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# Combined effects of okadaic acid and pectenotoxin-2, 13-desmethylspirolide C or yessotoxin in human intestinal Caco-2 cells

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#### 17 Abstract (150-250 words)

Lipophilic phycotoxins are secondary metabolites produced by phytoplanktonic species. They 18 19 accumulate in filtering shellfish and can cause human intoxications. Humans can be exposed to 20 combinations of several phycotoxins. The toxicological effects of phycotoxin mixtures on human 21 health are largely unknown. Published data on phycotoxin co-exposure show that okadaic acid (OA) is simultaneously found with pectenetoxin-2 (PTX-2), 13-desmethylspirolide C (also known as SPX-1), 22 23 or yessotoxin (YTX). Therefore, the aim of this study was to examine the effects of three binary 24 mixtures, OA/PTX-2, OA/SPX-1 and OA/YTX on human intestinal Caco-2 cells. A multi-parametric 25 approach for cytotoxicity determination was applied using a high-content analysis platform, including markers for cell viability, oxidative stress, inflammation, and DNA damage. Mixtures effects were 26 27 analyzed using two additivity mathematical models. Our assays revealed that OA induced cytotoxicity, 28 DNA strand breaks and interleukin 8 release. PTX-2 slightly induced DNA strand breaks, whereas 29 SPX-1 and YTX did not affect the investigated endpoints. The combination of OA with another toxin 30 resulted in reduced toxicity at low concentrations, suggesting antagonistic effects, but in increased 31 effects at higher concentrations, suggesting additive or synergistic effects. Taken together, our results 32 demonstrated that the cytotoxic effects of binary mixtures of lipophilic phycotoxins could not be 33 predicted by additivity mathematical models. In conclusion, the present data suggest that combined 34 effects of phycotoxins may occur which might have the potential to impact on risk assessment of these 35 compounds.

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37 Keywords: phycotoxins, mixtures, antagonism, synergism, genotoxicity, inflammation

Abbreviations<sup>1</sup>: ABC: ATP-binding cassette transporter; ADME: absorption, distribution,
metabolism, excretion; CI: combination index; CYP: cytochrome P450; DSP: diarrhetic shellfish
poisoning; EFSA: European Food Safety Authority; IL-8: interleukin 8; OA: okadaic acid; P-gp: Pglycoprotein; PP2A: protein phosphatase 2A; PTX-2: pectenotoxin-2; PXR: pregnane-X-receptor;
ROS: reactive oxygen species; SM: shellfish meat; SPX-1: 13-desmethylspirolide C; TEF: toxicity
equivalent factor; YTX: yessotoxin.

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#### 45 **1. Introduction (750 words)**

46 Marine biotoxins are secondary metabolites produced by specific phytoplanktonic species (Visciano et al., 2016). Among the lipophilic marine biotoxins, four main families have been described: okadaic 47 acid (OA) and dinophysistoxins (DTXs), pectenotoxins (PTXs), yessotoxins (YTXs) and finally cyclic 48 49 imines (including spirolides, pinnatoxins, pteriatoxins and gymnodimines). OA and DTXs are 50 responsible for diarrhetic shellfish poisoning (DSP), characterized by diarrhea, nausea, abdominal pain 51 and vomiting (Valdiglesias et al., 2013). The OA group of toxins inhibits serine/threonine protein 52 phosphatase 2A (PP2A) and, to a lesser extent, also PP1 (Takai et al., 1992). The group of PTXs, principally PTX-2, is not anymore considered a DSP toxin since its implication in gastro-intestinal 53 symptoms is not clear (Ito et al., 2008). At the cellular level, PTX-2 provokes actin depolarization 54 55 leading to a disruption of the cytoskeleton (Alligham et al., 2007). The group of YTXs has no recorded 56 effects in humans but has been frequently found together with OA in shellfish (Alarcan et al., 2018). YTX displays in vitro toxicity, for instance apoptosis induction and disturbance of the calcium flux 57 (Tubaro et al., 2008). Its mechanism of action is still unknown but some studies suggest a possible 58 59 connection with autophagy (Fernández-Araujo et al., 2015). Finally, potent neurological effects in 60 mice have been reported for the group of cyclic imines (Munday et al., 2012). SPX-1 is a selective 61 inhibitor of nicotinic acetylcholine receptors (Araoz et al., 2015).

