

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

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

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4 Karine Lebaron^{1,2}, Lilia Mechiri¹, Simone Richard¹, Annabelle Austruy³, Jean-Luc Boudenne²,
5 Stéphane Coupé¹

¹ Université Toulon, CNRS/INSU, IRD, MIO UM 110, Mediterranean Institute of Oceanography, La
 Garde, France.

8 ² Aix-Marseille-Université, CNRS, LCE UMR7476, 13331 Marseille, France

³ Institut Ecocitoyen pour la Connaissance des Pollutions, Centre de vie, La Fossette, 13270 Fos sur-Mer, France

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 urchin, *Paracentrotus lividus*, Ecotoxicology, Genotoxicity.

14

15 Abstract

Water chlorination is the most widely used technique to avoid microbial contamination and 16 biofouling. Adding chlorine in bromide-rich waters leads to the rapid oxidation of bromide ions and 17 18 leads to the formation of brominated disinfection by products (bromo-DBPs) that exert adverse 19 effects on various biological models. Bromo-DBPs are regularly encountered within industrialized 20 embayments, potentially impacting marine organisms. Of these, bromoform, tribromoacetic acid 21 and tribromophenol are among the most prevalent. In the present study, we tested the potential 22 toxicity and genotoxicity of these disinfection by-products, using sea urchin, Paracentrotus lividus, embryos. We highlighted that tribromophenol showed higher toxicity compared to bromoform 23 and tribromoacetic acid. Furthermore, a synergistic effect was detected when tested in 24 25 combination. Pluteus cells exposed for 1h to mixtures of DPBs at several concentrations 26 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 27 28 locally adapted. This hypothesis remains to be tested in order to better understand the obvious impact of bromo-DBPs environments on marine wildlife. 29

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- 31

32 Introduction

Chlorination has been used worldwide, for decades, and is one of the most effective treatment for 33 34 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 35 cooling or heating purposes, as well as to prevent and control biofouling and corrosion of pipelines. 36 Hence, in some industrialized embayment areas, huge volumes of coastal seawater are 37 continuously being pumped and chlorinated before being re-released into the original embayment 38 39 (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 40 and chloro-brominated disinfection-by-products (DBPs). Finally, benthic organism, resident in the 41 embayment, are exposed to these DBPs (Singer 1999; Westerhoff et al. 2004; Halpern et al. 2008). 42 Although the diversity and occurrence of brominated DBPs in contaminated coastal seawater is 43 relatively undocumented (Richardson et al. 2007; Manasfi et al. 2017), bromoform (halomethane), 44 45 tribromoacetic acid (haloacetic acid) and tribromophenol (halophenol) are among the most 46 prevalent molecules, often measured at relatively high concentrations (i.e. μ g/L) (Manasfi et al. 2018). 47

Most of the DBPs so far tested have been found to be toxic and genotoxic in diverse model systems, 48 49 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, 50 51 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, 52 suggesting that toxic and genotoxic risks are limited (Teixidó et al. 2015). Nevertheless, recent 53 54 epidemiological research has suggested that lifetime exposure to DBPs, mainly through ingestion, 55 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 56 57 surveillance agencies to enact maximum concentration limits (MCL). For instance, the US EPA 58 recommends MCLs for total halomethane and haloacetic acid of 80 μ M and 60 μ M, respectively, 59 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 orFrance (CAREX Canada 2009; Mouly et al. 2009).

There are currently no MCLs for environmental waters that harbor functional ecosystems, and 62 which are potentially chronically exposed to DBPs. Studies that assess the toxicity of DBPs in 63 aquatic animals remain scarce and to our knowledge, only two have tested brominated DBPs in 64 recent years, using a zebrafish embryo model (Teixidó et al. 2015; Hanigan et al. 2017). These two 65 studies tested a total of 11 brominated DBPs, including bromoform (BMF) and tribromoacetic acid 66 67 (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 68 wastewater did not have any significant toxic effect. Finally, a genotoxic effect was only detected 69 for chlorodibromomethane and sodium borate at their EC₅₀ levels, 160 μ M and 0.049 μ M, 70 71 respectively.

72 In this study, we were interested in broadening the understanding of the toxicological impact of 73 DBPs on coastal marine organisms, using the sea urchin, Paracentrotus lividus, as a proxy for the 74 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. 75 Furthermore, they can be encountered in numerous contrasted habitats that include differences 76 77 in temperature, salinity and chemical contamination (Bellas et al. 2008). While adults are benthic 78 and relatively sedentary, the larvae produced at each reproductive event are pelagic for up to 4 79 weeks, thus ensuring species dispersion. The sea urchin is a relevant model organism to assess the 80 effect of potentially harmful molecules. It is more closely related to human than the mussel (Tu et 81 al. 2006), with two species having been fully sequenced. Reproduction is easily performed in vitro, 82 resulting in high numbers of translucid larvae. Sea urchin larvae have been extensively used to 83 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; 84 85 Messinetti et al., 2018; Periera et al., 2018; Dorey et al., 2018).

