Chlorination and ozonation differentially reduced the microcystin content and tumour promoting activity of a complex cyanobacterial extract

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ABSTRACT

Despite intensive research and management efforts in the past decades, cyanobacterial blooms and their toxins, such as microcystins (MCs), continue to represent a major ecological and health problem in fresh waters throughout the world. Our objective was to compare the efficacy of two commonly used drinking water treatment technologies, chlorination and ozonation, in removing MCs and in reducing tumour promotion-related effects of cyanobacteria, such as inhibition of gap junctional intercellular communication (GJIC) and activation of mitogen activated protein kinases (MAPKs) in a rat liver epithelial stem-like cell line (WB-F344). This combined chemical and bioassay approach demonstrated that ozone effectively removed all MCs from an extract of a globally important bloom-forming cyanobacterium, Microcystis sp. Ozone also significantly reduced the overall tumour promotional potency of the cyanobacterial extract, as indicated by a substantial reduction in the ability of the extract to inhibit GJIC and activate extracellular receptor kinase 1/2 (ERK1/2). Although comparable reduction of total organic carbon was achieved by ozone and chlorine treatment, chlorination was much less effective in removing MCs and reducing the effects on GJIC. Chlorination had a biphasic effect with an observed decrease of extract-induced activation of ERK1/2 at the lower chlorine doses; whereas at high doses of chlorine the by-products of chlorination actually induced the activation of ERK1/2. The extracts induced p38 activation, and chlorination was not effective in reversing this effect, while ozone did reverse this effect, albeit not as much as the activation of ERK1/2. Thus, ozone was effective in reducing the toxicity of cyanobacterial extracts while chlorination was not only lacking efficacy, but at high doses of chlorination further produced by-products that were equally toxic as the untreated samples. Our study indicates the value of using an effect-based approach to assess the efficacy of water treatment systems in removing toxins, and more specifically demonstrates that ozone was more effective at reducing the toxic potential of cyanobacterial-contaminated water.

Key words: Cyanobacteria; water treatment; toxicity; chlorination; ozonation; microcystin.

Received: 11 October 2016. Accepted: 14 February 2017.

INTRODUCTION

Despite intensive research identifying causes, consequences and possible preventive measures of cyanobacterial mass proliferations, cyanobacterial blooms represent a major problem in fresh waters throughout the world. The adverse consequences include water quality degradation, accumulation and microbial decay of bloom biomass followed by lowering of oxygen content in water (Wiegand and Pflugmacher, 2005). In addition, a wide spectrum of toxins and secondary metabolites produced by cyanobacteria have been shown to adversely affect aquatic organisms (Codd *et al.*, 2005; Zanchett and Oliveira-Filho, 2013), livestock (McGorum *et al.*, 2015) and human health (Kuiper-Goodman *et al.*, 1999; Zanchett and Oliveira-Filho, 2013).

While different cyanobacterial taxons can contribute to the formation of dense cyanobacterial water blooms depending on geographical and ecological conditions, Microcystis sp. represents cosmopolitan and pervasive cyanobacterial genera, which is frequently reported to dominate water blooms in freshwaters of all continents except Antarctica (Harke et al., 2016). Microcystis species (such as Microcystis aeruginosa) are also among the prominent producers of the most broadly studied cyanobacterial toxins microcystins (MCs) (Bláha et al., 2009). MCs are known liver tumour promoters (Nishiwaki-Matsushima et al., 1992), with the most common structural variant microcystin-LR (MC-LR) being classified by the International Agency for Research on Cancer as a possible human carcinogen (IARC2B) (Grosse et al., 2006). MCs have been shown to act via inhibition of serine/threonine protein phos-



phatases (PPs) (Campos and Vasconcelos, 2010) and induction of oxidative stress followed by damage to cellular macromolecules (Mathe *et al.*, 2016). Chronic exposures to cyanobacteria and their toxins – *e.g.*, via contaminated drinking water – have been associated with increased occurrence of liver and colorectal cancer (Yu, 1995; Zhou *et al.*, 2002; Svircev *et al.*, 2009). Other (often yet unidentified) compounds produced by complex cyanobacterial blooms can also induce or contribute to different adverse effects (Oberemm *et al.*, 1997; Berry *et al.*, 2009) including tumour promotion (Bláha *et al.*, 2010).

Occurrence of toxic cyanobacteria in surface water presents a challenge to drinking water treatment facilities. Preventive measures as well as various water treatment technologies used to minimize human health risks caused by cyanobacteria and their toxins have been recently summarized and critically discussed (Westrick et al., 2010; Merel et al., 2013; Roegner et al., 2014; Hiskia et al., 2016; Ibelings et al., 2016). The available approaches able to remove MCs to different extents include coagulationflocculation-sedimentation, standard oxidation and disinfection by chlorine or permanganate, ozonation and UV disinfection, sorption by activated carbon, nano- and ultrafiltration, and advanced oxidation processes (AOP) (Westrick et al., 2010; Merel et al., 2013; Roegner et al., 2014). Different drinking water treatment technologies are applied in different countries and contexts, and evaluation of their treatment efficiency should focus on i) removal of targeted priority pollutants (e.g., MCs in case of cyanobacterial toxins) to comply with current treatment goals; as well as ii) removal of other potentially harmful and toxic components of the complex material which may not be fully chemically characterized; and iii) formation of new harmful metabolites/toxic by-products during the application of the water treatment technology (Upham et al., 1994; Upham et al., 1995; Prasse et al., 2015).

To address these different aspects of drinking water treatment, complementary chemical and biological tools (*i.e.* instrumental analyses and bioassays) should be included in the monitoring plans (Maier *et al.*, 2015). This ongoing effort can be highlighted by current implementation of the effect-based tools into the monitoring guidelines for water quality assessment (Wernersson *et al.*, 2014), which combines several bioassays targeting different toxic modes of action (MoAs) to provide additional information to classical chemical analyses and thus a more integrative view.