62 For some of them, regulatory limits were set to protect consumers from their potential harm as they can contaminate shellfish and be responsible for foodborne outbreaks (EFSA report 2009). OA and 63 64 PTX-2 together share a limit of 160 µg /kg shellfish meat (SM), and for YTX and its analogues the 65 limit is set to 3.75 mg YTX eq/kg SM. However, in the absence of reports on human intoxications, no regulatory limits have been established for the group of cyclic imines, even though they are frequently 66 detected in shellfish and can be very toxic to mice following intra-periteonal or oral administration 67 (Munday et al., 2012). Mixtures of marine biotoxins are only regarded by the abovementioned EFSA 68 69 opinion in case of toxin analogs with established toxicity equivalent factors (TEFs). However, studies regarding deleterious effects generated by the mixtures of toxins from different chemical groups are 70 lacking. Therefore the impact of toxin co-occurrence on toxicity needs to be more deeply investigated. 71

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A high number of different approaches have been described to analyze the combined effects of
 mixtures (Foucquier and Guedj 2015). Even though antagonism and synergism are well-established

<sup>1</sup>**Abbreviations:** ABC: ATP-binding cassette transporter; ADME: absorption, distribution, metabolism, excretion; CI: combination index; CYP: cytochrome P450; DSP: diarrhetic shellfish poisoning; EFSA: European Food Safety Authority; IL-8: interleukin 8; OA: okadaic acid; P-gp: P-glycoprotein; PP2A: protein phosphatase 2A; PTX-2: pectenotoxin-2; PXR: pregnane-X-receptor; ROS: reactive oxygen species; SPX-1: 13-desmethylspirolide C; TEF: toxicity equivalent factor; YTX: yessotoxin.

75 concepts, the definition of additivity is broad (Foucquier and Guedj, 2015) resulting in several proposed models. Generally speaking, additivity models allow predicting the additive effects of a 76 77 mixture from the toxicity of each component. They assume that each component exerts an effect without influencing the action of the other ones. Deviation from predicted additivity is considered as 78 79 synergism or antagonism. The two main concepts used are concentration addition (CA) and 80 independent action (IA). The CA concept relies on the assumption that the components in the mixture act on the same biological sites, by the same mechanisms of action and differ only in their potency 81 (Loewe and Muischnek, 1926). The IA concept established by Bliss (1939) relies on the hypothesis 82 that a combined effect can be calculated from the responses of the individual mixture components, 83 84 assuming that each one acts independently without interfering with the others (different sites of 85 action).

So far, only few studies have been undertaken with mixtures of phycotoxins. No combined effects were reported in rodents except in the distribution into internal organs where a reduced level was observed in case of mixtures (Aasen et al., 2011; Sosa et al., 2013). In human neuroblastoma cells, OA co-incubated with YTX or SPX-1 showed additive effect towards cell viability (Rodriguez et al., 2015). However, on proliferative human intestinal epithelial cells, binary combinations of azaspiracid-1 (AZA-1)/OA and YTX/OA resulted in an increasing antagonistic effect on viability when toxin concentrations raised, whereas AZA-1/YTX mixtures showed dose-dependent synergism (Ferron et col. 2016).

93 al., 2016).

In this study we chose relevant mixtures based on our recently published review (Alarcan et al., 2018),
highlighting that OA was predominantly found as the main toxin combined with PTX-2, YTX or SPX-

96 1. Our objective was to complete the previous data by assessing a panel of different toxicity endpoints

- 97 (cell viability, oxidative stress, inflammation, and DNA strand breaks) in the human Caco-2 cell line
- 98 with binary mixtures of lipophilic toxins.
- 99

## 100 2. Materials and methods

#### 101 2.1 Establishment of mixtures

Three binary mixtures were selected in the following molar ratios according to published literature on 102 lipophilic phycotoxins co-exposure (Alarcan et al., 2018): 3:1 for OA/PTX-2 and OA/YTX and 9:1 for 103 OA/SPX-1. Concentrations ranges were established based on preliminary cell viability experiments 104 and published data. PTX-2, YTX and SPX-1 did not show any cytotoxicity (NRU, DAPI cell count) 105 after 24h on differentiated Caco-2 cells up to 500, 200 and 510 nM, respectively (unpublished data). 106 107 Based on the cytotoxicity results (MTT assay) for OA reported by Ehlers et al. (2011), a maximum 108 concentration of 600 nM was chosen in order to study biological responses in both cytotoxic and non-109 cytotoxic conditions. Then, based on the ratios of toxins previously indicated, we deduced the 110 concentrations for PTX-2, YTX and SPX-1 from those chosen for OA.

