

# **Assessment of individual and mixed toxicity of bromoform, tribromoacetic-acid and 2,4,6 tribromophenol, on the embryo-larval development of Paracentrotus lividus sea urchin**

Karine Lebaron, Lilia Mechiri, Simone Richard, Annabelle Austruy, Jean-Luc Boudenne, Stéphane Coupé

## **To cite this version:**

Karine Lebaron, Lilia Mechiri, Simone Richard, Annabelle Austruy, Jean-Luc Boudenne, et al.. Assessment of individual and mixed toxicity of bromoform, tribromoacetic-acid and 2,4,6 tribromophenol, on the embryo-larval development of Paracentrotus lividus sea urchin. Environmental Science and Pollution Research, 2019, 26 (20), pp.20573-20580.  $10.1007/s11356-019-05279-8$ . hal-02282339

# **HAL Id: hal-02282339 <https://hal.science/hal-02282339v1>**

Submitted on 9 Sep 2019

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Assessment of individual and mixture toxicity of bromoform, tribromoacetic-acid and 2,4,6

2 tribromophenol, on the embryo-larval development of *Paracentrotus lividus* sea urchin.

4 Karine Lebaron<sup>1,2</sup>, Lilia Mechiri<sup>1</sup>, Simone Richard<sup>1</sup>, Annabelle Austruy<sup>3</sup>, Jean-Luc Boudenne<sup>2</sup>, 5 Stéphane Coupé<sup>1</sup>

 $6<sup>-1</sup>$  Université Toulon, CNRS/INSU, IRD, MIO UM 110, Mediterranean Institute of Oceanography, La Garde, France.

8 <sup>2</sup> Aix-Marseille-Université, CNRS, LCE UMR7476, 13331 Marseille, France

 Institut Ecocitoyen pour la Connaissance des Pollutions, Centre de vie, La Fossette, 13270 Fos-sur-Mer, France

 Keywords: Bromoform, Tribromoacetic acid, Tribromophenol, Disinfection by-products, Sea urchin, *Paracentrotus lividus*, Ecotoxicology, Genotoxicity.

#### Abstract

 Water chlorination is the most widely used technique to avoid microbial contamination and biofouling. Adding chlorine in bromide-rich waters leads to the rapid oxidation of bromide ions and leads to the formation of brominated disinfection by products (bromo-DBPs) that exert adverse effects on various biological models. Bromo-DBPs are regularly encountered within industrialized embayments, potentially impacting marine organisms. Of these, bromoform, tribromoacetic acid and tribromophenol are among the most prevalent. In the present study, we tested the potential toxicity and genotoxicity of these disinfection by-products, using sea urchin, *Paracentrotus lividus*, 23 embryos. We highlighted that tribromophenol showed higher toxicity compared to bromoform and tribromoacetic acid. Furthermore, a synergistic effect was detected when tested in combination. Pluteus cells exposed for 1h to mixtures of DPBs at several concentrations demonstrated significant DNA damage. Finally, when compared to a non-exposed population, sea urchins living in a bromo-DPB-polluted area produced more resistant progenies, as if they were 28 locally adapted. This hypothesis remains to be tested in order to better understand the obvious impact of bromo-DBPs environments on marine wildlife.

- 
- 

#### Introduction

 Chlorination has been used worldwide, for decades, and is one of the most effective treatment for water disinfection in treatment plants for the production of tap water. Chlorination is also employed in the management and upkeep of industrial installations, where water is often used for cooling or heating purposes, as well asto prevent and control biofouling and corrosion of pipelines. Hence, in some industrialized embayment areas, huge volumes of coastal seawater are continuously being pumped and chlorinated before being re-released into the original embayment (Allonier et al. 1999; Boudjellaba et al. 2016; Manasfi et al. 2018). As chlorine is very reactive in the presence of natural organic matter, seawater chlorination generates a complex set of brominated and chloro-brominated disinfection-by-products (DBPs). Finally, benthic organism, resident in the embayment, are exposed to these DBPs (Singer 1999; Westerhoff et al. 2004; Halpern et al. 2008). Although the diversity and occurrence of brominated DBPs in contaminated coastal seawater is relatively undocumented (Richardson et al. 2007; Manasfi et al. 2017), bromoform (halomethane), tribromoacetic acid (haloacetic acid) and tribromophenol (halophenol) are among the most prevalent molecules, often measured at relatively high concentrations (i.e. µg/L) (Manasfi et al. 2018).