Gap junctional intercellular communication (GJIC) plays a fundamental role in maintaining tissue homeostasis, and provides an excellent biological endpoint to assess potential adverse health effects of many anthropogenic toxicants and natural toxins (Vinken *et al.*, 2009). GJIC is a critical cellular process for the coordination of different intra-, extra-, and inter-cellular signalling pathways re-

quired for proper cell behaviour, tissue development, tissue function and maintenance of tissue homeostasis. Most chemical carcinogens and tumour promoters inhibit GJIC in in vitro assays, and demonstrated to be a representative marker of tumour promoting potency (Rosenkranz et al., 1997). Recently cyanobacterial extracts and exudates were determined to be potent in vitro inhibitors of GJIC (Bláha et al., 2010; Novakova et al., 2011). Inhibition of GJIC by these extracts were independent of the well-recognized tumour promoting cyanotoxins, MCs or cylindrospermopsin, indicating the existence of not-yet-identified toxic compounds (Novakova et al., 2013). Aquatic contaminants, such as polycyclic aromatic hydrocarbons (PAHs) or polychlorinated biphenyls synergistically potentiated the inhibitory effects of cyanobacterial extracts on GJIC (Novakova et al., 2012), which further highlights the need to bioassay these mixtures for adverse effects in the complex assessment of drinking water quality, and efficacy of treatment technologies.

In fact, in vitro assessment of GJIC has been successfully used along with chemical analysis as a principal bioassay to study tumour promoting activity of water chlorination by-products (Hakulinen et al., 2004; Nishikawa et al., 2006). Similarly, GJIC assay has been applied to evaluate the efficiency of removing different anthropogenic contaminants and their toxic by-products with ozone, such as polycyclic aromatic hydrocarbons (Upham et al., 1994; Upham et al., 1995; Herner et al., 2001; Luster-Teasley et al., 2002; Luster-Teasley et al., 2005) and various pesticides (Upham et al., 1997; Masten et al., 2001; Wu et al., 2007). The in vitro bioassays used in these studies to assess GJIC were based on the scrapeload dye transfer (SL-DT) technique (El-Fouly et al., 1987). This SL-DT assay provides fast (minutes of exposure) and integrative responses, which reflect dysregulations of different cell processes and multiple signalling pathways controlling GJIC (Upham et al., 2016). Dysregulation of GJIC is an epigenetic, phenotypic marker for determining tumour promotional activity, which contrasts and compliments the more commonly used genotoxic and specific nuclear receptor transactivation assays (Leusch and Snyder, 2015). GJIC can be assessed in vitro using diverse non-tumorigenic cells, with a rat liver epithelial cell line, WB-F344, being one of the most widely used GJIC model for the assessment of tumour promoting activity, as well as determining chemopreventive effects of chemicals (Upham et al., 1998; Sovadinova et al., 2015; Babica et al., 2016b). To further validate tumorigenic activity, compounds that dysregulate GJIC are often tested for effects on signal transduction pathways implicated in neoplastic transformation, such as mitogen-activated protein kinases (MAPK-ERK1/2 and MAPK-p38) (Upham et al., 2008; Osgood et al., 2014; Babica et al., 2016a).

Our objective was to evaluate and compare the effi-

cacy of two broadly used drinking water treatment oxidation technologies, namely chlorination and ozonation, on the removal of known cyanotoxin MC concentrations as well as on changes in biological effects that are independent of the MC content (*i.e.*, removal of overall cytotoxicity and tumour promotional potency). We used a natural bloom sample that was dominated by the cosmopolitan and environmentally relevant bloom-forming cyanobacteria *M. aeruginosa*. Extracts were prepared and characterized for content of MC, total organic carbon (TOC) concentration, initial cytotoxicity, effect on GJIC and modulation of signalling kinases (MAPKs). These cyanobacterial extracts were treated by chlorine or ozone, and evaluated for the changes in the toxin content, TOC, and *in vitro* cytotoxicity and tumour promoting activity.

METHODS

Cyanobacterial sample

The sample of toxic cyanobacterial water bloom was collected from a lake located within the campus of Michigan State University (East Lansing, MI, USA; 42°40'50.09"N, 84°29'14.27"W) in September 2008 using a 20 µm plankton net. The bloom was dominated by Microcystis species: M. aeruginosa (>50% of the cell counts) accompanied by M. flos-aquae (~20%) and M. ichthyoblabe (~20%). The biomass was freeze-dried and 38 g of dry weight (DW) was extracted by 20 min sonication (Fisher Sonic Dismembrator Model 300; Fisher Scientific, Pittsburgh, PA, USA) while stirring on ice with 566 mL of 50% methanol (i.e., 66.7 g DW L⁻¹ equivalent). The samples were centrifuged at $31,000 \times g$ and the supernatant fraction was collected and dried using a vacuum evaporator. The dry extract was dissolved in 47.2 mL of MilliQ water (MilliQ Synthesis A10; Millipore, Billerica, MA, USA) to obtain the final concentrated extract corresponding to 800 g DW of original biomass per one litter of water.

Chlorination and ozonation

The extract was aliquoted into 4 mL fractions to be treated by chlorination or ozonation as summarized in Tab. 1. Chlorination was carried out with sodium hypochlorite (NaOCl) in a phosphate buffer (0.5 M K₂HPO₄-0.293 M NaOH, pH 7.0±0.2) according to the Method 5710C (Clesceri *et al.*, 1998). The sample was treated for 30 min to 24 h with NaOCl at different concentrations of free chlorine (7 to 1000 mg L⁻¹) corresponding to contact time (CT) values 0.21×10^3 , 7×10^3 , 50×10^3 and 1440×10^3 mg min L⁻¹. The oxidation reaction was stopped by addition of 10% w/v of NaHSO₃. Vehicle controls were prepared from MilliQ water equally treated with chlorine and quenched. No residual chlorine was present in the vehicle controls after the quenching. To increase the weight ratio of chlorine to dry

weight of the extract (or TOC or MC concentration), the concentration of the original biomass was also diluted fourand eight times (*i.e.*, " $\frac{1}{4}$ " or " $\frac{1}{8}$ ") before the chlorination step with 500 mg L⁻¹ of chlorine for 100 min.

