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1	Chloramination of wastewater effluent: Toxicity and formation of disinfection
2	byproducts
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#### 19 Abstract

20 The reclamation and disinfection of waters impacted by human activities (e.g., wastewater 21 effluent discharges) are of growing interest for various applications but has been associated with 22 the formation of toxic nitrogenous disinfection byproducts (N-DBPs). Monochloramine used as 23 an alternative disinfectant to chlorine can be an additional source of nitrogen in the formation of 24 N-DBPs. Individual toxicity assays have been performed on many DBPs, but few studies have 25 been conducted with complex mixtures such as wastewater effluents. In this work, we compared 26 the cytotoxicity and genotoxicity of wastewater effluent organic matter (EfOM) before and after 27 chloramination. The toxicity of chloraminated EfOM was significantly higher than the toxicity of 28 raw EfOM, and the more hydrophobic fraction (HPO) isolated on XAD-8 resin was more toxic 29 than the fraction isolated on XAD-4 resin. More DBPs were also isolated on the XAD-8 resin. N-30 DBPs (i.e., haloacetonitriles or haloacetamides) were responsible for the majority of the 31 cytotoxicity estimated from DBP concentrations measured in the XAD-8 and XAD-4 fractions 32 (99.4% and 78.5%, respectively). Measured DBPs accounted for minor proportions of total 33 brominated and chlorinated products, which means that many unknown halogenated compounds 34 were formed and can be responsible for a significant part of the toxicity. Other non-halogenated 35 byproducts (e.g., nitrosamines) may contribute to the toxicity of chloraminated effluents as well. 36 **Keywords** 

# Disinfection byproducts, Chloramination, Toxicity, Wastewater, XAD resins, Haloacetonitriles, Haloacetamides

#### 39 Introduction

40 The disinfection of waters impacted by human activities (e.g., agriculture or wastewater
41 effluent discharges) has been associated with the formation of nitrogenous disinfection

42	byproducts (N-DBPs) due to their enrichment in nitrogen-containing compounds (e.g., ammonia
43	or organic nitrogen such as amino acids and peptides) (Bond et al., 2011; Westerhoff and Mash,
44	2002). N-DBPs generally form in lower concentrations than non-nitrogenous regulated DBPs
45	(i.e., trihalomethanes, THMs, and haloacetic acids, HAAs), but may present a higher health risk
46	(Muellner et al., 2007; Plewa et al., 2004). In vitro mammalian cell assays have demonstrated
47	that N-DBPs such as haloacetonitriles (HANs) (Muellner et al., 2007), halonitromethanes
48	(HNMs) (Plewa et al., 2004) and haloacetamides (HAcAms) (Plewa et al., 2007) exhibit orders
49	of magnitude higher levels of cytotoxicity and genotoxicity than THMs and HAAs (Plewa et al.,
50	2008). Reclaiming wastewater for various agricultural, industrial or municipal applications is of
51	growing interest but requires the practice of disinfection to prevent outbreaks of waterborne
52	diseases. Dissolved organic matter isolated from wastewater effluent (EfOM) was found to be
53	significantly more enriched in nitrogen (i.e., to be a potential source of N-DBPs) than organics
54	recovered from surface waters (e.g., N/C mass ratios up to 0.17 compared to 0.01-0.06 for river
55	waters) (Drewes and Croue, 2002; Le Roux et al., 2016; Zheng et al., 2014).
56	Moreover, the presence of bromide and iodide ion in wastewater (especially at locations where
57	potable water is produced through the desalination of seawater or brackish water) favors the
58	formation of brominated and iodinated byproducts that are often more toxic than their
59	chlorinated analogues (Plewa et al., 2008; Richardson et al., 2008, 2007). Brominated and
60	iodinated N-DBPs are among the most cytotoxic/genotoxic disinfection by-products known
61	today (Muellner et al., 2007; Plewa et al., 2008). Water utilities, especially in the USA, have
62	been increasingly switching chlorine disinfection to monochloramine to reduce the concentration
63	of regulated THMs and HAAs (U.S. Environmental Protection Agency, 2006), however,
64	monochloramine can be an additional source of nitrogen in the formation of N-DBPs (Kimura et

al., 2013; Le Roux et al., 2016). Chloramines can also be formed unintentionally from the
reaction between free chlorine and ammonia during chlorination, which may increase the risk of
N-DBP formation when high ammonia concentrations are present.

68 While toxicity assays have been conducted for many individual DBPs, few studies have been 69 performed with complex mixtures such as natural waters, drinking waters or wastewater 70 effluents. Many of the >500 DBPs reported in the literature were not analyzed for toxicological 71 effects (Richardson and Postigo, 2015). Similarly, many studies characterized DBP occurrence 72 from various sources and their formation conditions, but the evaluation of DBP formation in 73 conjunction with toxicity assays has not been extensively explored. Richardson (2011) published 74 a protocol for DBP extraction, analysis and toxicity assessment, consisting in the extraction of 75 disinfected waters by XAD resins (XAD-8 and XAD-2 in series), followed by an elution with 76 ethyl acetate. The extract is then either directly analyzed by gas chromatography coupled with 77 mass spectrometry (GC-MS) or evaporated and exchanged to dimethylsulfoxide (DMSO) for further genotoxicity/cytotoxicity analyses. This method has been used for swimming pool waters 78 79 (Liviac et al., 2010; Plewa et al., 2011; Richardson et al., 2010) and drinking waters disinfected 80 with chlorine, ozone or chlorine dioxide (Jeong et al., 2012) and was recently applied to 81 disinfected (i.e., chlorinated and ozonated) wastewater effluents (Dong et al., 2016). N-DBPs are 82 compounds of interest because of their potential toxicity, and the chloramination of EfOM is 83 expected to favor the production of this class of DBPs because nitrogen can be incorporated both 84 from monochloramine and from the nitrogenous moieties present in EfOM. 85 As a result, the aim of this work was (i) to compare the cytotoxicity and genotoxicity of EfOM

86 resin isolates recovered before and after chloramination, (ii) to analyze the toxicity of resin

87 extracts obtained from chloraminated wastewater effluent in relation with the formed DBPs, and

(iii) to estimate the contribution of N-DBPs to the toxicity of chloraminated wastewatereffluents.