## 111 2.2 Chemicals

112 Toxin standards were purchased from the National Research Council Institute for Marine Biosciences

113 (Halifax, NS Canada). DCFH-DA, potassium bromate (KBrO<sub>3</sub>), sulfuric acid (H<sub>2</sub>SO<sub>4</sub>) and TNFα were

- 114 purchased from Sigma-Aldrich (St. Louis, MO, USA). Primary anti-interleukin 8 (IL-8) antibody
- 115 (M801), biotin-conjugated human IL-8 (M802B), HRP-conjugated streptavidin (N100), and 3,3',5,5'-
- tetramethylbenzidine (TMB) were obtained from Thermo Fisher Scientific (Waltham, MA, USA).
- 117 Mouse monoclonal anti γH2AX S19 (ab26350) and goat pAb to Ms IgG Alexa Fluor 647 (ab150115)

were purchased from Abcam (Cambridge, UK). All other chemicals including ethanol (EtOH),
methanol (MeOH) and dimethyl sulfoxide (DMSO) were of analytical grade and purchased from
Fisher Scientific (Leicestershire, England). Deionised water was prepared using a Milli-Q system
(Millipore, Bedford, MA, USA).

#### 122 **2.3 Cell culture**

Caco-2 cells were obtained from the American Type Culture Collection HTB-37 (LGC Standards, 123 Molsheim, France). Cells (passages 30-38) were seeded at 10,000 cells/cm<sup>2</sup> in 96-well plates in 124 culture medium (minimum essential medium, supplemented with 10% fetal calf serum, 1% non-125 126 essential amino acids, 50 IU/ml penicillin and 50 µg/ml streptomycin) and maintained at 37 °C under 5% CO2. Media, serum, amino acids solution and antibiotics were all from Invitrogen (Saint Aubin, 127 France). For differentiation into an intestinal epithelial-like monolayer, cells were cultured for 3 weeks 128 129 with renewal of medium every 2 to 3 days (Sambuy et al., 2005). All the assays were performed on differentiated Caco-2 cells. 130

#### 131 2.4 Measurement of ROS production

132 ROS were measured using the probe DCFH-DA. Caco-2 cells were pre-treated with 25 µM DCFH-

133 DA for 30 min. The culture medium was then removed and medium containing the toxins was added

for 24 h of incubation at 37°C. Fluorescence was measured at  $\lambda$  ext = 485 nm and  $\lambda$  em = 520 nm using a Fluostar Omega microplate reader (BMG Labtek).

#### 136 **2.5 Cell viability and DNA strand breaks**

After 24 h of treatment with the toxins, Caco-2 cells were fixed with 4% paraformaldehyde in 137 phosphate-buffered saline (PBS) for 10 min and permeabilized with 0.2% Triton X-100 for 15 min. 138 139 Plates were then incubated in blocking solution (PBS with 1% BSA and 0.05% Tween-20) for 30 min 140 before the addition of primary antibody prepared in blocking solution and filtered with a 0.2 µm 141 syringe filter. Mouse monoclonal anti  $\gamma$ H2AX S19 (1:1000 dilution) was incubated with the cells for 1.5 h at room temperature. After washing with PBS + 0.05% Tween-20, goat pAb to Ms IgG Alexa 142 Fluor 647 (1:1000) was added for 45 min incubation at room temperature. Nuclear DAPI (1 µg/mL) 143 144 staining was used for automated cell identification by high content analysis. Plates were scanned with 145 the Thermo Scientific ArrayScan VTI HCS Reader (Thermo Scientific, Waltham, USA) and analyzed using the Target Activation module of the BioApplication software. For each well, 10 fields (10X 146 magnification) were scanned and analyzed for immunofluorescence quantification. Cell counts were 147 148 determined by DAPI staining. y-H2AX was quantified in the nuclei and expressed as fold increase compared to solvent controls. No y-H2AX data are provided for toxin concentrations resulting in a cell 149 viability < 40%, to ensure a reliable measurement of DNA damage. 150

#### 151 2.6 IL-8 release by ELISA

152 After treatment with toxins, cell culture supernatants were collected and analyzed for IL-8 release.

153 Samples were transferred to 96-well microplates coated overnight at 4°C with 1  $\mu$ g/mL human

154 recombinant anti-IL-8 primary antibody. Following addition of 0.1 µg/mL of the biotin-conjugated

human IL-8 antibody for 1 h, streptavidin peroxidase (1:10000) was added for 45 min. After washing