The original extract dissolved in MilliQ water was ozonated for 30 min using a commercial ozone generator (OZ2PCS-V; Ozotech, Yreka, CA, USA) at the concentration of O_3 (gas)=5.0 mg L⁻¹ with a gas flow rate of one L min⁻¹ and temperature of 20.0±0.5°C to maximize ozone dissolution.

Microcystin analysis

The concentrations of MCs in the original biomass as well as in the samples after the chlorination and ozonation treatment were analysed by HPLC-UV/DAD following the procedure described earlier (Babica *et al.*, 2006). Toxins were identified based on their retention times and characteristic UV absorption spectra and quantified using the calibration curves of standards of MC-RR, -YR, -LR, -LW, -LF, and nodularin. An example of the HPLC chromatograph of the cyanobacterial extract recorded at 238 nm is in Supplementary Fig. 1.

TOC analysis

TOC of the extract before and after chlorination and ozonation was determined using LiquiTOC analyzer (Elementar Analysensysteme, Hanau, Germany) where measurements were made by high temperature oxidation of the carbon (850-900°C) and detection of CO_2 by an NDIR photometer.

Bioassays

WB-F344 rat liver non-tumorigenic stem-like cells (Tsao et al., 1984) were cultured in D-media (Kao et al., 1997) with 5% v/v of fetal bovine serum (Gibco, Life Technologies, Grand Island, NY, USA) at 37°C and 5% CO₂. Cells were cultured to full confluence for 48 h in 35 mm tissue culture dishes (Costar; Cambridge, MA, USA). These confluent cells were used for the various time and dose related experiments. The vehicle controls were water chlorinated or ozonated using the same conditions as applied for studied samples. A sample or vehicle was added directly to the cell culture medium in the dish and gently mixed. Non-treated cells were used as negative controls. The final concentrations of extracts in the bioassays were expressed as the original weight of dry biomass used for extract preparation per unit volume (g DW L⁻¹). Viability/cytotoxicity was tested after 30-min and 24-h exposures using the neutral red assay as reported before (Babica et al., 2016b). The method determines viable cells capable of neutral red inclusion into lysosomes (Borenfreund and Puerner, 1985). Viability was expressed as the fraction of negative (non-treated) control (FOC).

Abbreviation	Treatment type and CT value (mg min L ⁻¹)	Dose of Cl or O ₃ (mg L ⁻¹)	Treatment duration	Total MC (µg g ⁻¹ DW)°	MC-LR (µg g ¹ DW)°	TOC 2 (µg g ⁻¹ DW)°	TOC 24-h Cell viability (ug g ⁻¹ DW)° 1C ₅₀ (95% CI) LOEC 1 (g DW L ⁻¹)	y 30-min GJIC IC ₅₆ (95%CI) LOEC (g DW L ⁻¹)	30-min ERK1/2 LOEC (g DW L ⁻¹)	30-min p38 LOEC (g DW L ⁻¹)
IN	no treatment	ł	56	517 (0.52-12.41) [§]	411 $(0.41-9.86)^{\$}$	104,761 (105-2514) [§]	$11.6^{\#}$ (10.2-13.1) 12	$7.8^{\#}$ (7.0 to 8.7) 8	8	4
Cl(7-30)	Chlorination 0.21×10 ³	٢	30 min	559 (0.56-13.42)	443 (0.44-10.63)	114,761 (114-2739)	14.8 (12.6-17.3) <i>12</i>	10.0 (9.3-10.7) 8	×	4
CI(70-100)	Chlorination 7×10 ³	70	100 min	447 (0.45-10.74)	352 (0.35-8.44)	88,355 (88-2121)	14.3 (12.2-16.5) <i>12</i>	10.7***^ (9.7-11.8) 8	16	4
CI(500-100)	Chlorination 50×10 ³	500	100 min	467 (0.47-11.20)	371 (0.37-8.91)	86,895 (87-2085)	9.1 (6.9-12.2) <i>12</i>	12.6^{***} (11.2-14.1) 8	16	4
¹ /4 Cl(500-100) ^s	Chlorination 50×10 ³	500	100 min	346 (0.35-8.31)	275 (0.28-6.60)	77,229 (77-1853)	10.4 (7.1-15.4) 12	9.6 (8.4-11.0) <i>8</i>	×	4
⅓ Cl(500-100)⁵	Chlorination 50×10 ³	500	100 min	412 (0.41-9.88)	331 (0.33-7.94)	81,193 (81-1949)	7.0 (7.0-15.5) 12	8.1 (6.7-9.7) <i>8</i>	n.a.	n.a.
Cl(1000-24h)	Chlorination 1440×10 ³	1000	24 h	367 (0.37-8.81)	295 (0.29-7.07)	78,754 (79-1890)	$ \begin{array}{c} 12.3 \\ (9.7-15.7) \\ 12 \end{array} $	12.7^{***} (10.7-15.0) 12	16	4
03	Ozonation	5 °°	30 min	n.d.	n.d.	70,622 (71-1695)	>24 >24	>24 24	>16	4
CT values: the concentration of free chlorine multiplied by t biomass before methanol extraction. # Plain text represents 1 actual MC and TOC concentrations (mg L ⁻¹) in the experime P<0.05). *Before chlorination, the concentration of original 1 an individual MC variant=1.3 µg g ⁻¹ DW; n.a., not analysed.	ntration of free chlon anol extraction. #Plε concentrations (mg 1 orination, the concen iant=1.3 μg g ⁻¹ DW;	rine multiplied by the ain text represents IC L^{-1}) in the experiment atration of original bit ; n.a., not analysed.	CT values: the concentration of free chlorine multiplied by the contact time with the sample being disinfected; it is expressed in units of mg min L^{-1} . ^o The concentration is calculated per g DW of original biomass before methanol extraction. [#] Plain text represents IC ₃₀ values with 95% confidence intervals in parentheses, italicized numbers show LOEC values. [§] Values in parentheses indicate the range of actual MC and TOC concentrations (mg L^{-1}) in the experiments (calculated for the range of tested concentrations of 1-24 g DW L^{-1}). [°] Significantly different than IC ₃₀ value of NT extract (Student's <i>t</i> -test, P<0.05). [§] Before chlorination, the concentration of original biomass was diluted four (¹ / ₄) or eight times (¹ / ₈). ^{oo} Gas flow rate of 1 L min ⁻¹ for 30 min; n.d., not detected: minimal detection limit (MDL) for an individual MC variant=1.3 µg g ⁻¹ DW; n.a., not analysed.	ample being disin fidence intervals i nge of tested conce (\mathcal{V}_{4}) or eight times	fected; it is express in parentheses, ital antrations of 1-24 § ()/s). °°Gas flow ra	sed in units of n icized numbers g DW L ⁻¹). 'Sig te of 1 L min ⁻¹	ig min L ⁻¹ . ^o The show LOEC val iffcantly differe for 30 min; n.d.,	concentration is c ues. [§] Values in pa nt than IC ₅₀ value not detected: min	alculated per g I rentheses indica of NT extract (S imal detection li	W of original e the range of tudent's <i>t</i> -test, mit (MDL) for