90

#### 91 **1. Materials and methods**

#### 92 1.1. Materials

93 Analytical or laboratory grade reagents and were used without further purification. MilliQ 94 water was produced with a Millipore system (18.2 MΩ.cm). Sodium hypochlorite (NaOCl, 5.65-95 6%, Fisher Scientific) and ammonium chloride (Acros Organics, 99.6%) were used to prepare 96 chloramine solutions. Methyl tert-butyl ether (MTBE) and ethyl acetate (> 99%, Fisher 97 Scientific) were used for DBP extractions without further purification. A THM calibration mix 98 (chloroform - TCM, dichlorobromomethane - CHCl<sub>2</sub>Br, chlorodibromomethane - CHClBr<sub>2</sub>, and 99 bromoform - TBM), a mixed standard (EPA 551B Halogenated Volatiles Mix) containing 100 HANs, trichloronitromethane (TCNM, or chloropicrin) and haloketones (HKs), and a mixed 101 standard containing 9 HAAs (EPA 552.2 Methyl Ester Calibration Mix) were supplied from 102 Supelco (Sigma-Aldrich). Chloro-, bromo-, dichloro-, and trichloroacetamide (CAcAm, BAcAm, 103 DCAcAm and TCAcAm, respectively) were obtained from Sigma-Aldrich. Other HAcAms (i.e., 104 dibromoacetamide - DBAcAm, tribromoacetamide - TBAcAm, bromochloroacetamide -105 BCAcAm, chloroiodoacetamide - CIAcAm, bromoiodoacetamide - BIAcAm and 106 diiodoacetamide - DIAcAm) were purchased from Cansyn Chem. Corp. Haloacetaldehydes 107 (HAcAls) were obtained from TCI America, Cansyn Chem. Corp. and Sigma-Aldrich. 108 Decafluorobiphenyl (99%, Sigma-Aldrich, Supelco) was used as a surrogate standard. 2 109 bromopropionic acid (Fluka Analytical) was used as a surrogate for HAA extractions and 110 analyses.

#### 112 **1.2. Analytical methods**

113 Total Organic Carbon (TOC) and Total Nitrogen (TN) concentrations were measured using a 114 TOC analyzer equipped with a TN detection unit (TOC-VCSH, Shimadzu). Three-dimensional 115 fluorescence excitation - emission matrices (EEMs) were obtained by a Fluoromax fluorometer 116 (Horiba Scientific, Japan). Samples for Adsorbable Organic Halide (AOX) analyses were 117 processed through adsorption on activated carbon columns using a TOX sample preparatory unit 118 (TXA-03, Mitsubishi Chemical Analytech Co., Ltd., Japan). AOX were then transformed into 119 hydrogen halides by combustion (950 °C) of the activated carbon for at least 30 min via an 120 AOX-200 adsorbable halogen analyzer and then collected in Milli-Q water as chloride and 121 bromide ions. Offline quantification of chloride and bromide ions was performed by a Dionex 122 1600 reagent free ion chromatograph (IC) equipped with a conductivity detector and a Dionex 123 IonPac AS-15 column ( $2 \times 250$  mm) and using an online KOH eluent (30 mM) generator at a 124 flow rate of 0.4 mL/min. AOCl and AOBr concentrations were determined from the respective 125 Cl<sup>-</sup> and Br<sup>-</sup> concentrations. Free chlorine and total chlorine concentrations in the sodium 126 hypochlorite stock solutions were determined by spectrophotometric measurement at 292 nm. 127 NH<sub>2</sub>Cl and NHCl<sub>2</sub> concentrations in stock solutions were determined by spectrophotometric 128 measurement using their respective molar extinction coefficients at 245 nm and 295 nm. Residual oxidant during chloramination reactions was analyzed by DPD colorimetric method. 129 130 Four THMs, four HANs, two HKs, chloropicrin and seven HAcAls were extracted and analyzed 131 following EPA method 551, which consists of a liquid-liquid extraction using MTBE followed 132 by gas chromatography coupled with electron capture detector (GC-ECD) or GC-MS (Munch 133 and Hautman, 1995). Nine HAAs were extracted and analyzed following EPA method 552.2,

134 which is based on a liquid-liquid extraction with MTBE in acidic conditions followed by

135 derivatization to methyl esters using acidic methanol, and analysis by GC-MS (Munch and

136 Munch, 1995). HAcAms were analyzed following the same EPA method 551 protocol, replacing

137 MTBE by ethyl acetate for the liquid-liquid extraction. Samples were also analyzed in GC-MS

138 full scan mode to search for other DBPs and unknown mass spectra were subjected to library

139 database searching (National Institute of Standards and Technology - NIST).