156 with PBS + 0.05% Tween-20, 3,3',5,5'-tetramethylbenzidine (TMB) solution was added to initiate the

- 157 colorimetric reaction. The reaction was stopped by 1 M H<sub>2</sub>SO<sub>4</sub> and absorbance was measured at 405
- nm using a Fluostar Omega microplate reader (BMG Labtek).
- 159 2.7 Statistics/Data analysis

GraphPad Prism 5 (GraphPad Software, Inc) was used for statistical analyses. Two-way ANOVA 160 161 followed by Bonferroni's post-hoc tests were used to compare the effects of toxins and solvent control All error bars denote standard error of the mean (SEM). Symbols \*, \*\*, \*\*\*, \*\*\*\* and #, ##, ###, 162 #### indicate statistical significance between toxin and solvent control or OA and mixture 163

respectively (p < 0.05, p < 0.01, p < 0.001, p < 0.0001 respectively). 164

#### 165 2.8 Mixture effects predictions using additivity models

Since the tested phycotoxins do not share any known common mechanism of action, using the CA 166 concept is therefore not relevant. Instead, the mathematical definition of additivity, also known as 167 168 theoretical additivity, response additivity or linear interaction effect, was employed (Foucquier and Guedi, 2015). The independent action concept was also employed. 169

#### 170 2.8.1 Theoretical additivity method

171 The theoretical additivity (TA) method was used as described by Weber et al., (2005). This method compares theoretical predicted values calculated from the results of each individual compound to the 172 measured values obtained with mixtures. In our study of binary phycotoxin combinations, the 173

174 predicted mixture effect values were calculated as follows:

175

Mix(A + B) predicted value = (mean value A + mean value B) – mean value solvent control 176

177 For data expressed as fold change compared to solvent control, mean value solvent control = 1 while for cell viability analysis, the mean value for the solvent control was set to 100. 178

179 Combination index (CI), a standard measure of combination effect, was calculated as indicated by 180 Foucquier and Guedj (2015):

181 
$$CI = \frac{Mix(A+B)predicted value}{Mix(A+B)measured value}$$

182 Threshold of the computed CI of CI <  $0.9, 0.9 \le$  CI  $\le 1.1$ , and CI > 1.1 were set to indicate synergism, additive effects and antagonism, respectively. In order to get cell viability CI values that respect the 183

combination effect range, ratios  $\frac{1}{CI}$  were calculated. 184

#### 2.8.2 Independent action concept 185

186 The independent action (IA) concept was used as described by Foucquier and Guedj (2015). This method established by Bliss (1939) relies on the hypothesis that a combined effect can be calculated 187 from the responses of the individual mixture components, assuming that each one acts independently 188 without interfering with the others (different sites of action). In our study, for binary phycotoxin 189 190 combinations, the predicted mixture effect values were calculated as follows:

- 191 Mix(A + B) predicted value
- = (mean value A + mean value B) (mean value  $A \times$  mean value B) 192
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The CI were calculated as indicated by Foucquier and Guedi (2015): 194

195 
$$CI = \frac{Mix(A+B)predicted value}{Mix(A+B)measured value}$$

To establish the predictions, the raw data of ROS production, IL-8 release and  $\gamma$ -H2AX 196 197 phosphorylation, originally expressed as fold change compared to the respective solvent control, were processed to obtain data suitable for IA analysis, i.e. comprised between 0 and 1. For each endpoint, 198 the condition among the phycotoxins tested individually and the binary mixture showing the highest 199 effect was attributed the value of 1 (Amax). All other conditions were then normalized to the Amax. 200 201 This data processing was described previously by Alassane-Kpembi et al. (2016) and allows assessing combined effects for endpoints other than cell viability. The raw data of cell viability were first 202 divided by 100 to get data comprised between 0 and 1. Cell viability values superior to 100 were set to 203 1. Cell viability CI values were calculated as described for TA. 204

205 **3. Results** 

#### 206 3.1 Toxic effects following OA/PTX-2 mixtures treatment in Caco-2 cells

#### 207 3.1.1 PTX-2 exacerbates the cytotoxicity of OA in Caco-2 cells

208 PTX-2 alone was not toxic for Caco-2 cells, whereas OA showed concentration-dependent toxicity 209  $(IC_{50} = 331 \text{ nM})$  (Fig. 1a). The binary OA/PTX-2 mixture induced strong cytotoxicity in Caco-2 cells, 210 with a tremendous decrease of cells number (80% for the condition D) (Fig. 1a), indicating that the 211 addition of PTX-2 increased the cytotoxicity of OA.