 Most of the DBPs so far tested have been found to be toxic and genotoxic in diverse model systems, such as bacteria, mammalian cells or zebrafish embryos. Furthermore, brominated DBPs have generally been shown to be more toxic than their chlorinated analogues (Richardson et al. 2007, 2010; Yang and Zhang 2013; Hanigan et al. 2017). It is worth noting that at the single molecule level, visible effects have always been following exposure to high concentrations of DBPs, suggesting that toxic and genotoxic risks are limited (Teixidó et al. 2015). Nevertheless, recent epidemiological research has suggested that lifetime exposure to DBPs, mainly through ingestion, would significantly increase the risk of bladder cancer in humans (Villanueva et al. 2004, 2006). Hence, there are great concerns about the level of human exposure, causing health and sanitary surveillance agencies to enact maximum concentration limits (MCL). For instance, the US EPA 58 recommends MCLs for total halomethane and haloacetic acid of 80  $\mu$ M and 60  $\mu$ M, respectively, in safe drinking water. It is notable that these values are relatively consistent with the maximum

 concentrations measured in the effluents of water treatment plants from the USA, Canada or France (CAREX Canada 2009; Mouly et al. 2009).

 There are currently no MCLs for environmental waters that harbor functional ecosystems, and which are potentially chronically exposed to DBPs. Studies that assess the toxicity of DBPs in aquatic animals remain scarce and to our knowledge, only two have tested brominated DBPs in recent years, using a zebrafish embryo model (Teixidó et al. 2015; Hanigan et al. 2017). These two studies tested a total of 11 brominated DBPs, including bromoform (BMF) and tribromoacetic acid (TBAA). Most of the brominated DBPs that were compared, proved toxic for larval development, though only at high concentration (i.e. >100 µM). Moreover, short-term exposure to chlorinated wastewater did not have any significant toxic effect. Finally, a genotoxic effect was only detected 70 for chlorodibromomethane and sodium borate at their EC<sub>50</sub> levels, 160  $\mu$ M and 0.049  $\mu$ M, respectively.

 In this study, we were interested in broadening the understanding of the toxicological impact of DBPs on coastal marine organisms, using the sea urchin, *Paracentrotus lividus,* as a proxy for the echinoderm class. *Paracentrotus lividus* lives in the first few meters of coastal areas. The species is widely distributed along the north-eastern Atlantic coast and on all Mediterranean coasts. Furthermore, they can be encountered in numerous contrasted habitats that include differences in temperature, salinity and chemical contamination (Bellas et al. 2008). While adults are benthic and relatively sedentary, the larvae produced at each reproductive event are pelagic for up to 4 weeks, thus ensuring species dispersion. The sea urchin is a relevant model organism to assess the effect of potentially harmful molecules. It is more closely related to human than the mussel (Tu et al. 2006), with two species having been fully sequenced. Reproduction is easily performed in vitro, resulting in high numbers of translucid larvae. Sea urchin larvae have been extensively used to assess the embryo-toxicity and embryo genotoxicity of molecules (Hose 1985; McGibbon and Moldan 1986; Gharred et al. 2016; Morroni et al. 2016; Trifuoggi et al., 2017; Neves et al., 2018; Messinetti et al., 2018; Periera et al., 2018; Dorey et al., 2018).

 In this study, our objective was to generate additional data in a model marine organism, of the toxic influence of three major brominated contaminants found in coastal seawater that is subjected to massive anthropogenic disturbance. With this aim in mind, we used sea urchin

 embryos to assess the toxicity and genotoxic potential of bromoform (halomethane), tribromoacetic acid (haloacetic acid) and tribromophenol (halophenol), tested either alone or in combination, on two populations of *P. lividus*, either chronically exposed or unexposed.