140

#### 141 **1.3. Chloramination conditions**

142 Monochloramine stock solutions were prepared by adding sodium hypochlorite (NaOCl) to a 143 continuously-stirred ammonium chloride solution adjusted to pH 8.5 with sodium hydroxide, at a 144 Cl:N molar ratio of 1:1.4. The concentration of monochloramine stock solutions was adjusted to 145 a desired concentration. Chloramination reactions were performed at pH 7.7 without addition of 146 buffering agents, and doses were calculated as:  $NH_2Cl (mg/L as Cl_2) = 3 \times DOC$  (dissolved 147 organic carbon, mg C/L) (Dotson et al., 2009; Krasner et al., 2007). Preformed monochloramine 148 was used to avoid breakpoint reactions that could occur with residual ammonium concentration 149 of the effluent (0.2-1 mgN/L after prolonged aeration activated sludge). The pH remained stable 150  $(7.7 \pm 0.5)$  during all the reaction time (72 hr). At the end of the contact time, samples were 151 acidified at pH 2 with concentrated HCl and immediately subjected to the XAD resins extraction 152 protocol. No additional quenching was performed in order to preserve the disinfection 153 byproducts formed (no residual oxidant was measured after acidification), and samples were 154 directly passed through XAD resins for fractionation and extraction. 155

#### 156 **1.4. Effluent organic matter (EfOM) extraction procedure**

157 Wastewater secondary effluent samples were collected at three different periods (WW1, 2 and 158 3, Table 1) from a wastewater treatment plant (Jeddah, Saudi Arabia) before chlorine 159 disinfection. Samples were transferred to the laboratory, filtered on cartridge filters (Polygard<sup>®</sup> 160 10 µm pore size, Millipore) prior to the chloramination and fractionation experiments to remove 161 any particulate matter and stored at 4 °C for maximum 7 days. The hydrophobic (HPO) and 162 transphilic (TPI) fractions of the non-chloraminated and chloraminated (laboratory conditions) 163 wastewater EfOM were isolated using a comprehensive isolation protocol described elsewhere 164 (Croué, 2004; Leenheer et al., 2000). Briefly, wastewater effluent samples (before or after 72 hr 165 chloramination) were acidified at pH 2 with hydrochloric acid and passed through XAD-8 and 166 XAD-4 macroporous resins in series at a flow rate of 5.2 L/hr. Resins were then rinsed with 167 formic acid at pH 2 in order to remove the inorganic species present in the void volume without 168 eluting the organics adsorbed onto the resins, which were recovered afterward with either 169 acetonitrile/water (75:25 v/v), acetonitrile or ethyl acetate. During the adsorption phase, 100 mL 170 samples were collected from the output of the resins for DOC and DBP measurements. For 171 effluent WW1, organics adsorbed onto the two XAD resins and eluted with the same volume of 172 acetonitrile/water were concentrated using a rotary evaporator to remove acetonitrile. Equal 173 volumes (50:50 v/v) of the fractions obtained from the two resins were also combined before the 174 rotary evaporation step to obtain a mixed XAD-8+XAD-4 extract. The water concentrates were 175 lyophilized to obtain the XAD-8, XAD-4 and mixed XAD-8+XAD-4 solid organic fractions for 176 use in toxicity analyses. For effluent WW2, the elution of both resins was performed with pure 177 acetonitrile to perform direct DBP identification by full-scan GC-MS on a small volume (i.e., 1.5 mL) of the extracts. Milli-Q<sup>®</sup> water was then added to the acetonitrile extracts to reach 75:25 v/v 178 179 acetonitrile/water condition (same as used for effluent WW1) before rotary evaporation (removal

180 of acetonitrile) and lyophilization. With this approach the cytotoxicity and genotoxicity values of 181 samples WW1 and WW2 could be expressed as per mg of EfOM. For effluent WW3, both resins 182 were eluted with ethyl acetate and the chloraminated EfOM extracts were concentrated under a 183 stream of  $N_2$  to a final volume of 100 mL to allow the direct quantification of DBPs and the 184 analysis of toxicity from the same extract, as previously described in the literature (Jeong et al., 185 2012; Richardson, 2011). It should be noted that these concentration approaches (i.e., 186 lyophilization or N<sub>2</sub> evaporation) are usually suitable for semi- or non-volatile chemicals but not 187 for volatile compounds (Weinberg, 2009). Nevertheless, volatile DBPs (e.g., THMs) analyzed in 188 the ethyl acetate extracts after concentration were in accordance with concentrations analyzed in 189 the wastewater effluent samples (see Section 2.3.4). Since the two chloraminated wastewater 190 effluents (WW2 and WW3) were not sampled at the same time (and exhibited different DOC 191 values) and fractions were obtained through different processes (as described above), the toxicity 192 results of these two types of samples were not comparable.

193

#### 194 **1.5. Analytical biology**

#### 195 **1.5.1. Sample preparation**

The lyophilized or liquid organic fractions were received and stored at -20 °C. Mg amounts of each lyophilized fraction were placed into separate sterile Supelco sample vials. To sterilize the sample, DMSO was added to each sample such that for each mg of sample 1 µL of DMSO was added. After the sample was mixed (vortexed) in DMSO, 1 µL of sterile ddH<sub>2</sub>O was added for each mg of the lyophilized fraction. The stock solution for each sample was completely dissolved in this DMSO-water (1:1 v/v) solvent and this was stored at -20°C. Ethyl acetate organic extracts (liquid form) obtained from WW3 were solvent exchanged into DMSO so that 203 the organic material from 1 L of wastewater effluent was concentrated into 10  $\mu$ L of DMSO 204 (10<sup>5</sup>× concentration) and stored in Teflon-sealed sample vials at -20°C. For each experiment a 205 50× dilution of each sample stock solution was prepared in F12 cell culture medium+5% fetal 206 bovine serum (FBS).

#### 207 **1.5.2.** Chinese hamster ovary cells

208 Chinese hamster ovary (CHO) cell line AS52, clone 11–4–8 was used for the biological assays

209 (Wagner et al., 1998). CHO cells were maintained on glass culture plates in Ham's F12 medium

210 containing 5% FBS, 1% antibiotics (100 U/mL sodium penicillin G, 100 μg/mL streptomycin

sulfate, 0.25 µg/mL amphotericin B in 0.85% saline), and 1% glutamine at 37 °C in a humidified

atmosphere of 5% CO<sub>2</sub>.