#### 212 3.1.2 PTX-2 decreases ROS production induced by OA

After a 24 h treatment, OA not-statistically increased ROS production while PTX-2 had no effect (Fig.

214 1b). We observed that low and intermediate concentrations of OA/PTX-2 (conditions B and C)

reduced the level of ROS production compared to that induced by OA alone (Fig. 1b).

#### 216 3.1.3 PTX-2 decreases IL-8 release induced by OA only at low concentrations

Only OA induced IL-8 release in Caco-2 cells (conditions B to E) (Fig. 1c). For low concentrations the
OA/PTX-2 mixture resulted in a reduction of IL-8 release compared to that induced by OA alone,
which was statistically significant for 37.5/12.5 nM (condition B). PTX-2 did not decrease IL-8

220 production induced by OA at higher concentrations (Fig. 1c).

#### 221 3.1.3 PTX-2 decreases y-H2AX phosphorylation induced by OA at low concentrations

PTX-2 alone slightly induced the phosphorylation of  $\gamma$ -H2AX (1.8-fold, not statistically significant) whereas OA concentration-dependently induced the phosphorylation of  $\gamma$ -H2AX even at low concentration (37.5 nM condition B, or higher) (Fig. 1d). For 37.5/12.5 nM (condition B), the mixture resulted in a lower phosphorylation of  $\gamma$ -H2AX compared to that induced by OA alone (Fig. 1d). This mixture effect is also illustrated by the immunostaining in Fig. 2. No data are provided for the three highest concentrations due to the high cytotoxicity that prevents from a reliable measurement of DNA damage.

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nM	solo		mixtures	
	OA	PTX-2	OA/PTX-2	
Α	18.75	6.25	18.75/6.25	
В	37.5	12.5	37.5/12.5	
С	75	25	75/25	
D	150	50	150/50	
Е	300	100	300/100	
F	600	200	600/200	

Fig. 1 Effects of OA/PTX-2 mixtures on a panel of toxicity endpoints in differentiated Caco-2 cells. Cells were pre-treated with the ROS probe DCFH-DA (25  $\mu$ M) for 30 min before incubation with the toxin(s) for 24 h. After fluorometric measurement of ROS production, cell culture supernatant was collected for the quantification of IL-8 release, while cells were fixed for DAPI and y-H2AX immunostaining: (a) cell viability, (b) ROS production, (c) IL-8 release and (d) phosphorylation of  $\gamma$ -H2AX are shown. KBrO<sub>3</sub> (500  $\mu$ M) and TNF $\alpha$ (20 ng/ml) were used as positive controls. KBrO<sub>3</sub> increased ROS production by 1.5-fold and induced the phosphorylation of  $\gamma$ -H2AX by 2.7-fold. TNF $\alpha$ failed to induce IL-8 release. Results were obtained from three independent experiments. Data represents means  $\pm$  SEM of fold change compared to solvent control. \*, \*\*, \*\*\* and #, ##, ### indicate statistical significance between toxin and solvent control or OA and mixture respectively (p < 0.05, p < 0.01, p <0.001 respectively) after two-way ANOVA followed by Bonferroni's post-hoc test. §: no data are provided due to high cytotoxicity that prevents from a reliable measurement of DNA damage.



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**Fig. 2** Representative images (10X magnification) of DNA damage induced by OA/PTX-2 mixtures. Cells were incubated with toxins for 24 h and labeled with antibodies against  $\gamma$ -H2AX. The nuclei were stained with DAPI and images were captured with an Arrayscan VTi. White bar = 100  $\mu$ m.

#### 279 3.2 Toxic effects following OA/SPX-1 mixtures treatment in Caco-2 cells

#### 280 3.2.1 SPX-1 reduces the cytotoxicity effect of OA in Caco-2 cells at low concentrations

SPX-1 was not toxic for Caco-2 cells in the tested range concentration. In mixture condition, a lower
toxicity was depicted for low and intermediate concentrations (conditions B and C) even if not
statistically significant (Fig. 3a).

#### 284 3.2.2 SPX-1 decreases ROS production induced by OA

Whereas SPX-1 failed to induce ROS production, we observed that low and intermediate concentrations of OA/SPX-1 (conditions A, C and D) reduced the level of ROS production compared to that induced by OA alone (Fig. 3b).