#### Materiel and methods

#### Sea urchins

- Adult *Paracentrotus lividus* sea urchins were harvested by scuba diving, on the morning of each
- experiment, at a non-polluted site (NPS) near the Toulon (Var, France) embayment (Garonne Bay:
- 43.098503-6.018430)and at a polluted site (PS) where sweater is chronically chlorinated (Manasfi
- et al., 2018). *P. lividus* were induced to spawn by gentle shaking.
- Sperms and eggs were individually collected, respectively dried with a micropipette and kept on ice and in 100mL of filtered seawater (FSW). Eggs were microscopically observed to verify maturity
- before adding dry sperm.
- Ten independent (i.e. unrelated) larval populations were produced by arbitrarily mixing one sperm with one egg suspension, in 50mL of FSW at 20°C at a concentration of 500 eggs/mL, under agitation for 45 min, then the fertilization rates were assessed.
- Observed fertilization rates were 100% in every experiment.
- 

#### Chemicals

- Bromoform (CAS 75-25-2), tribromoacetic acid (CAS 75-96-7) and tribromophenol (CAS 118-79-16)
- were purchased from SimgaAldrich. Stock solutions of bromoform and tribromoacetic acid were
- prepared by direct dissolution in filtered sea water (FSW). Tribromophenol was first dissolved in
- dimethylsulfoxide solvent (DMSO) to compensate for its very low water solubility, then it was
- dissolved in FSW, with a final DMSO concentration of 0.1%.
- 
- Experimental design
- *Toxicological assessments*

 Ten unrelated larval populations were used and tested in triplicates, as follows: in 24-wells microplates, suspensions of 500 fertilized eggs in 2 mL FSW were exposed for 48 hours to 7  concentrations of each chemical (Table 1). At the end of the exposure time, larvae were fixed by adding ethanol at 15% final concentration and kept at 4°C until microscopic observations wre carried out. Normal and abnormal (i.e. delayed growth, developmental anomalies) Pluteus larvae were recorded.

 Percentages of normal Pluteus larvae were plotted against chemical concentrations. We then 123 determined the dose-response curves using R software and deduced the half maximal effective 124 concentrations, EC<sub>50</sub>, here considered as the concentration of chemical at which we could observe only 50% of normal and viable larvae among the pool. The no and lowest observed effective concentrations (respectively NOEC and LOEC) were deduced from the chemical concentrations used in the experiments.

 We then assessed the toxicity of the chemicals combined in ten other independent experiments. 129 Specifically, we tested three mixtures named mix NOECs, mix LOECs and mix EC<sub>50</sub>, that is each 130 chemical within a mixture is at its specific endpoints (NOEC, LOEC or EC<sub>50</sub>). Depending on the chemical tested, controls were either exposed to FSW alone or FSW containing 0.1% DMSO.

*Genotoxicity test*

 The genotoxicity of the mix NOECs, mix LOECs and twice mix LOECs was assessed using the comet assays, based on the Tice et al. (2000) procedure.

 Five thousand, 48-h, Pluteus were exposed as described above for 1 h to the mixtures. After exposure, larvae were collected by centrifugation at 1500g for 10 min at 4°C. The pellet was resuspended in 1mL of FSW, then gently mixed with 1 mL of glycine 1M and incubated on ice for 5 min to allow for complete dissociation of cells. Fifty thousand cells were then collected and placed in 1 mL of phosphate buffer saline (PBS) 1× then centrifuged at 3000g for 5 min at 4°C. Pelleted 141 cells were then collected and gently mixed with 50  $\mu$ L of low melting point agarose at 37 $\degree$ C, then plated onto a pre-coated laboratory microscope slide (Tice et al. 2000).

143 Once prepared, slides were immersed in a lysis buffer (2.5 M NaCl, 100 mM Na<sub>2</sub>EDTA, 2H2O, 10 mM Tris pH10, 10 g N-Lauroylsarcosine, 10% DMSO and 1% Triton X100) for 90 min at 4°C. Slides 145 were then placed in an electrophoresis tank containing an alkaline solution (200 mM  $Na<sub>2</sub>EDTA$ 2H2O, 10 N NaOH) for 20 min at room temperature, to allow DNA to denature. This was followed

 by electrophoresis which was conducted for 20 min at 25V and 350mA. Thereafter, slides were plunged in a neutralization buffer (0.4 M Tris-HCl pH 7.5) for 20 min. Finally, slides were fixed with 100% methanol and dried at room temperature overnight.

 DNA was stained with a solution of SyberGreen (Sigma-Aldrich) deposited onto each slide and left to stand for 20 min in the dark. Slides were then read using an epifluorescence microscope equipped with a digital camera and dedicated software. Whole cells and comets observed within several randomly selected microscopic fields were captured, irrespective of the length and the shape of the comet (Gyori et al. 2014). Pictures were then analysed using Open Comet v1.3.1 (cometbio.org) implemented in ImageJ (SciJava) program (Gyori et al. 2014).