Tab. 1. Conditions of chlorination and ozonation of the studied cyanobacterial extract.

Tumour promotion assay based on GJIC assessment used only the concentrations that were determined to be noncytotoxic using the neutral red assay. Cells were exposed for 30 min to the tested samples or corresponding vehicle. Treatment with 1-methylanthracene (70 µM, 30 min) was used as a positive control inducing complete inhibition of GJIC. GJIC was assessed using modified SL-DT technique (El-Fouly et al., 1987; Babica et al., 2016c; Upham et al., 2016). The migration of the dye through gap junctions was visualized with a Nikon Eclipse TE3000 phase contrast/fluorescent microscope and the images digitally captured with Nikon EZ Cool Snap CCD camera (Nikon Instruments, Melville, NY, USA), where three representative images were acquired from each dish. The area of dye transfer was measured for each image using ImageJ (https://imagej.nih.gov/ij/). The measured areas were adjusted by subtracting an area of the dye transfer in the positive control with completely inhibited GJIC. Adjusted areas from each image were compared with an averaged adjusted area of the negative control and expressed as FOC.

Activation of regulatory kinases MAPK ERK1/2 and p38 after a 30 min exposure to the sample was determined by Western blotting. Western blot analyses were done as reported previously (Babica et al., 2016a). Briefly, the proteins were extracted with 20% SDS solution containing inhibitors of proteases and phosphatases, and the protein concentration of the cell lysates was determined with DC assay kit (Bio-Rad, Hercules, CA, USA). The proteins (20 µg per lane) were separated on 12.5% SDS-PAGE (Laemmli, 1970) and then electrophoretically transferred to a 0.45 µm PVDF membrane (Millipore). To visualize activated, i.e. phosphorylated ERKs and p38, we used rabbit phospho-specific polyclonal antibodies directed to ERK-1 phosphorylated at Thr 202/Tyr204, and ERK-2 phosphorylated at Thr185/Tyr187 (Cell Signaling #9101S, Danvers, MA, USA) and directed to p38 phosporylated at Thr180/Tyr182 (Zymed #36-8500, San Francisco, CA, USA), and secondary donkey anti-rabbit IgG conjugated with horse radish peroxidase (Amersham Bioscience # NA934V, Life Science, Denver, CO, USA). Levels of glyceraldehyde 3-phosphate dehydrogenase (GAPDH) was used as a housekeeping protein, and determined with mouse anti-GAPDH antibodies (Chemicon #MAB374, Millipore) and secondary sheep anti-mouse IgG conjugated with horse radish peroxidase (Amersham Bioscience #NA931V; Life Science). The ERK, p38 and GAPDH protein bands were detected using the ECL SuperSignal West Pico Chemiluminesence detection kit (Pierce, Arlington Heights, IL, USA) and Bio-Rad Image Analyzer.

Data analyses and statistics

At least three independent experiments were done for each treatment (except for Western blot analyses), and the mean \pm standard deviations from independent experi-

ments were calculated. IC_{50} concentrations causing 50% inhibition of the studied effects and their 95% confidence intervals were derived using non-linear regression in GraphPad (GraphPad software Inc., La Jolla, CA, USA). One-way Analysis of Variance (ANOVA) followed by Dunnett's *post-hoc* test was used to identify treatments significantly different from the control. Differences between IC₅₀ values were assessed by Student's *t*-test. P-values less than 0.05 were considered statistically significant.

RESULTS

The original biomass contained 517 μ g of total MC per g of DW with MC-LR being the dominant variant (411 μ g g⁻¹ DW, 79%, Tab. 1). The other variants present included MC-RR (2%), MC-LW (2%), MC-LF (3%) and two other structurally unidentified MC variants (14%). The total MC concentration in the original extract (corresponding to 800 g biomass DW L⁻¹) was 414 mg L⁻¹ (MC-LR concentration: 329 mg L⁻¹). TOC concentration in the non-treated extract of 800 g biomass DW L⁻¹ was 83.81 g L⁻¹, which represents 105 mg of extractable total organic carbon per g of biomass DW.

Ozonation of cyanobacterial samples had direct effects on the concentrations of organic matter in the biomass. As shown in Tab. 1, the 30-min treatment with ozone caused complete degradation of MCs and 33% reduction of TOC compared to non-treated extract. The chlorination with low dose of chlorine (7 mg L^{-1}) for short time (30 min) had no effect on MC and TOC concentrations. The higher doses (70 to 1000 mg L-1) of chlorine and longer treatments (100 min to 24 h) were more effective in reducing MC levels (by 10 to 33%), and this reduction was dependent on chlorine concentrations and treatment times. There was no observed shift in the proportion of individual MC congeners with the MC-LR being the dominant variant. TOC degradation by chlorine (by 16 to 25%) correlated well with MC degradation. The increase of the weight ratio of chlorine to original biomass by diluting nontreated extract before chlorination (Tab. 1, variants 1/4 or 1/8 Cl(500-100)) slightly improved reduction of MC and TOC amount, when the results were adjusted by the corresponding dilution factor and then compared to the nondiluted extract chlorinated under the same conditions (Cl(500-100)).