#### 213 **1.5.3. CHO cell chronic cytotoxicity assay**

214 This assay measures the reduction in cell density on flat-bottom 96-well microplates as a 215 function of the concentration of the test sample over a period of approximately 72 hr (~3 cell 216 cycles). Microliters of the sample were diluted with F12 +FBS medium to analyze a range of 217 concentration factors. This assay was calibrated; the detailed procedure was published (Plewa 218 and Wagner, 2009). For each sample concentration factor, 4-8 replicate wells were analyzed. 219 The experiments were repeated 2-3 times. A concentration-response curve was generated for 220 each sample. The LC<sub>50</sub> values were calculated from the regression analysis and represent the 221 sample concentration factor that induced a 50% reduction in cell density as compared to the 222 concurrent negative controls. The cytotoxicity index (CTI) value was a metric that expressed increasing values with increased cytotoxic damage  $(LC_{50}^{-1})(10^3)$  and allowed for easy 223 224 comparisons among the wastewater samples isolated in the same conditions.

#### 225 **1.5.4. CHO cell single cell gel electrophoresis assay**

226 Single cell gel electrophoresis (SCGE, or Comet) assay quantitatively measures genomic 227 deoxyribonucleic acid (DNA) damage in individual nuclei induced by a test agent (Tice et al., 228 2000). We employed a microplate methodology (Wagner and Plewa, 2009). The SCGE metric 229 for genomic DNA damage was the %Tail DNA value which is the amount of DNA that migrated 230 from the nucleus into the microgel (Kumaravel and Jha, 2006). Within each concentration factor 231 range with >70% cell viability, a concentration-response curve was generated and regression 232 analysis was used to fit the curve. The concentration factor inducing a 50% Tail DNA (TDNA) 233 value was calculated using a regression analysis for each concentration-response curve. The 234 genotoxic index (GTI) was a metric that expressed increasing values with increased genotoxic 235 damage  $(50\% \text{ TDNA}^{-1})(10^3)$  and allowed for easy comparisons among the evaluated wastewater 236 samples.

#### 237 1.5.5. Statistical analyses

238 For the cytotoxicity assay, a one-way analysis of variance (ANOVA) test was conducted to 239 determine if the sample induced a statistically significant level of cell death as compared to the 240 concurrent negative controls at a specific concentration factor. If a significant F value ( $P \le 0.05$ ) 241 was obtained, a Holm-Sidak multiple comparison versus the control group analysis was 242 performed to identify the lowest cytotoxic concentration factor. The power of the test statistic 243  $(1-\beta)$  was maintained as  $\geq 0.8$  at  $\alpha = 0.05$ . For the SCGE assay, the %Tail DNA values were not 244 normally distributed which limits the use of parametric statistics (Box et al., 1978). The mean 245 %Tail DNA value for each microgel was calculated and these values were averaged among all 246 the microgels for each wastewater sample concentration factor. Averaged mean values express a 247 normal distribution according to the central limit theorem. An ANOVA test was conducted on

- these averaged %Tail DNA values (Lovell and Omori, 2008). If a significant F value of  $P \le 0.05$
- 249 was obtained, a Holm-Sidak multiple comparison versus the control group analysis was
- 250 conducted (power  $\geq 0.8$ ;  $\alpha = 0.05$ ).
- 251
- 252 **2. Results and discussion**
- 253 2.1. EfOM extraction recovery
- 254 The EfOM extraction conditions and recovery for the three different batches of wastewater
- effluents (WW1, 2 and 3) are presented in Table 1 and 2.

Wastewater effluent sample	Chloramination	Extracted volumes (L)	DOC before / after chloramination (mg C/L)	Elution solvent	Fractions phase
WW1	No	150	4.65 / N.A.	Acetonitrile/H <sub>2</sub> O	Solid
WW2	Yes	69	4.68 / 4.55	Acetonitrile	Solid
WW3	Yes	76	4.97 / 4.60	Ethyl acetate	Liquid
N.A. = Not Applicable; DOC: dissolved organic carbon					

256 Table 1. Summary of experiments and extraction conditions

257

258 The first experiment consisted in the extraction of 150 L non-chloraminated wastewater

effluent (WW1, 4.65 mg C/L) followed by an elution with acetonitrile/H<sub>2</sub>O (75:25, v/v). 69.5%

260 DOC was retained on both XAD resins (2.13 mg C/L on XAD-8, 1.10 mg C/L on XAD-4)

261 (Table 2). 1.42 mg C/L was measured at the output of the XAD-4 resin (consisting in the

- 262 hydrophilic fraction of EfOM). The second extraction was performed with 69 L wastewater
- effluent (WW2, 4.68 mg C/L) after 72 hr of chloramination (14.2 mg Cl<sub>2</sub>/L as preformed
- 264 monochloramine, pH 7.7), followed by an elution with acetonitrile (no water was added to be

265	able to analyze the recovered fractions by GC-MS, see Section 2.3.1). Less DOC (i.e., humic-
266	like substances) was adsorbed onto the XAD-8 resin (1.56 mg C/L) than from the non-
267	chloraminated effluent (2.13 mg C/L), while WW1 and WW2 showed similar DOC content (4.65
268	and 4.55 mg C/L, respectively). Similar amount of DOC was adsorbed onto the XAD-4 resin,
269	i.e., 1.10 and 1.05 mg C/L for WW1 and WW2, respectively. The observed lower retention of
270	chloraminated EfOM onto the XAD-8 resin could be attributed to the breakdown of large organic
271	macromolecules (i.e., humic-like substances) into smaller hydrophilic molecules during
272	oxidation by chloramines. Accordingly, fluorescence EEMs obtained before and after
273	chloramination of wastewater effluent WW2 (Fig. 1) exhibited a decrease of 26.5% in signal
274	intensity after chloramination at the wavelengths showing the most intense response (i.e.,
275	excitation: 350 nm, emission: 420 nm), indicating that EfOM degradation occurred through the
276	elimination of aromatic structures that usually exhibit fluorescence signal at those wavelengths
277	(Chen et al., 2003). The third extraction was performed with 76 L of wastewater effluent (WW3,
278	4.97 mg C/L) after 72 hr of chloramination (14.9 mg Cl <sub>2</sub> /L, pH 7.7) followed by an elution with
279	ethyl acetate. As observed for WW2, the DOC content of WW3 slightly decreased after
280	chloramination (i.e., 4.97 vs 4.60 mg C/L). The total amount of DOC adsorbed onto the two
281	XAD resins (64.6%) was comparable to WW1 and WW2.
282	