 DNA damage was assessed considering the olive tail moment (OTM) that consider both the length and the distribution of DNA within the comet's tail (Olive et al. 1991).

158 The control experiment consisted in larval suspensions exposed to FSW containing 700 $\mu$ M of H<sub>2</sub>O<sub>2</sub>. This concentration has been chosen after a preliminary experiment in which we tested the 160 genotoxicity of H<sub>2</sub>O<sub>2</sub> for a range of concentrations ( $R^2$  = 0.9967), as previously described (Nahon et al. 2008).

 DNA damage was assessed via the olive tail moment (OTM), which takes into consideration both 163 the length and the distribution of DNA within the comet's tail (Olive et al;, 1991).

164 The positive control experiment consisted of larval suspensions to FSW containing 500  $\mu$ M of H<sub>2</sub>O<sub>2</sub>.

This concentration was chosen after a preliminary experiment in which we tested the genotoxicity

of H2O2 for a range of concentrations (*R*=0.9983), as previously performed (Nahon et al., 2008).

#### Statistical analysis

 Statistics were performed with Rstudio (version 1.1.442) and XLSTAT (Addinsoft, 2017). Dose-170 response curves were produced using regression models. Effective concentrations (i.e.  $EC_{10}$ ,  $EC_{50}$ 171 and  $EC_{90}$ ) values were generated with the "estimating effective doses" (ED.drc) from R package. Prior to Kruskal-Wallis test, ecotoxicological and comet assay data were tested for normality. The Mann-Whitney U test was proceeded to compare data from non-polluted site (NPS) to polluted site (PS). Differences between each condition were considered significant when *p*-value < 0.05, data were expressed as mean ±S.E.M (standard error of the mean).

#### Results and discussion

- Assessment of BMF, TBAA and TBP toxicity on early embryo development
- In this study, embryos were exposed to a range of bromoform (BMF), tribromoacetic acid (TBAA)
- and tribromophenol (TBP) concentrations for 48 h (Table 1), from fertilization to the Pluteus stage.
- The control with 0.1% DMSO was not toxic for larvae, and we found typical dose-response curves,
- 181 from which  $EC_{50}$  and  $EC_{90}$  were extrapolated (Figure 1 and Table 2).



# Figure 1: Effect of tribromophenol, bromoform and tribromoacetic acid singly, on the percentage of normal pluteus growth after 48h

 TBP (halophenol) was by far the more toxic molecule compared to BMF (halomethane) and TBAA (haloacetic acid) (Table 2). BMFwas found more toxic than TBAA. This order of toxicity has regularly been reported in several aquatic organisms living in fresh or marine water (Yoshioka et al. 1985; Yang and Zhang 2013; Liu and Zhang 2014; Teixidó et al. 2015; Hanigan et al. 2017).

190 Table 1: Different endpoints ( $\mu$ M) for three DBPs, tribromophenol, bromoform and tribromoacetic acid obtain on pluteus larva after 48 hours of exposure



 Like other aquatic organisms (Yoshioka et al. 1985; Delacroix et al. 2013; Liu and Zhang 2014; Teixidó et al. 2015; Hanigan et al. 2017), *P. lividus* embryos were sensitive to relatively high 195 concentrations of the tested molecules. The  $EC_{50}$  was 14.5  $\mu$ M for TBP, 526.31  $\mu$ M for BMF and 923.5 µM for TBAA, and the lowest LOEC was 3 µM withTBP (Table 2). Except for bromoform, *P. lividus* appears to be more sensitive to TBP and TBAA, than the embryos of the marine polychaete, *Platynereis dumerilii* (Yang and Zhang 2013), or the phytoplanktonic green algae, *Tetraselmis*  199 marina (Liu and Zhang 2014). Indeed, according to the three previous studies, the EC<sub>50</sub> for bromoformin *P.lividus* is situated between the EC50 in zebrafish (200 µM (Teixidó et al. 2015)) and 201 the EC<sub>50</sub> in polychaete embryos (730  $\mu$ M (Yang and Zhang 2013)). With regard to TBP and TBAA, our model seems 4 to 14 times more sensitive to TBP and 4 to 6 times more sensitive to TBAA, when compared with other, different, biological systems (Yang and Zhang 2013; Liu and Zhang 2014; Teixidó et al. 2015).