#### 288 3.2.3 SPX-1 decreases IL-8 release induced by OA only at low concentrations

SPX-1 did not induce the release of IL-8. For low concentrations the OA/SPX-1 mixture resulted in a
reduction of IL-8 release compared to that induced by OA alone, which was statistically significant for
37.5/12.5 nM (condition B). SPX-1 did not decrease IL-8 production induced by OA at higher
concentrations (Fig. 3c).

#### 293 3.2.4 SPX-1 decreases γ-H2AX phosphorylation induced by OA at low concentrations

No induction of  $\gamma$ -H2AX phosphorylation was observed after treatment with SPX-1. For 37.5/12.5 and 600/200 nM (conditions B and F), the OA/SPX-1 mixture resulted in a reduced phosphorylation of  $\gamma$ -H2AX compared to that induced by OA alone (Fig. 1d). This mixture effect is also illustrated by the immunostaining in Fig. 4.

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nM	s	olo	mixtures	
	OA	SPX-1	OA/SPX-1	
Α	18.75	2.1	18.75/2.1	
В	37.5	4.2	37.5/4.2	
С	75	8.3	75/8.3	
D	150	16.7	150/16.7	
Ε	300	33.4	300/33.4	
$\mathbf{F}$	600	66.7	600/66.7	

Fig. 3 Effects of OA/SPX-1 mixtures on a panel of toxicity endpoints in differentiated Caco-2 cells. Cells were pre-treated with the ROS probe DCFH-DA (25  $\mu$ M) for 30 min before incubation with the toxin(s) for 24 h. After fluorometric measurement of ROS production, cell culture supernatant was collected for the quantification of IL-8 release, while cells were fixed for DAPI and y-H2AX immunostaining: (a) cell viability, (b) ROS production, (c) IL-8 release and (d) phosphorylation of  $\gamma$ -H2AX are shown. KBrO<sub>3</sub> (500  $\mu$ M) and TnF $\alpha$ (20 ng/ml) were used as positive controls. KBrO3 increased ROS production by 1.5-fold and induced the phosphorylation of  $\gamma$ -H2AX by 2.7-fold. TNF $\alpha$ failed to induce IL-8 release. Results were obtained from three independent experiments. Data represents means  $\pm$  SEM of fold change compared to solvent control. \*, \*\*, \*\*\* and #, ## indicate statistical significance between toxin and solvent control or OA and mixture respectively (p < 0.05, p < 0.01, p < 0.001 respectively) after two-way ANOVA followed by Bonferroni's post-hoc test.



#### 354 3.3 Toxic effects following OA/YTX mixtures treatment in Caco-2 cells

#### 355 3.3.1 YTX reduces the cytotoxicity effect of OA in Caco-2 cells at low concentrations

YTX was not toxic for Caco-2 cells in the tested range concentration. In mixture condition, a lower
toxicity was depicted for low and intermediate concentrations (conditions A and B) even if not
statistically significant (Fig. 5a).

#### 359 3.3.2 YTX decreases ROS production induced by OA

Whereas YTX failed to induce ROS production, we observed that low and intermediate concentrations
of OA/YTX (conditions A and C) reduced the level of ROS production compared to that induced by
OA alone.

#### 363 3.3.3 YTX increases IL-8 release induced by OA at high concentrations

If YTX failed to induce IL-8 release in Caco-2 cells, the mixtures with a high concentration of YTX
showed an increase of IL-8 release compared to that induced by OA alone (conditions D and F) (Fig.
5c).

#### 367 3.3.4 YTX decreases y-H2AX phosphorylation induced by OA at low concentrations

368 No induction of  $\gamma$ -H2AX phosphorylation was observed after treatment with YTX. Similarly to what 369 was observed for the OA/SPX-1 mixture, for 37.5/12.5 and 600/200 nM (conditions B and F), the 370 OA/YTX mixture resulted in a reduced phosphorylation of  $\gamma$ -H2AX compared to that induced by OA 371 alone (Fig. 1d). This mixture effect is also illustrated by the immunostaining in Fig. 6.