The original extract (non-treated, NT) showed no significant cytotoxic effects on WB-F344 during 30-min exposure within the range of concentrations tested (up to 24 g DW L⁻¹; negative data not shown). After longer exposure time (24 h), the sample significantly decreased viability of WB-F344 cells (Fig. 1A) with the calculated IC_{50} =11.6 g DW L⁻¹ (Tab. 1). The lowest observed effect concentration (LOEC) inducing statistically significant reduction of the cell viability was 12 g DW L⁻¹ (Fig. 1A).

inhibition of intercellular communication was 8 g DW L^{-1} (containing 4.1 mg L^{-1} MCs) (Fig. 2A). The tumour promotional activity of the NT sample indicated by its effect on GJIC was also confirmed by additional experiments assessing phosphorylation of signalling protein

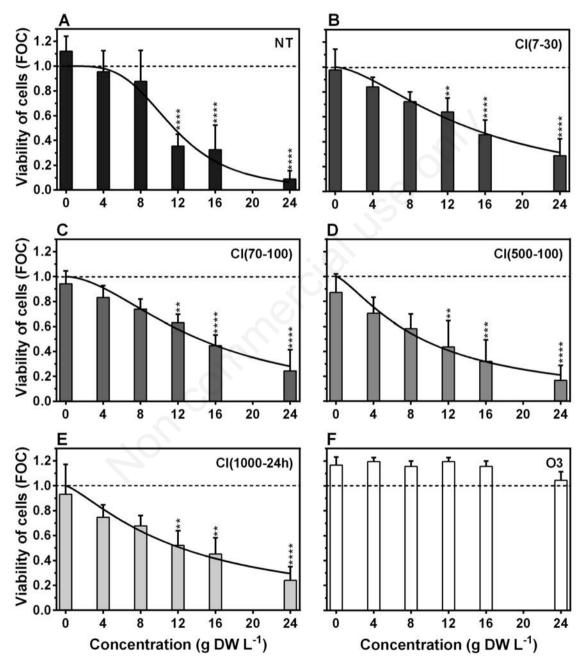


Fig. 1. The effect of chlorination (B-E) and ozonation (F) of the tested cyanobacterial extract on viability in WB-F344 cells after 24-h exposure to 6 different concentrations (4-24 g DW L⁻¹). Data are fractions of controls (FOC) as means \pm standard deviations of independent repetitions of the experiment (n \ge 3). Significant differences from the vehicle control (A) are indicated by asterisks (one-way ANOVA followed by Dunnet's *post-hoc* test; *P \le 0.05; **P \le 0.01; ***P \le 0.001; ***P \le 0.001). NT, original non-treated extract; Cl, chlorinated extracts (in parentheses - free Cl concentrations 7, 70, 500 and 1000 mg L⁻¹; treatment durations 30 min, 100 min, 24 h); O3, ozonated extract (30 min, the flow rate of 1 L min⁻¹ with 5 g O₃ m⁻³).

kinases in WB-F344 cells (MAPK-ERK 1/2 and p38, Fig. 3A). Rapid and clear concentration-dependent activation of both MAPKs was observed after 30-min exposure. The LOEC values for hyperphosphorylation of MAPK- ERK 1/2 and p38 were 8 or 4 g DW L^{-1} of the NT extract, respectively, with maximal effects observed at the highest tested concentration of 16 g DW L^{-1} (Fig. 3A).

Ozonation and chlorination of the original sample had a pronounced effect on the biological activity. The cytotoxic effect was completely eliminated by the ozonation of the extract (Fig. 1F). At the same time, none of the chlorination experimental protocols had any significant effect on cytotoxicity of the original NT extract (Figs. 1B-E), with the estimated IC_{50} values ranging between 9.1 and 14.8 g DW L⁻¹ and not being significantly different (P<0.05) from the IC₅₀ value for the NT extract (Tab. 1). Also, the LOEC values for viability of cells exposed to chlorinated extracts remained at 12 g DW L⁻¹. Interestingly, slight but non-significant increase of IC₅₀ values was associated with lower rather than higher chlorine doses. The increase of weight ratio of chlorine to the original biomass by diluting non-treated extract before chlorination did not decrease the cytotoxicity with the IC_{50} values of 10.4 g DW L⁻¹ for 1/4 Cl(500-100) and 7.0 g DW L^{-1} for $\frac{1}{8}$ Cl(500-100) (Tab. 1, Supplementary Fig. 2). The corresponding vehicle controls for these chlorination conditions caused 20 to 30% significant decrease in WB-F344 cell viability when compared to non-treated control (Supplementary Fig. 2).

The ozone application was also highly potent in eliminating tumour promotional activity of the original NT extract, as shown in Fig. 2F. After 30-min ozonation, the LOEC value for GJIC inhibition was 24 g DW L⁻¹, *i.e.* three times higher than in NT extract. GJIC was reduced only by 30% at 24 g DW L⁻¹ concentration (Fig. 2F), while nearly 100% inhibition of GJIC was observed at 16 to 24 g DW L⁻¹ of NT extract (Fig. 2A).

In contrast, the chlorination treatments had much less pronounced effects on the inhibition of GJIC, with the IC₅₀ values of the chlorinated extracts ranging between 10.0-12.7 g DW L⁻¹. Although these IC₅₀ values were relatively similar to the IC_{50} estimated for NT extract, they were significantly higher except for the lowest free chlorine doses and shortest treatment (P<0.05), and their increase depended on the chlorine dose- and treatment duration (Tab. 1). When the weight-ratio of chlorine to the original biomass was increased by diluting the non-treated extract before chlorination (Supplementary Fig. 3), there was no observable decrease in the dysregulation of GJIC induced as compared to the original NT extract, with the IC_{50} values after chlorination being 8.1 to 9.6 g DW L⁻¹ (Tab. 1, Supplementary Fig. 2). Both ozone and chlorine treatments significantly attenuated the activation of MAPK ERK1/2 induced by NT extract (Fig. 3). Ozonation was more effective, when no activations of the ERK1/2 kinase were observed at concentrations between 4 to 16 g DW L⁻¹. Chlorination also resulted in reduced levels of MAPK ERK1/2 activation (Fig. 3, Tab. 1). The protective effects had biphasic character, when initially increased but then became less apparent with an increase in chlorine dose and chlorination time (Fig. 3, Tab. 1). The effects of both ozone and chlorine treatments on the activation of p38 MAPK were much less pronounced when compared to MAPK ERK1/2, and activation of p38 was still apparent for the ozonated extract as well as most of the chlorinated extracts even at the lowest experimental concentration of 4 g DW L⁻¹ (Fig. 3).