 The first observable effect (LOEC) of the three chemicals in our model was the abnormal development of a fraction of Plutei. At higher concentrations, we observed both abnormal and delayed development, mostly between the morula and gastrula stages, although in different relative proportion, depending upon the molecule and the concentration. Of the developmental anomalies triggered by all three chemicals, we mainly observed crossed spicules, which are among the most regularly reported malformations in sea urchin larvae (Gharred et al. 2016). Interestingly, 211 TBAA-induced mortality was detected early, since we observed a black-coloured dead eggs (*unpublished data*).

### Table 2: Percentage of normal pluteus larva growth when exposed to NOEC and LOEC of bromoform, tribromophenol and tribromoacetic acid singly and combined



 Exposure for 48h to mix NOECs and mix LOECs critically reduced the proportion of normal Pluteus, 217 to 63%  $\pm$  2 and 18%  $\pm$ 1, respectively, while none survived exposure to mix EC<sub>50</sub> condition, demonstrating the combined effect of these chemicals (Table 2). The results obtained with a

 mixture of chemicals were entirely expected. However, in zebrafish embryos, the influence of chlorinated water had no detectable phenotypic impact after 5 days of exposure, suggesting that 221 a complex water environment, containing a wide diversity of DBPs at much lower concentrations than those tested, may not be deleterious for the population (Hanigan et al. 2017). Nevertheless, chronic exposure of adult fish to 0.9 nM of TBP could potentiallyhave a significant effect over a number of generations, on survival, larval development and the malformation rate (Deng et al. 2010).

#### Table 2: Percentage of normal pluteus larva growth when exposed to NOEC and LOEC of bromoform, tribromophenol and tribromoacetic acid singly and combined



 

230 We further tested the impact of embryo exposure at polluted site (PS), with embryos generated by *P. lividus* adults chronically exposed to chlorinated sea water. These PS embryos were exposed 232 to TBP, BMF and TBAA at their respective LOEC, EC<sub>50</sub> and EC<sub>90</sub>, and to mix LOECs condition. They were found slighly less sensitive to TBP and TBAA than NPS embryos, while very resistant to BMF, which is the most prevalent molecule found in seawater where genitors were harvested (Figure 2) (Manasfi et al. 2018). Moreover, exposure to mix LOECs reduced the proportion of normal Pluteus by 40% instead of 80% for the NPS embryos. This increased PS embryo resistance could be explained by either natural selection of larva, with higher fitness for the polluted site, or a parental effect (Ross et al., 2016).



239

240 Figure 2: Effect of bromoform (pink), tribromophenol (blue), tribromoacetic acid (orange), singly 241 and in combination (grey), on the percentage of normal pluteus larva growth from a polluted site, 242 after 48h of exposure. \*\*\* represent statistically significant differences at p < 0.05 from Dunnet's 243 test, between the combine molecules toxicity on larva from a non-polluted site and from a DBPs 244 polluted site.

245

#### 246 Evaluation of mixtures' genotoxicity

- 247 The alkaline comet assay has been widely used to test potential genotoxic effects of environmental
- 248 pollutants, in several model organisms such as animal embryos and plant roots, as it is an efficient
- 249 method for detecting single- and double-strand DNA breaks (Yıldız et al. 2009; Liman et al. 2011).
- 250 The results obtained from the comet assay are summarized in Figure 3.



#### 

 Figure 3: Distribution of OTM (A) and % of DNA within the comet tail (B) according to the level of exposure of Pluteus larvae to the chemicals' mixture. Boxplots encompass the 1st and the 3rd quartile. The black line within the boxplot represents the median. Tukey's whisker extents are presented (defined as 1.5 times the 1st and 3rd quartiles). The red horizontal line indicates the 95% percentile of the control condition used as a reference, and the numbers in red indicate the proportion of comets presenting a higher value than the reference