86 In this study, our objective was to generate additional data in a model marine organism, of the 87 toxic influence of three major brominated contaminants found in coastal seawater that is 88 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.

92

93 Materiel and methods

94 <u>Sea urchins</u>

- 95 Adult Paracentrotus lividus sea urchins were harvested by scuba diving, on the morning of each
- 96 experiment, at a non-polluted site (NPS) near the Toulon (Var, France) embayment (Garonne Bay:
- 97 43.098503-6.018430) and at a polluted site (PS) where sweater is chronically chlorinated (Manasfi
- 98 et al., 2018). *P. lividus* were induced to spawn by gentle shaking.
- 99 Sperms and eggs were individually collected, respectively dried with a micropipette and kept on
 100 ice and in 100mL of filtered seawater (FSW). Eggs were microscopically observed to verify maturity
- 101 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.
- 105 Observed fertilization rates were 100% in every experiment.
- 106

107 <u>Chemicals</u>

- **108** Bromoform (CAS 75-25-2), tribromoacetic acid (CAS 75-96-7) and tribromophenol (CAS 118-79-16)
- 109 were purchased from SimgaAldrich. Stock solutions of bromoform and tribromoacetic acid were
- 110 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%.
- 113
- 114 Experimental design
- **115** *Toxicological assessments*

Ten unrelated larval populations were used and tested in triplicates, as follows: in 24-wellsmicroplates, suspensions of 500 fertilized eggs in 2 mL FSW were exposed for 48 hours to 7

118 concentrations of each chemical (Table 1). At the end of the exposure time, larvae were fixed by 119 adding ethanol at 15% final concentration and kept at 4°C until microscopic observations wre 120 carried out. Normal and abnormal (i.e. delayed growth, developmental anomalies) Pluteus larvae 121 were recorded.

Percentages of normal Pluteus larvae were plotted against chemical concentrations. We then determined the dose-response curves using R software and deduced the half maximal effective concentrations, EC₅₀, 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. Specifically, we tested three mixtures named mix NOECs, mix LOECs and mix EC₅₀, that is each chemical within a mixture is at its specific endpoints (NOEC, LOEC or EC₅₀). Depending on the chemical tested, controls were either exposed to FSW alone or FSW containing 0.1% DMSO.

132

133 *Genotoxicity test*

The genotoxicity of the mix NOECs, mix LOECs and twice mix LOECs was assessed using the cometassays, 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 cells were then collected and gently mixed with 50 μL of low melting point agarose at 37°C, then plated onto a pre-coated laboratory microscope slide (Tice et al. 2000).

Once prepared, slides were immersed in a lysis buffer (2.5 M NaCl, 100 mM Na₂EDTA,2H2O, 10
mM Tris pH10, 10 g N-Lauroylsarcosine, 10% DMSO and 1% Triton X100) for 90 min at 4°C. Slides
were then placed in an electrophoresis tank containing an alkaline solution (200 mM Na₂EDTA
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 lengthand the distribution of DNA within the comet's tail (Olive et al. 1991).

The control experiment consisted in larval suspensions exposed to FSW containing 700 μ M of H₂O₂. This concentration has been chosen after a preliminary experiment in which we tested the genotoxicity of H₂O₂ for a range of concentrations (R² = 0.9967), as previously described (Nahon et al. 2008).

DNA damage was assessed via the olive tail moment (OTM), which takes into consideration boththe 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 μ M of H₂O₂.

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

of H_2O_2 for a range of concentrations (*R*=0.9983), as previously performed (Nahon et al., 2008).

167

168 <u>Statistical analysis</u>

Statistics were performed with Rstudio (version 1.1.442) and XLSTAT (Addinsoft, 2017). Doseresponse curves were produced using regression models. Effective concentrations (i.e. EC_{10} , EC_{50} 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).

176 Results and discussion

- 177 Assessment of BMF, TBAA and TBP toxicity on early embryo development
- 178 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.
- 180 The control with 0.1% DMSO was not toxic for larvae, and we found typical dose-response curves,
- 181 from which EC₅₀ and EC₉₀ were extrapolated (Figure 1 and Table 2).