Table 2. Percentages of dissolved organic carbon (DOC) isolated on XAD-8 and XAD-4 resins
from wastewater effluent and chloraminated wastewater effluent (72 hr chloramination at 14.2
mg Cl<sub>2</sub>/L and at pH 7.7)

Wastewater effluent sampleDOC before<br/>extraction<br/>(mg C/L)% DOC isolated<br/>on XAD-8 resin% DOC isolated<br/>on XAD-4 resin% DOC not<br/>isolated

WW1 (Non-chloraminated)	4.65	45.8	23.7	30.5
WW2 (Chloraminated)	4.55	34.3	23.1	42.6
WW3 (Chloraminated)	4.60	42.4	22.2	35.4



Fig. 1. Fluorescence EEMs of a) wastewater effluent WW2 and b) wastewater effluent WW2
after 72h chloramination (14.2 mg Cl<sub>2</sub>/L) at pH 7.7

289

#### 290 **2.2. Analytical biology results**

#### 291 **2.2.1. CHO cell cytotoxicity results**

From the data presented in Table 3 and Fig. 2, the XAD-8 EfOM isolate recovered from non-

293 chloraminated wastewater was more cytotoxic than the XAD-4 EfOM extract of the same

294 wastewater. According to the isolation protocol used, hydrophobic EfOM (i.e., XAD-8 isolate,

295 humic substances-like) was more cytotoxic than transphilic EfOM (i.e., XAD-4 isolate). The

296 mixed XAD-8+XAD-4 fraction (50:50 v/v) exhibited an intermediate cytotoxicity, consistent

- with its composition (see Section 1.4) and with the toxicity of the XAD-8 and XAD-4 fractions.
- 298 The structural characterization performed on similar isolates obtained from the effluent collected
- from the same wastewater treatment plant by Zheng et al. (2014) indicated that the XAD-8

300	isolate was more enriched in aromatic structures including phenolic moieties than the XAD-4
301	extract that also incorporated carbohydrates. From the chloraminated wastewater effluent WW2,
302	the XAD-8 extract was also more cytotoxic than the XAD-4 extract, and was significantly more
303	cytotoxic than the non-chloraminated XAD-8 extract (WW1). The XAD-4 extract of WW2 was
304	more cytotoxic than that of WW1 as well. This indicates a significant increase in cytotoxicity of
305	the XAD-8 and XAD-4 extractable EfOM after chloramination. Since the amount of DOC
306	isolated on each resin was lower from the non-chloraminated effluent than from the
307	chloraminated one (i.e., 69.5 and 57.4%, respectively), the cytotoxicity of the chloraminated
308	effluent was even higher when normalized to the total amount of DOC extracted on each resin
309	(Appendix A, Fig. S1). Cytotoxicity results of chloraminated effluent WW3 obtained from ethyl
310	acetate extracts (liquid form) exhibited a greater difference between XAD-8 and XAD-4 EfOM
311	toxicity (i.e., XAD-8 EfOM was 7.6× more cytotoxic). These results are further discussed in
312	paragraph 3.3.4.

Table 3. Comparative CHO cell chronic cytotoxicity of EfOM fractions isolated from wastewatereffluents before (WW1) and after (WW2) chloramination.

Sample	Fraction	Conc. Range (µg/µL)	Chronic Cytotoxicity (LC <sub>50</sub> ) (µg/µL) <sup>a</sup>	r <sup>2 b</sup>	Lowest Significant Toxic Concentration (µg/µL) <sup>c</sup>	ANOVA Test Statistic <sup>d</sup>
WW1	XAD-8	0.05-0.75	0.215	0.99	0.15	$F_{10, 50} = 144.2$ : $P \le 0.001$
WW1	XAD-4	0.05-1.15	0.746	0.99	0.35	$F_{10,55} = 57.6$ : $P \le 0.001$
WW1	XAD-8 + XAD-4	0.05-0.75	0.338	0.99	0.20	$F_{10, 51} = 174.1$ : $P \le 0.001$
WW2 (+NH <sub>2</sub> Cl)	XAD8	.005-0.4	0.060	0.97	0.025	$F_{10, 53} = 81.1: P \le 0.001$
WW2	XAD-4	0.05-0.5	0.124	0.99	0.05	$F_{10, 53} = 105.0: P \le 0.001$

#### $(+NH_2Cl)$

CHO: Chinese Hamster Ovary; EfOM: effluent organic matter; ANOVA: analysis of variance. <sup>a</sup> The LC<sub>50</sub> value is the concentration of the wastewater EfOM sample (as  $\mu g/\mu L$ ), determined from a regression analysis of the data, that induced a cell density of 50% as compared to the concurrent negative controls. <sup>b</sup>  $r^2$  is the coefficient of determination for the regression analysis upon which the LC<sub>50</sub> value was calculated. <sup>c</sup> Lowest cytotoxic concentration was the lowest concentration of the wastewater EfOM sample in the concentration-response curve that induced a statistically significant reduction in cell density as compared to the concurrent negative controls. <sup>d</sup> The degrees of freedom for the between-groups and residual associated with the calculated *F*-test result and the resulting probability value.



315



317 chloraminated wastewater effluent (WW1) and chloraminated wastewater effluent (WW2).

318 XAD-8+XAD-4 fraction from WW1 was obtained by mixing equal volumes of XAD-8 and

319 XAD-4 fractions (50:50 v/v). CHO: Chinese hamster ovary.



322 The results of the genomic DNA damage analyses are presented in Table 4 and the genotoxicity

index (GTI) values are presented in Fig. 3. The GTI is defined as (50%Tail DNA<sup>-1</sup>) $(10^3)$  and the

324 larger the GTI value the greater the genotoxic damage.

