4. Carbon fractions and yields values from DBP's formation correlation

C or N fraction (F)	Yields, $\mu\text{g}_F / \mu\text{g}_{\text{THM or HAA}}$	
	THM	HAA
DOC	27	43
DON	0.9	0.6
HS	20	28
BP	6.9	4.7
BB	3.8	7.2
LMW Ac	0.51	1.2
LMW Neutral	1.9	6

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