DISCUSSION

Adverse effects of toxic cyanobacteria on human health remain a major issue for both researchers and water managers. Our study confirmed that rapid inhibitions of GJIC and activations of MAPK-ERK1/2 might be a common effect induced by bloom-forming cyanobacteria. The effective concentration for 30-min GJIC inhibition $(IC_{50}=8 \text{ g DW } L^{-1})$ was similar to previously reported values for other extracts from natural blooms dominated by *Microcystis* sp. (IC₅₀=4 or 6 g DW L⁻¹, respectively) (Bláha et al., 2010; Novakova et al., 2011). Modulation of these cellular events by chemicals in vitro are considered to be relevant biomarkers of tumour promoting potency in vivo, as was demonstrated e.g. for tumour promoting phorbol esters like TPA (Madhukar et al., 1996), organochlorine pesticides (Trosko et al., 1987), PCBs (Kang et al., 1996), low molecular weight PAHs (Bláha et al., 2002), clofibrate, phenobarbital, perfluorooctanoic acid, or organic peroxides (Upham et al., 2007; Upham et al., 2009; Vinken et al., 2009). MCs and other cyanotoxins like cylindrospermopsin have been shown to modulate development of tumours (Nishiwaki-Matsushima et al., 1992; Falconer and Humpage, 2005; Svircev et al., 2010; Zegura et al., 2011; de la Cruz et al., 2013). However, neither MC-LR nor cylindrospermopsin had any direct effect on rapid inhibition of GJIC or activation of MAPK ERK 1/2 indicating that other metabolites in cyanobacteria might be responsible for their GJIC-dependent tumour promoting activity (Bláha et al., 2010; Novakova et al., 2011). However, these metabolites have not been identified yet.

In addition, our present study demonstrates, for the first time, a rapid activation of another MAPK-p38 by cyanobacterial environmental extract in the cell line WB-F344, which possesses characteristics of liver progenitor cells (Babica *et al.*, 2016a, b). MAPK-p38 is a critical participant in cellular stress responses and has a key role in inflammation, as well as in tissue homeostasis, by controlling cell proliferation, differentiation, death, survival and the migration of specific cell types (DiDonato *et al.*, 2012). In contrast to MAPK-ERK1/2, which are activated by mitogens or growth factors, MAPK-p38 is activated by environmental and genotoxic stresses including hypoxia, UV, ROS, hyperosmolarity, and heat shock, and its activation

has been linked to protein phosphatase 2 (PP2A) (Nebreda and Porras, 2000; Wagner and Nebreda, 2009). Indeed, MC-LR, a known PP2A inhibitor, has been previously reported to activate MAPKs ERK1/2, p38 or JNK in rodent liver *in vivo* and also in experiments with various cell lines

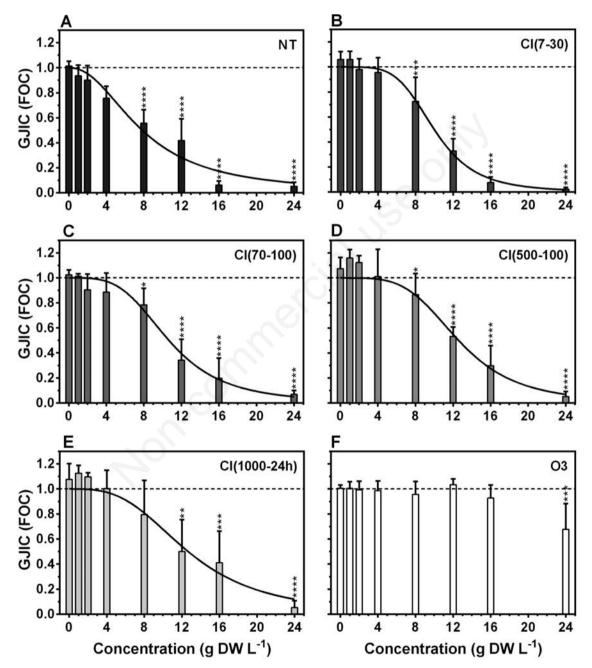


Fig. 2. The effect of chlorination (B-E and ozonation (F) of the tested cyanobacterial extract on gap-junctional intercellular communication (GJIC) in WB-F344 cells after 30-min exposure to 6 different concentrations (1-24 g DW L⁻¹). Data are fractions of controls (FOC) as means \pm standard deviations of independent repetitions of the experiment (n \geq 3). Significant differences from the NT extract (A) are indicated by asterisks (one-way ANOVA followed by Dunnet's post hoc test; *P \leq 0.05; **P \leq 0.01; ***P \leq 0.001; ***P \leq 0.0001). NT, original non-treated extract; Cl, chlorinated extracts (in parentheses - free Cl concentrations 7, 70, 500 and 1000 mg L⁻¹; treatment durations 30 min, 100 min, 24 h); O3, ozonated extract (30 min, the flow rate of 1 L min⁻¹ with 5 g O₃ m⁻³).

in vitro. However, MC-LR effects on MAPKs seem to be dependent on the kinase type, a cell type, and probably also exposure times and concentrations. Depending on the study, MC-LR was found to activate ERK1/2 but not p38 (Dias *et al.*, 2010; Zhang *et al.*, 2013; Adamovsky *et al.*, 2015), p38 but not ERK1/2 (Meng *et al.*, 2011; Lezcano *et al.*, 2012), and both ERK1/2 and p38 (Komatsu *et al.*, 2007; Daily *et al.*, 2010; Sun *et al.*, 2011; Chen *et al.*, 2012; Liu *et al.*, 2016; Wang *et al.*, 2017). The effective concentrations of MC-LR in these studies were typically between 1 and 10 μ M (~1-10 mg L⁻¹).