 Here we tested whether 1h exposure to mixtures of TBP, BMF and TBAA, at their respective NOEC, LOEC and twice their LOEC concentrations, could induce DNA strand breakages in Pluteus larvae. 261 It is likely that given the short-term exposures to the mixtures, we observed the obvious potential 262 of our mixtures to induce DNA strand breakages, prior to any activation of DNA repair mechanisms that would counteract the induced DNA damage. The level of genotoxicity was assessed in positive and negative controls and exposed groups according to the OTM and the percentage of DNA

265 (%DNA) measured in the comet tail. Furthermore, we only assessed the proportion of cells within 266 each group that exceeded a threshold value, corresponding to the 95<sup>th</sup> percentile of the control 267 group for the OTM and %DNA. Under controlled conditions, the 95<sup>th</sup> percentile for OTM and %DNA 268 were 2.6 and 16.5%, respectively.

 For the two parameters considered, we found normal data distributions within the negative control groups (Shapiro-Wilk, *p*=0.1) but not in the tested groups or in the positive control (Shapiro-271 Wilk, *p*<0.0001). The non-parametric test did not reveal any influence of exposure level based on 272 the OTM but demonstrated that the group exposed to the mixture at twice the LOEC concentration 273 (mix LOEC\_ X 2), had significantly more DNA within the comet tail, compared to all other groups (Kruskal-Wallis, *p*<0.033). This result would suggest that this last condition only had a genotoxic 275 effect. However, the discrepancy in the results between controls and tested groups for normality 276 of data distribution indicates exposure had an influence on a fraction of the whole cells within each group, as previously reported (Taban et al. 2004). Indeed, the proportion of cells with an OTM higher than 2.6 regularly increases with the concentration and gives higher OTM values than observed in the negative control group (Figure 3, A). This increase of the proportion of cells has 280 also been observed for the %DNA parameter, although it is clearly noticeable at the mix LOEC\_X2 condition (Figure 3, B). Hence, overall, our results suggest a dose response when taking into consideration both the effect on DNA and the occurrence of responding cells. Similar results have already been reported in short-term exposure experiments, such as those performed on CHO (Chinese hamster ovary) cells exposed to TBA for 4h (Plewa et al. 2008) and occurrence (Taban et al. 2004).

286 It is interesting to note that at the mix LOEC X2 condition, we found that 7% of the cells were 287 distributed with more than 90 % of their DNA within the comet tail. Among these, a few cells also 288 displayed a high OTM value, which relates to the length of the comet. A similar result was obtained 289 with some cells from the positive control group (Figure 3, B). This could be indicative of ongoing 290 apoptosis (Ríos et al. 2003; Collins et al., 2004). Evidence for this has already been demonstrated 291 in other studies. It has been shown that hydrogen peroxide can induce apoptosis, causing cell death 292 in less than 2 h in 293T cells exposed to more than 400  $\mu$ M of H<sub>2</sub>O<sub>2</sub> (Xiang et al. 2016). A recent 293 study has also suggested that halogenated DBPs might trigger apoptosis at high concentration (Ríos  et al. 2003; Collins, 2004). Thus, in our study, the mixtures might have the potential to also trigger apoptosis in a fraction of the larval cells.

 This distribution pattern is difficult to explain at this stage, and it would be interesting to assess whether a certain cell type would be more sensitive to mixtures than others, or if the external cells are more impacted than those embedded within the larval body, for instance. This has already been reported in zebrafish exposed for a much longer time (i.e. 72 h) to BMF (Teixidó et al. 2015).

 Finally, we have highlighted with our experimental conditions that a significant genotoxic effect, observed with high doses of a cocktail of DBPs, are generally consistent with previous findings (Teixidó et al. 2015; Hanigan et al. 2017).

#### Conclusion

 This study has furthered our knowledge of the impact of bromoform, tribromophenol and tribromoacetic acid, tested alone and in combination, on sea urchin, *Paracentrotus lividus,* development. We have revealed that all three molecules could impair embryo development with the order of toxicity TBP>BMF>TBAA. TBP is 10 to 30 times more toxic compared to the two others molecules. The mixture of the three DBPs also significantly impaired development and were proved weakly genotoxic at low doses (i.e. mix NOECs). Increasing the concentration of the mixture to mix LOECs significantly increases cytotoxicity, as well as increasing the level of DNA damage and the proportion of cells affected.

313 In this study, we have highlighted a potential genome by environment interaction, particularly regarding BMF. Further experiments will be required in order to assess the relative influence of a potential parental effect, as well as local genetic adaptation in the observed resistance.