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nM	solo		mixtures	
	OA	YTX	OA/YTX	
Α	18.75	6.25	18.75/6.25	
В	37.5	12.5	37.5/12.5	
С	75	25	75/25	
D	150	50	150/50	
Ε	300	100	300/100	
F	600	200	600/200	

Fig. 5 Effects of OA/YTX mixtures on a panel of toxicity endpoints in differentiated Caco-2 cells. Cells were pre-treated with the ROS probe DCFH-DA (25  $\mu$ M) for 30 min before incubation with the toxin(s) for 24 h. After fluorometric measurement of ROS production, cell culture supernatant was collected for the quantification of IL-8 release, while cells were fixed for DAPI and y-H2AX immunostaining: (a) cell viability, (b) ROS production, (c) IL-8 release and (d) phosphorylation of  $\gamma$ -H2AX are shown. KBrO<sub>3</sub> (500  $\mu$ M) and TnF $\alpha$ (20 ng/ml) were used as positive controls. KBrO3 increased ROS production by 1.5-fold and induced the phosphorylation of  $\gamma$ -H2AX by 2.7-fold. TNF $\alpha$ failed to induce IL-8 release. Results were obtained from three independent experiments. Data represents means  $\pm$  SEM of fold change compared to solvent control. \*, \*\*, \*\*\* and #, ##, ### indicate statistical significance between toxin and solvent control or OA and mixture respectively (p < 0.05, p < 0.01, p < 0.001 respectively) after two-way ANOVA followed by Bonferroni's post-hoc test.

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421 Fig. 6 Representative images (10X magnification) of DNA damage induced by OA/YTX mixtures. 422 Cells were incubated with toxins for 24 h and labeled with antibodies against  $\gamma$ -H2AX. The nuclei 423 were stained with DAPI and images were captured with an Arrayscan VTi. White bar = 100 µm.

#### 430 **3.4** Predictions of mixture effects using the theoretical additivity (TA) method and the 431 independent action (IA) concept

#### 432 3.4.1 OA/PTX-2 mixtures treatment on Caco-2 cells

433 Regarding cytotoxicity, TA model predicted additivity or antagonism for the two low concentrations while synergism occurred at higher concentrations except for the highest concentration where 434 additivity was shown (Fig. 7a). IA model predicted additivity for the two low concentrations and 435 synergism at higher concentrations. Using IA model, antagonism was observed for all concentrations 436 towards ROS production while TA predicted additivity for the two highest concentrations. For  $\gamma$ -437 438 H2AX phosphorylation, no predictions at the three highest concentrations were made due to the high cytotoxicity that prevents from a reliable measurement of DNA damage. Both models gave similar 439 440 predictions of antagonism for A and B conditions while C condition showed additivity. Antagonism 441 towards IL-8 release was obtained for low concentrations, compared to slight synergism or additivity at higher concentrations with both models. 442

#### 443 3.4.2 OA/SPX-1 mixtures treatment on Caco-2 cells

Regarding cytotoxicity, no clear tendency emerges but additivity seems to be the main effect occurring using TA model (Fig. 7b). IA model predicted additivity for the two low concentrations and synergism at higher concentrations. For IL-8 release, all kinds of effects were observed from antagonism at the lowest concentrations, to additivity or synergism at the intermediate and highest concentrations respectively. Using IA model, antagonism was observed for all concentrations towards ROS production while TA predicted additivity for the highest concentration. For  $\gamma$ -H2AX phosphorylation, both models predicted antagonism except for E condition where additivity was described.

#### 451 3.4.3 OA/YTX mixtures treatment on Caco-2 cells

452 Regarding cytotoxicity, no clear tendency emerges but additivity seems to be the main effect occurring 453 using TA model (Fig. 7c). IA model predicted additivity for the two low concentrations and synergism 454 at higher concentrations. ROS measurement and  $\gamma$ -H2AX phosphorylation showed antagonism for 455 almost all concentrations, additivity being sometimes described (e.g D condition for  $\gamma$ -H2AX 456 phosphorylation or F condition for ROS with TA). Antagonism towards IL-8 release was obtained for 457 low concentrations, compared to synergism at higher concentrations.

**Fig. 7** Analysis of mixtures using the theoretical additivity (TA) method and the independent action (IA) concept on a panel of toxicity endpoints in differentiated Caco-2 cells. CI < 0.9,  $0.9 \le CI \le 1.1$  and CI > 1.1 indicate respectively synergism, additive effect and antagonism. Data represents means  $\pm$  SEM from three independent experiments. Dashed lines indicate lower and upper limits of additivity.



#### 470 **4. Discussion (1500 words)**

In this study, we investigated the effects of binary mixtures of lipophilic phycotoxins on a variety of toxicity endpoints in a human intestinal cell model. The mixture compositions and ratios were selected according to a review of the literature on phycotoxin mixtures (Alarcan et al., 2018). Three main mixtures were tested, all of them featuring OA as the major constituent and reflecting realistic exposure scenarios. Irrespective of the phycotoxin combined with OA, the effects observed were, according to the chosen mathematical approach, mainly antagonistic at low concentrations and, in a few cases, synergistic at higher concentrations.