In the present study, cytotoxicity, GJIC and MAPKs were affected by the cyanobacterial extract diluted to 4-12 g DW L⁻¹ containing MCs with concentrations between 2 and 6 mg L^{-1} , which corresponds to 2 to 6 μ M range and is quite comparable with the other in vitro studies reporting MAPK activation by MCs. However, we previously demonstrated that tumour promoting events, such as rapid GJIC inhibition and MAPK ERK1/2 activation in rat liver progenitor cells, are induced by other cyanobacterial metabolites but not MC-LR or cylindrospermopsin (Bláha et al., 2010). Our results suggest that MAPK p38 can also be activated by transformation or degradation products of MCs, other compounds of cyanobacterial origin and/or their transformation or degradation products, since p38 was activated not only by MCcontaining non-treated cyanobacterial extract, but also by ozonated extract without detectable levels of MCs. These findings suggest that progenitor cells, in comparison with differentiated hepatocytes, might be less prone to the effects of MCs, possibly due to limited expression of key proteins involved in MCs uptake and metabolism, such as organic-anion transporting polypeptides (OATPs). Nevertheless, progenitor cells can be apparently a target of other compounds present in cyanobacterial biomass, which are capable to induce toxic and tumour promoting effects in this specific population of liver cells (Bláha et al., 2010) known to play a critical role in the maintenance of liver tissue homeostasis, liver regeneration and hepatocarcinogenesis (Canovas-Jorda et al., 2014).

The evidence supporting the existence of other components of cyanobacterial biomass contributing to the tumour promoting and toxic effects of complex cyanobacterial samples emphasize the need for effectbased evaluation of the efficacy of water treatment technologies in addition to chemical analyses. Different physicochemical purification processes employed in DWTPs may have different efficacies in removing the target contaminant, such as well recognized toxicants (such as MCs) *vs.* elimination of the overall toxicity. Our results demonstrate that ozone effectively and rapidly removes the MC fraction of the complex cyanobacterial samples. Although the highest chlorine dose resulted in a decrease of TOC that was comparable to the ozone treatment, reactive chlorine was not as effective compared to ozone in removal of MCs. Similarly, removal of cytotoxicity and overall epigenetic toxicity of the studied sample by ozonation appeared to be much more effective than chlorination, although ozone had a less pronounced effect in decreasing p38 activation as compared to ERK1/2 and GJIC.

With regard to chlorine the literature demonstrate that its application removes MCs but the efficiency of removal decreases with increases in pH and dissolved organic material along with formation of less effective oxidant ClO⁻ (Merel *et al.*, 2010). Several studies agreed that 0.5 mg of residual chlorine per liter should efficiently remove pure MCs in distilled water during 5-30 min (depending on MC concentration) at pH lower than 8 (Nicholson *et al.*, 1994; Newcombe and Nicholson, 2004; Acero *et al.*, 2005). Similar scenarios are also expected at DWTP at environmentally relevant concentrations of MCs (Merel *et al.*, 2009).

Several previous studies using PP-inhibition as a biomarker of toxicity as well as other bioassays reported a decrease in MC concentration and toxicity after chlorination of cyanobacterial samples (Nicholson et al., 1994; Tsuji et al., 1997; Rodriguez et al., 2008; Merel et al., 2010). For example, the chlorination of *M. aeruginosa* extract (16 mg DW L^{-1}) with the dose of 1 mg L^{-1} chlorine and contact time of 30 min (CT value=30) effectively removed 95% of MCs (initial concentration: 192 μ g L⁻¹) and completely eliminated the acute toxicity of this extract in a mouse bioassay (Nicholson et al., 1994). Interestingly, our study showed that chlorination of cyanobacterial extract was less effective. The dose of 7 mg of chlorine per liter for 30 min at pH 7.2 did not decrease MCs nor TOC concentrations. Higher doses and longer treatment times removed up to 10-30% of the original MCs and TOC levels, and caused only moderate reduction of toxicity (cytotoxicity, GJIC inhibition and MAPK activations). Chlorination apparently removed some compounds responsible for the inhibition of GJIC and activation of ERK1/2 as reflected by the slight, yet statistically significant, increase in respective IC₅₀ or LOEC values, and had only a minor effect on cytotoxicity and p38 activation. The lower effectiveness of chlorination observed in our study could be explained by the interactions of chlorine with the relatively higher concentrations of organic matter, which might have reduced the effectiveness of the oxidation process due to the competition between the toxins and the dissolved organic carbon reacting with the oxidant (Rodriguez et al., 2008).

An important problem associated with the chlorine application is the formation of by-products such as halogenated organic compounds, especially in the presence of high amounts of organic matter. These by-products can have toxic or potential carcinogenic potencies (Neale *et al.*, 2012); and also for MCs, chlorine was shown to cause substitutions and modifications of the toxic Adda moiety