### Acknowledgment

 This work was co-funded by the French National Research Agency (ANR) within the project FOS- SEA (ANR-16-CE34-0009) and by Conseil regional Provence Alpes Côte d'Azur (Reginal Council SUD).

## References

 Allonier, A.-S. et al. 1999. Determination of dihaloacetonitriles and halophenols in chlorinated sea water. *Talanta*

 Bellas, J. et al. 2008. Integrative assessment of coastal pollution in a Ría coastal system (Galicia, NW Spain): Correspondence between sediment chemistry and toxicity. *Chemosphere* 72(5), pp. 826–835. doi: 10.1016/j.chemosphere.2008.02.039.

 Boudjellaba, D. et al. 2016. Chlorination by-product concentration levels in seawater and fish of an industrialised bay (Gulf of Fos, France) exposed to multiple chlorinated effluents. *Science of The Total Environment* 541, pp. 391–399. doi: 10.1016/j.scitotenv.2015.09.046.

 CAREX Canada 2009. *Les sous-produits de chloration*. School of Environmental Health University of British Columbia. Available at: https://www.carexcanada.ca/fr/chloration.pdf [Accessed: 12 December 2018].

 Delacroix, S. et al. 2013. Disinfection by-products and ecotoxicity of ballast water after oxidative treatment – Results and experiences from seven years of full-scale testing of ballast water management systems. *Marine Pollution Bulletin* 73(1), pp. 24–36. doi: 10.1016/j.marpolbul.2013.06.014.

 Deng, J. et al. 2010. Chronic exposure to environmental levels of tribromophenol impairs zebrafish reproduction. *Toxicology and Applied Pharmacology* 243(1), pp. 87–95. doi: 10.1016/j.taap.2009.11.016.

 Gharred, T. et al. 2016. Assessment of the individual and mixture toxicity of cadmium, copper and oxytetracycline, on the embryo-larval development of the sea urchin Paracentrotus lividus. *Environmental Science and Pollution Research* 23(18), pp. 18064–18072. doi: 10.1007/s11356-016- 6988-3.

- Gyori, B.M. et al. 2014. OpenComet: An automated tool for comet assay image analysis. *Redox Biology* 2, pp. 457–465. doi: 10.1016/j.redox.2013.12.020.
- Halpern, B. et al. 2008. A Global Map of Human Impact on Marine Ecosystems. *Science* 319, pp. 948–952. doi: 10.1126/science.1149345.

 Hanigan, D. et al. 2017. Zebrafish embryo toxicity of 15 chlorinated, brominated, and iodinated disinfection by-products. *Journal of Environmental Sciences* 58, pp. 302–310. doi: 10.1016/j.jes.2017.05.008.

 Liman, R. et al. 2011. Determination of genotoxicity of Fenaminosulf by Allium and Comet tests. *Pesticide Biochemistry and Physiology* 99(1), pp. 61–64. doi: 10.1016/j.pestbp.2010.10.006.

 Liu, J. and Zhang, X. 2014. Comparative toxicity of new halophenolic DBPs in chlorinated saline wastewater effluents against a marine alga: Halophenolic DBPs are generally more toxic than haloaliphatic ones. *Water Research* 65, pp. 64–72. doi: 10.1016/j.watres.2014.07.024.

 Manasfi, T. et al. 2017. Assessing the genotoxicity of two commonly occurring byproducts of water disinfection: Chloral hydrate and bromal hydrate. *Mutation Research/Genetic Toxicology and Environmental Mutagenesis* 813, pp. 37–44. doi: 10.1016/j.mrgentox.2016.11.009.

 Manasfi, T. et al. 2018. Characterization of chlorination byproducts in marine waters and sediments in a semi-enclosed bay exposed to multiple industrial chlorinated effluents.

 Mouly, D. et al. 2009. Les sous-produits de chloration dans l'eau destinée à la consommation humaine en France., p. 76.

 Nahon, S. et al. 2008. Improved Comet assay for the assessment of UV genotoxicity in Mediterranean sea urchin eggs. *Environmental and Molecular Mutagenesis* 49(5), pp. 351–359. doi: 10.1002/em.20391.