325

#### 326

327 Table 4. Comparative CHO cell genotoxicity of EfOM fractions isolated from wastewater

328 effluents before (WW1) and after (WW2) chloramination.

Sample	Fraction	Conc. Range (µg/µL)	Acute Genotoxicity (50% Tail DNA) μg/μL <sup>a</sup>	r <sup>2 b</sup>	Lowest Significant Toxic Concentration (µg/µL) °	ANOVA Test Statistic <sup>d</sup>
WW1	XAD-4	0.3-5.0	4.91	0.98	4.25	$F_{8,23} = 23.0: P \le 0.001$
WW1	XAD-8	0.2-4.0	3.69	0.99	2.5	$F_{9,28} = 49.0: P \le 0.001$
WW2+NH <sub>2</sub> Cl	XAD-4	2.0-3.4	2.45	0.98	2.0	$F_{11, 28} = 153.0: P$ $\leq 0.001$
WW2+NH <sub>2</sub> Cl	XAD-8	0.075- 0.9	0.79	0.96	0.5	$F_{10, 28} = 20.8: P \le 0.001$

CHO: Chinese hamster ovary; EfOM: effluent organic matter; ANOVA: analysis of variance. <sup>a</sup> The SCGE 50% Tail DNA value is the EfOM sample concentration (as  $\mu g/\mu L$ ), determined from a regression analysis of the data, that induced the migration of 50% SCGE Tail DNA value. <sup>b</sup>  $r^2$  is the coefficient of determination for the regression analysis upon which the SCGE 50% Tail DNA value was calculated. <sup>c</sup> The lowest genotoxic concentration was the lowest concentration of the wastewater sample in the concentration-response curve that induced a statistically significant amount of genomic DNA damage as compared to the negative control. <sup>d</sup> The degrees of freedom for the between-groups and residual associated with the calculated *F*-test result and the resulting probability value.



331 Fig. 3. Comparison of the CHO cell genotoxicity index values for each fraction of non-

332 chloraminated wastewater effluent (WW1) and chloraminated wastewater effluent (WW2).

333 CHO: Chinese hamster ovary.

334

335 The genotoxicity data indicate that the wastewater EfOM isolated from the XAD-8 resin was

336 more genotoxic than the XAD-4 extracts of the same wastewaters. This suggests that

337 hydrophobic EfOM comprised more potent inducers of genomic DNA damage than transphilic

338 EfOM. Chloramination had a major effect on the resulting genotoxicity of the XAD-8 extracted

339 organic material, significantly more that for the XAD-4 isolate. These data indicate that the

340 chloraminated EfOM extracted over XAD-8 or XAD-4 resins were more genotoxic than non-

341 chloraminated EfOM.

#### 344 **2.3. DBP formation**

#### 345 **2.3.1. DBP identification**

346 A small volume of the acetonitrile extracts obtained from the two resins that received the

347 chloraminated wastewater effluent WW2 was evaporated under a stream of N<sub>2</sub> and transferred to

348 ethyl acetate for direct full-scan GC-MS analyses in order to characterize the main products

349 formed by chloramination and retained onto XAD resins. Tribromoacetaldehyde (TBAcAl) was

350 the most predominant identified peak in the chromatograms (Appendix A, Fig. S2). TBM,

351 DBAcAm, dibromoacetaldehyde (DBAcAl), chlorodibromoacetaldehyde (CDBAcAl), BIAcAm,

and BCAcAm were also present in the chromatograms. Many other brominated and chlorinated

353 peaks were detected but could not be identified by direct comparison of mass spectra with NIST

database. Brominated species were predominant because of the presence of bromide ion in the

355 wastewater effluent (2.52 mg/L as Br<sup>-</sup>), and some iodinated products (e.g., BIAcAm) were

detected in the XAD-8 fraction (though iodide ion could not be detected by IC analysis because

357 of matrix effects). Some of these brominated and iodinated DBPs (i.e., TBAcAl, DBAcAl,

BIAcAm) are among the most cytotoxic and genotoxic DBPs identified today (Plewa et al.,

359 2008). DBPs and AOX were not quantified during this experiment, and lyophilized fractions

360 were only analyzed for cytoxicity and genotoxicity.

#### 361 **2.3.2. DBP quantification and estimated retention on XAD resins**

362 During the XAD resins extraction conducted on WW3 (initial DOC 4.97 mg C/L, 14.9 mg

363 Cl<sub>2</sub>/L), aqueous samples were collected before and after each XAD resin (Appendix A, Table S1)

- 364 in order to quantify THMs, HAAs, HANs, HAcAms and HAcAls and evaluate the retention of
- 365 each class of DBPs onto both resins (calculated as the difference between DBP concentrations at
- 366 the inlet and at the outlet of a resin column). Results indicated a high retention of total THMs

367	and HANs on XAD-8 resin (94% and 97%, respectively), while HAAs, HAcAls and HAcAms
368	exhibited a poor retention on both resins (Fig. 4). The hydrophobic character of DBP classes are
369	in the order HAcAm < HAA < HAcAl < HAN < THM when estimating logD values at pH 2 for
370	both chlorinated and brominated species using ChemAxon Marvin Suite. The XAD-8 resin is
371	known to adsorb less hydrophilic molecules with higher molecular weights than molecules
372	retained on the XAD-4 resin. This is in agreement with the retention of more hydrophobic DBPs
373	(THMs and HANs) on the XAD-8 resin, possibly because they may exhibit stronger affinity with
374	the hydrophobic macromolecules retained on this resin, rather than because of their direct
375	adsorption on the resin. More hydrophilic compounds were better isolated on the XAD-4 resin
376	(as part of the transphilic/TPI fraction) or not retained by both resins (i.e., hydrophilic/HPI
377	fraction).