182

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

185

TBP (halophenol) was by far the more toxic molecule compared to BMF (halomethane) and TBAA
(haloacetic acid) (Table 2). BMF was 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).
Table 1: Different endpoints (μM) for three DBPs, tribromophenol, bromoform and tribromoacetic

191	acid c	btain	on p	luteus	larva	after	48	hours of	exposure	

	Tribromophenol	Bromoform	Tribromoacetic acid
LOEC µM	3,02E+00	1,98E+02	3,37E+02
$\text{EC}_{\text{50}}\mu\text{M}$	1,45E+01	5,26E+02	9,23E+02
EC ₉₀ μM	3,14E+01	1,05E+03	1,78E+03

192

193 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 194 concentrations of the tested molecules. The EC₅₀ was 14.5 μ M for TBP, 526.31 μ M for BMF and 195 196 923.5 μ M for TBAA, and the lowest LOEC was 3 μ M withTBP (Table 2). Except for bromoform, P. *lividus* appears to be more sensitive to TBP and TBAA, than the embryos of the marine polychaete, 197 198 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 EC₅₀ for 200 bromoform in *P.lividus* is situated between the EC_{50} in zebrafish (200 μ M (Teixidó et al. 2015)) and the EC₅₀ in polychaete embryos (730 μM (Yang and Zhang 2013)). With regard to TBP and TBAA, 201 202 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 203 2014; Teixidó et al. 2015). 204

205 The first observable effect (LOEC) of the three chemicals in our model was the abnormal 206 development of a fraction of Plutei. At higher concentrations, we observed both abnormal and 207 delayed development, mostly between the morula and gastrula stages, although in different relative proportion, depending upon the molecule and the concentration. Of the developmental 208 209 anomalies triggered by all three chemicals, we mainly observed crossed spicules, which are among 210 the most regularly reported malformations in sea urchin larvae (Gharred et al. 2016). Interestingly, TBAA-induced mortality was detected early, since we observed a black-coloured dead eggs 211 (unpublished data). 212

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

	% pluteus larva		
	NOEC	LOEC	
Bromoform (bmf)	99 ± 1	94 ± 5	
Tribromophenol (tbp)	96 ± 1	90 ± 8	
Tribromoacetic acid (tbaa)	94 ± 1	80 ± 4	
Cocktail	63 ± 3 🕂 -33%	18±6 🕂 -70%	

215

Exposure for 48h to mix NOECs and mix LOECs critically reduced the proportion of normal Pluteus, to $63\% \pm 2$ and $18\% \pm 1$, respectively, while none survived exposure to mix EC₅₀ 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 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

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Cocktail	63 ± 3 🕂 -33%	18±6 ↓ -70%	

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- 229

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 231 232 to TBP, BMF and TBAA at their respective LOEC, EC₅₀ and EC₉₀, and to mix LOECs condition. They 233 were found slighly less sensitive to TBP and TBAA than NPS embryos, while very resistant to BMF, 234 which is the most prevalent molecule found in seawater where genitors were harvested (Figure 2) 235 (Manasfi et al. 2018). Moreover, exposure to mix LOECs reduced the proportion of normal Pluteus 236 by 40% instead of 80% for the NPS embryos. This increased PS embryo resistance could be 237 explained by either natural selection of larva, with higher fitness for the polluted site, or a parental 238 effect (Ross et al., 2016).



239

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



251

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

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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.
It is likely that given the short-term exposures to the mixtures, we observed the obvious potential
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

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

For the two parameters considered, we found normal data distributions within the negative 269 control groups (Shapiro-Wilk, p=0.1) but not in the tested groups or in the positive control (Shapiro-270 271 Wilk, p < 0.0001). The non-parametric test did not reveal any influence of exposure level based on the OTM but demonstrated that the group exposed to the mixture at twice the LOEC concentration 272 273 (mix LOEC_ X 2), had significantly more DNA within the comet tail, compared to all other groups 274 (Kruskal-Wallis, p < 0.033). This result would suggest that this last condition only had a genotoxic effect. However, the discrepancy in the results between controls and tested groups for normality 275 of data distribution indicates exposure had an influence on a fraction of the whole cells within each 276 277 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 278 observed in the negative control group (Figure 3, A). This increase of the proportion of cells has 279 280 also been observed for the %DNA parameter, although it is clearly noticeable at the mix LOEC X2 281 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 282 already been reported in short-term exposure experiments, such as those performed on CHO 283 284 (Chinese hamster ovary) cells exposed to TBA for 4h (Plewa et al. 2008) and occurrence (Taban et 285 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 μ M of H₂O₂ (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 triggerapoptosis 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).

303

304 Conclusion

This study has furthered our knowledge of the impact of bromoform, tribromophenol and 305 tribromoacetic acid, tested alone and in combination, on sea urchin, Paracentrotus lividus, 306 307 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 308 309 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 310 311 LOECs significantly increases cytotoxicity, as well as increasing the level of DNA damage and the proportion of cells affected. 312

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.

316

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