 Plewa, M.J. et al. 2008. Comparative Mammalian Cell Toxicity of N-DBPs and C-DBPs. In: Karanfil, T. et al. eds. *Disinfection By-Products in Drinking Water*. Washington, DC: American Chemical Society, pp. 36–50. Available at: http://pubs.acs.org/doi/abs/10.1021/bk-2008-0995.ch003 [Accessed: 7 June 2018].

 Richardson, S. et al. 2007. Occurrence, genotoxicity, and carcinogenicity of regulated and emerging disinfection by-products in drinking water: A review and roadmap for research. *Mutation Research/Reviews in Mutation Research* 636(1–3), pp. 178–242. doi: 10.1016/j.mrrev.2007.09.001.

 Richardson, S.D. et al. 2010. What's in the Pool? A Comprehensive Identification of Disinfection By-products and Assessment of Mutagenicity of Chlorinated and Brominated Swimming Pool Water.

*Environmental Health Perspectives* 118(11), pp. 1523–1530. doi: 10.1289/ehp.1001965.

378 Ríos, J.C. et al. 2003. Tribromophenol induces the differentiation of SH-SY5Y human neuroblastoma cells in vitro. *Toxicology in Vitro* 17(5–6), pp. 635–641. doi: 10.1016/S0887-2333(03)00110-3.

Singer, P.. 1999. Humic substances as precursors for potentially harmful disinfection by-products.

 *Water Science and Technology* 40(9). Available at: http://wst.iwaponline.com/content/40/9/25 [Accessed: 6 June 2018].

*Some drinking-water disinfectants and contaminants, including arsenic.* 2004. Lyon: IARC.

- Taban, I.C. et al. 2004. Detection of DNA damage in mussels and sea urchins exposed to crude oil using comet assay. *Marine Environmental Research* 58(2), pp. 701–705. doi: 10.1016/j.marenvres.2004.03.018.
- Teixidó, E. et al. 2015. Developmental effects and genotoxicity of 10 water disinfection by-products in zebrafish. *Journal of Water and Health* 13(1), pp. 54–66. doi: 10.2166/wh.2014.006.
- Tice, R.R. et al. 2000. Single cell gel/comet assay: Guidelines for in vitro and in vivo genetic toxicology testing. *Environmental and Molecular Mutagenesis* 35(3), pp. 206–221. doi: 10.1002/(SICI)1098-2280(2000)35:3<206::AID-EM8>3.0.CO;2-J.
- Tu, Q. et al. 2006. Sea urchin Forkhead gene family: Phylogeny and embryonic expression. *Developmental Biology* 300(1), pp. 49–62. doi: 10.1016/j.ydbio.2006.09.031.
- Villanueva, C.M. et al. 2004. Disinfection Byproducts and Bladder Cancer: A Pooled Analysis. *Epidemiology* 15(3), pp. 357–367.
- Villanueva, C.M. et al. 2006. Bladder Cancer and Exposure to Water Disinfection By-Products through Ingestion, Bathing, Showering, and Swimming in Pools. *American Journal of Epidemiology* 165(2), pp. 148–156. doi: 10.1093/aje/kwj364.
- Westerhoff, P. et al. 2004. Reactivity of natural organic matter with aqueous chlorine and bromine. *Water Research* 38(6), pp. 1502–1513. doi: 10.1016/j.watres.2003.12.014.
- Xiang, J. et al. 2016. Is Hydrogen Peroxide a Suitable Apoptosis Inducer for All Cell Types? *BioMed Research International* 2016, pp. 1–6. doi: 10.1155/2016/7343965.
- Yang, M. and Zhang, X. 2013. Comparative Developmental Toxicity of New Aromatic Halogenated DBPs in a Chlorinated Saline Sewage Effluent to the Marine Polychaete *Platynereis dumerilii*. *Environmental Science & Technology* 47(19), pp. 10868–10876. doi: 10.1021/es401841t.
- Yıldız, M. et al. 2009. Determination of genotoxic effects of copper sulphate and cobalt chloride in Allium cepa root cells by chromosome aberration and comet assays. *Chemosphere* 75(7), pp. 934– 938. doi: 10.1016/j.chemosphere.2009.01.023.
- Yoshioka, Y. et al. 1985. Testing for the toxicity of chemicals with Tetrahymena pyriformis. *Science of The Total Environment* 43(1–2), pp. 149–157. doi: 10.1016/0048-9697(85)90037-3.