Fig. 4. Retention of a) THMs and HANs and b) HAAs, HAcAl and HAcAms onto XAD-8 and
XAD-4 resins used in series. DBP concentrations are expressed as the total sum of each
individual DBP in a given class of DBP. THMs: trihalomethanes; HANs: haloacetonitriles;
HAAs: haloacetic acids; HAcAls: haloacetaldehydes; HAcAms: haloacetamides; DBP:
disinfection byproduct.

**2.3.3. DBP contributions to AOX** 

387	AOBr and AOCl concentrations in chloraminated wastewater effluent were 1658.2 $\mu$ g/L as Br
388	and 1078.9 $\mu$ g/L as Cl, which is in accordance with the presence of a majority of brominated
389	DBPs detected in full-scan chromatograms. The proportions of AOX (AOCl and AOBr) retained
390	on each resin were calculated in the same manner than individual DBPs (i.e., by difference
391	between inlet and outlet concentrations). AOBr and AOCl were mostly retained on the XAD-8
392	resin (748.7 $\mu$ g/L as Br and 710.8 $\mu$ g/L as Cl), compared to the XAD-4 resin (563.8 $\mu$ g/L as Br
393	and 209.6 $\mu$ g/L as Cl), which may explain the higher toxicity observed in the isolated
394	XAD-8/hydrophobic (HPO) fraction than that of the XAD-4/transphilic (TPI) fraction (Fig. 5).
395	Overall, 85.3% of AOCl and 79.2% of AOBr were retained on both resins. All analyzed DBP
396	concentrations were expressed in $\mu$ g/L as Cl (Cl-DBP) or $\mu$ g/L as Br (Br-DBP) to obtain their
397	relative contribution to AOCl and AOBr concentrations (Fig. 5). Unknown AOCl (UAOCl) and
398	unknown AOBr (UAOBr) were determined from the differences between measured AOCl or
399	AOBr and the sum of chlorine or bromine-equivalent concentrations of measured specific DBPs.
400	Br-DBPs contributed to 23.1% of AOBr retained on the XAD-8 resin and to 9.2% of AOBr
401	retained on the XAD-4 resin. Bromoform, DBAN and DBCM were the three major Br-DBPs
402	contributing to AOBr retained on the XAD-8 resin, while DBAA, CDBAcAl and DBAcAm were
403	the major Br-DBPs retained on the XAD-4 resin. Br-DBPs present in the fraction not retained
404	onto the two resins exhibited concentrations of the same order of magnitude of Br-DBPs retained
405	onto the XAD-4 resin. Cl-DBPs only represented 2.2% and 4.5% of AOCl retained on XAD-8
406	and XAD-4 resins, respectively, which indicates that a majority of chlorine-containing
407	compounds were not quantified.
408	





- on concentrations analyzed in water samples at the inlet and outlet of each resin. UAOCl and
- UAOBr stand for unknown AOCl and unknown AOBr.

#### 2.3.4. DBP quantification in XAD resin-isolated fractions and associated toxicity

417 Ethyl acetate was used as the eluting solvent of the organics adsorbed onto the two resins from 418 the chloraminated wastewater effluent WW3, in order to directly quantify DBPs present in each 419 recovered fraction. The cytotoxicity of the two fractions was also assessed and expressed as  $LC_{50}$ 420 concentration factor as discussed previously (Appendix A, Table S2). Cytotoxicity index (CTI) 421 value derived from LC<sub>50</sub> concentration factor of the XAD-8 fraction was 222, i.e.,  $\sim$ 20-fold more 422 toxic than values reported from surface waters and secondary effluents subjected to various 423 oxidation processes (i.e., ozone, chlorine, UV or chlorine dioxide) and CTI value of XAD-4 424 fraction (CTI = 29) was in the same range than chlorinated secondary effluents (Dong et al., 425 2016; Jeong et al., 2012; Plewa et al., 2012). Small volumes of ethyl acetate extracts were diluted 426 in ethyl acetate (for HAcAms analyses) or in MTBE (for THMs and HAAs analyses) and spiked 427 with internal standards before injection in GC-MS. After correction with concentration factor of 428 extraction and evaporation steps (concentrations  $\times$  760), concentrations obtained by analysis of 429 the ethyl acetate extracts (Appendix A, Table S3) were in the same range as concentrations 430 calculated by the difference between inlet and outlet of the columns (i.e.,  $\sim$ 72% and 99% 431 recovery for XAD-8 and XAD-4 resins, respectively, losses possibly occurring during elution 432 and subsequent preparation of samples for analysis). More DBPs and AOX were quantified in 433 the chloraminated effluent isolated from the XAD-8 resin as compared to the XAD-4 resin. 434 Some DBPs (e.g., THMs) that were detected at high concentrations exert much lower toxicity 435 than others such as HANs. In order to compare DBP results with the measured cytotoxicity and 436 genotoxicity of each recovered fraction, the total DBP additive cytotoxicity (AC) was calculated 437 from the measured concentration of each individual DBP divided by its respective cytotoxicity 438 LC<sub>50</sub> value available in the literature (Eq. 1) (Allard et al., 2015; Smith et al., 2010).