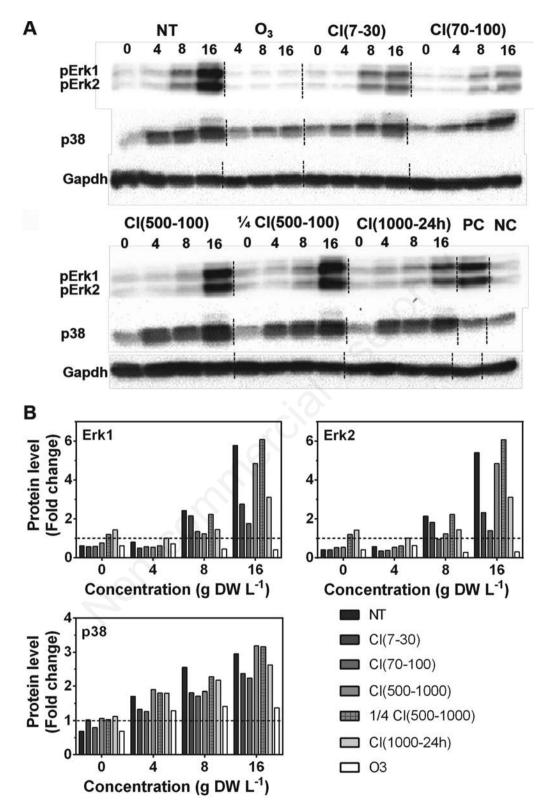


Fig. 3. Activation of mitogen-activated protein kinases (MAPKs) by studied samples after 30-min exposure to 3 different concentrations (4-16 g DW L⁻¹). Phosphorylation of extracellular receptor kinases 1 and 2 (ERK1/2) and p38 was determined by Western blotting (A). The bar graphs (B) show values from the densitometric image analysis normalized to negative control (NC=1). NT, original non-treated extract; Cl, chlorinated extracts (in parentheses - free Cl concentrations 7, 70, 500 and 1000 mg L⁻¹; treatment durations 30 min, 100 min, 24 h), O3 – ozonated extract (30 min, the flow rate of 1 L min⁻¹ with 5 g O₃ m⁻³); NC, negative control (no treatment of the cells); PC, positive control for ERK1/2 activation (12-O-tetradecanoyl phorbol-13-acetate, 10 nM, 30 min).

(Tsuji *et al.*, 1997; Merel *et al.*, 2009). In addition, *de novo* formation of chlorinated by-products with potencies to affect GJIC and activate intracellular signalling (Hakulinen *et al.*, 2004; Nishikawa *et al.*, 2006) should also be considered, and could be related to the weak efficiency in removal of GJIC inhibitions and MAPK activations during the chlorination as observed in the present study. Possible toxicity of chlorinated by-products could also explain the observed biphasic effect, when lower doses of chlorine were slightly more effective in the elimination of cytotoxicity and MAPKs activation than the higher doses, although concentrations of TOC and MCs were slightly but progressively reduced with increasing chlorine dose and treatment time.

We demonstrated that ozonation completely removed MCs, substantial fractions of TOC and protected against cytotoxicity, GJIC inhibition or activation of ERK1/2. These findings are in agreement with similar studies, which documented complete MC removal (5 mg L^{-1}) by 2 mg L⁻¹ O₃ within 2 min (Al Momani and Jarrah, 2010). Further improvements in kinetics could be achieved by increased O₃ doses and temperature, and decreased pH (Al Momani and Jarrah, 2010; Shawwa and Smith, 2001). Naturally, organic matter negatively reduces the efficiency of ozonation, but under realistic DWTP situations of levels as low as 0.05 mg L-1 of residual O3 assures MC removal (Newcombe and Nicholson, 2004; Brooke et al., 2006). Despite a high amount of organic material in our sample that also competes with toxins for ozone, ozonation was highly effective in MC reduction and elimination of toxicity even after short treatment.

A high efficiency of oxidation of MC is known to be mediated by hydroxyl radicals attacking conjugated diene structure in MC followed by the cleavage of the Adda side chain (responsible for PPase inhibition) and ultimately opening of the peptide ring (Al Momani and Jarrah, 2010; Miao et al., 2010). Biological assessments using PP-inhibition assay or mouse test confirmed elimination of the toxicity along with the described structural changes of MC (Brooke et al., 2006; Miao et al., 2010). Although ozone was quite efficient in removal of cytotoxic, GJIC inhibiting and ERK1/2 activating compounds in our study, it had only a partial effect on the removal of p38 activating components. This might indicate that p38 is not involved in GJIC inhibition and its activation was caused by metabolites with different modes of actions (Wagner and Nebreda, 2009). With respect to the critical role of p38 in cellular responses to different types of stress and also in controlling the proliferation, differentiation, survival, migration and inflammatory responses of specific cell types, further research should address interactions of cyanobacterial metabolites with this signalling pathway and evaluate its relevance as a biomarker of environmental and genotoxic stress induced by

cyanobacteria. Interestingly, activation of p38 was found to be the most sensitive endpoint in this study, where the increased levels of p38 phosphorylation were observed after 30-min exposure to the non-treated extract at concentration of 4 g DW L⁻¹, whereas significant inhibition of GJIC and activation of ERK1/2 occurred at concentrations 8 g DW L⁻¹ and higher, and significant reduction of cell viability required 24-h exposure to 12 g DW L^{-1} . Inhibition of GJIC and activation of MAPKs was induced by lower concentrations and after shorter exposures than the cytotoxic effects, which indicates that these cell signalling events were altered via rapid non-genotoxic and non-cytotoxic mechanisms. In vitro evaluation of GJIC and MAPKs thus represent a simple and sensitive bioassay for assessment of 'epigenetic toxicity' and tumour promoting potential of complex cyanobacterial extracts, which is also suitable for effect-based studies focusing on the elimination of these hazardous properties of contaminated water.

CONCLUSIONS

Ozonation of an extract of a Microcystis water bloom sample was shown to be a very effective method in the complete removal of MCs, as well as the substantial elimination of the overall cytotoxicity and tumour promotional potency. On the contrary, chlorination experiments, despite high doses and long exposures, were much less effective, and potentially led to the formation of by-products, which could add to the observed toxic effects. Our study also demonstrated strong activations of p38 MAPK by cyanobacterial samples, which were not effectively removed by chlorination and only partially by ozonation. With respect to the role of p38 in inflammation as well as maintenance of tissue homeostasis, further research should address interactions of cyanobacterial samples with this biomarker of cellular stress and evaluate its environmental relevance. In agreement with several recent reports, the study also demonstrates the need to enforce effect-based (bioassay) tools into the assessment of water quality and monitoring the efficacy of water treatment systems.

ACKNOWLEDGMENTS

Supported by the Czech Science Foundation project GA15-12408S; the Czech Republic Ministry of Education, Youth and Sports infrastructure projects No. LO1214 and LM2015051; long-term research development project RVO 67985939; and by National Institute of Environmental Health Sciences (NIEHS) grant #R01 ES013268-01A2 to Upham. We would like to thank Dr. Ondřej Mikeš for the help with TOC analysis.

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