439 
$$AC = \sum_{i}^{n} \frac{C_{i}}{LC_{50,i}}$$
 (1)

440	Where C <sub>i</sub> is the measured	concentration of a	i given DBP <i>i</i> i	n a mixture of <i>i</i>	i DBP, LC <sub>50,i</sub> is its
	1		0		, , , , , , , , , , , , , , , , , , , ,

441 LC<sub>50</sub> cytotoxicity value and the calculated AC is a dimensionless value. Following this approach,

442 the ratio between XAD-8 and XAD-4 resin calculated additive cytotoxicity was similar to the

- 443 ratio of measured cytotoxicity index values (Fig. 6). While THMs were the major DBPs present
- 444 in the XAD-8 fraction, most of the calculated additive cytotoxicity was attributed to HANs and
- 445 HAcAms. DBAN, BAcAm, BCAN and DBAcAm were the dominant cytotoxic DBPs,
- 446 accounting for 68.7%, 14.0%, 8.2% and 6.7% of the calculated AC, respectively. In the XAD-4
- 447 fraction, HAcAms and HAAs accounted for the majority of the calculated additive cytotoxicity,
- 448 with DBAcAm, BCAcAm and MBAA accounting for 64.8%, 16.5% and 13.4%, respectively.
- 449 Overall, N-DBPs (i.e., HANs and HAcAms) were responsible for the majority of the calculated
- 450 cytotoxicity in both fractions (99.4% and 78.5% for XAD-8 and XAD-4 fractions, respectively).



453 Fig. 6. DBP additive cytotoxicity (AC) calculated from measured concentrations of DBPs in the
454 XAD-8 and XAD-4 resin fractions and individual DBP cytotoxicity LC<sub>50</sub> values, compared to
455 the measured cytotoxicity index values of the two fractions.



458 isolated fraction according to a concentration addition model based on the individual cytotoxicity

- 459 LC<sub>50</sub> value of each analyzed DBPs and their respective measured molar fraction (calculated from
- 460 the total measured DBP concentration) (Eq. 2) (Tang et al., 2013; Zeng et al., 2016).

461 CCTI = 
$$\sum_{i=1}^{n} \frac{p_i}{LC_{50, i}}$$
 (2)

Where  $p_i$  is the molar fraction of DBP *i* in a mixture of *n* DBP and LC<sub>50,i</sub> is the individual LC<sub>50</sub> value of DBP *i*. The obtained CCTI values are expressed as  $(LC_{50})^{-1}$ , which facilitates the comparison with measured cytotoxicity index values (expressed as  $(LC_{50})^{-1}$ ).

466 Following this approach, the CCTI value of the XAD-8 fraction was still higher than that of the 467 XAD-4 fraction, in accordance with measured cytotoxicity index values, but the ratio between 468 the calculated cytotoxicity of XAD-8 and XAD-4 isolates was much lower than from the first 469 approach (Appendix A, Fig. S3), because this calculation is based on a number n of measured 470 DBPs, while many other unknown Cl-DBPs and Br-DBPs were measured in the XAD-8 fraction 471 as UAOCl and UAOBr. This result shows that analyzed DBPs did not significantly represent the global cytotoxicity of the samples, i.e., that other toxic products (e.g., chlorinated, brominated or 472 473 even iodinated byproducts) accounted for the difference observed between the XAD-8 and the 474 XAD-4 fraction. In order to correct the CCTI values to take into account the unknown 475 halogenated products, molar fractions of DBPs were calculated based on total concentrations of 476 measured AOX (expressed as mol/L as X) (Eq. 3).

477 
$$CCTI = \sum_{i=1}^{n} \frac{\frac{C_i}{AOX}}{LC_{50, i}}$$
(3)

478 Where Ci is the measured concentration (expressed as mol/L as X) of a given DBP *i* and LC<sub>50,i</sub> is 479 its LC<sub>50</sub> cytotoxicity value.

480 With this approach, the proportion of N-DBPs (HANs and HAcAms) in AOX was a good proxy

481 of the global toxicity measured in the samples (Fig. 7).



Calculated DBP Cytotoxicity Index (CCTI)  $(LC_{50})^{-1}$  (10<sup>3</sup>)



Fig. 7. DBP additive cytotoxicity index (CCTI) values calculated from molar fractions of each
DBP (as fractions of measured AOX), divided by their respective individual cytotoxicity LC<sub>50</sub>
values, compared to the measured cytotoxicity index values of the XAD-8 and XAD-4 fractions.
AOX: adsorbable organic halide.

#### 490 **3. Conclusions**

491 - The toxicity of chloraminated wastewater effluents was significantly higher than the

492 toxicity of raw wastewater effluents

493	-	More DBPs were isolated from the XAD-8 resin, i.e., hydrophobic (HPO) EfOM fraction,
494		which is in accordance with the higher toxicity observed for this fraction, when compared
495		to the results obtained with the XAD-4 resin.
496	-	Isolation recovery of DBPs onto XAD resins depends on their hydrophilicity: most
497		hydrophobic THMs and HANs were mostly retained on the XAD-8 resin (also showing a
498		larger proportion of DOC adsorbed), while more hydrophilic HAAs, HAcAms and HAcAls
499		were less retained on both resins, leading to higher concentrations in the organic fraction
500		that did not adsorb onto the two superposed resins. The toxicity of the non-retained organic
501		fraction needs to be evaluated.
502	-	Measured DBPs accounted for minor proportions of the brominated and chlorinated
503		products analyzed as AOBr and AOCl, which means that most AOX consisted in unknown
504		brominated and chlorinated compounds.
505	-	Estimated cytotoxicity of XAD-8 resin/hydrophobic (HPO) and XAD-4 resin/transphilic
506		(TPI) isolates calculated based on additive cytotoxicity of measured DBPs did not account
507		for the difference observed between the measured cytotoxicity of the two fractions. Other
508		products were probably responsible for the higher toxicity of the HPO fraction. These
509		products may include chlorinated, brominated and iodinated byproducts, or non-
510		halogenated byproducts (e.g., nitrosamines) and unknown high-molecular weight
511		compounds.
512	-	Among the measured DBPs, N-DBPs (i.e., HANs and HAcAms) contributed to the
513		majority of the estimated additive DBP cytotoxicity of each fraction (78.5% and 99.4% for

- 514 TPI and HPO fractions, respectively), and were a good proxy of the global cytotoxicity
- 515 when expressed as molar fractions of AOX.

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#### 521 Appendix A. Supplementary Material

522 Supplementary data to this article can be found online at 523 http://dx.doi.org/10.1016/j.jes.2017.04.022.

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