When investigating mixture effects, the choice of endpoints is a main concern. A specific endpoint can be easily selected when the compounds share a common mechanism of action. In our case, the toxins tested have different mechanisms of action, and therefore a single relevant common endpoint can hardly be chosen. Therefore, we designed a multi-parametric approach for an integrative analysis of toxic responses. To this end, we selected different endpoints among widely studied toxicological pathways.

484 Our study revealed that OA was cytotoxic, induced DNA strand-breaks and triggered the release of IL-8 from differentiated Caco-2 cells, similar to the effects reported with proliferative Caco-2 cells by 485 486 Ferron et al. (2014). These effects correspond to observations from in vivo studies in rodents that 487 reported erosion of the epithelium and release of inflammatory mediators following oral exposure to OA (Aune et al., 2012, Tubaro et al., 2003, Ito et al., 2002). PTX-2 did not affect Caco-2 cell viability 488 and only slightly induced DNA strand-breaks at the highest concentrations. SPX-1 and YTX did not 489 490 induce any effect on all tested endpoints. The absence of toxic effects correlates well with in vivo 491 studies in rodents that did not report particular effects on the gastro-intestinal tract following oral 492 exposure to these toxins (Aune et al., 2002, Tubaro et al., 2003, Munday et al., 2012). Irrespective of 493 the phycotoxin, no ROS production was detected. However we cannot exclude that ROS might have 494 been produced earlier and were already neutralized after 24h due to protective cellular anti-oxidant 495 systems

Two different conceptual approaches were employed to assess the combined effects of phycotoxins in 496 497 mixture. The TA method and IA concept offer the advantage to be easy to handle, fast and applicable 498 to all kind of data. We showed that under the present settings and parameters additivity was rarely predicted irrespective of the mixture and the endpoint tested, meaning that the phycotoxins tested 499 might be able to interact and modulate their toxic effects. Therefore, in absence of additivity, the two 500 501 approaches are not appropriate to predict the toxicity of phycotoxin mixtures. In this case, the combined effects of phycotoxins must be determined from experimental data. Besides, caution needs 502 to be taken with mixture simulations. For instance, in our study, IA analysis revealed strong synergism 503 504 towards cell viability for OA/SPX-1 and OA/YTX (Figure 7) whereas raw data revealed only a slight 505 or no increase of toxicity with mixtures (Figures 3a and 5a). It is noteworthy that a general assumption of additivity may not be reliable since OA is believed to target different serine/threonine protein 506 phosphatases (PP) depending on its concentration: the estimated  $IC_{50}$  for PP1 is 10-100 nM, while 507 being 100-fold inferior for PP2A (0.1-1 nM) (Cruz et al., 2013). Indeed, Gehringer (2004) suggested 508 509 that the PP inhibitors microcystin-LR and OA showed a dualistic response by inducing different 510 cellular effects depending on the concentration used, and therefore on the PP isoform inhibited.

511 Since few years, the Chou-Talalay method (Chou and Talalay 1984) has received considerable interest 512 in the context of food contaminant mixtures (Alassane-Kpembi et al., 2016, Ferron et al., 2016, Smith 513 et al., 2017, Karri et al., 2018). This easy-to-use method is a powerful tool which offers a quantitative 514 definition for additivity and the type of interaction (synergism or antagonism) in chemical mixtures.

- 515 Based on the median-effect equation, it requires that each compound alone and in mixtures follows the
- 516 principle of the mass-action law (e.g., r > 0.95). Therefore the method is not suitable when one drug 517 does not exert any effect (Chou, 2006). In our study, most endpoints were not affected by at least one
- 518 of the toxins, and consequently the Chou-Talalay method would not be appropriate.
- 519 Our study revealed that only OA provoked IL-8 release. However, we observed a greater effect at high
- 520 concentrations with the OA/YTX mixture, meaning that YTX may potentiate the OA response. A 521 similar hypothesis can be deduced for the cytotoxicity of OA/PTX-2 mixture. However, confirmation
- 522 of such observations through a mathematical model was not possible since none exists for testing
- 523 potentiation.
- Irrespective of the toxin, mixtures with OA always resulted in a diminished toxic response at low 524 525 concentrations (ROS production, IL-8 release and H2AX phosphorylation), whereas a higher toxic response was observed for some endpoints at high concentrations. A diminished ROS production 526 527 might be linked to a diminished DNA damage response, as it is known that ROS can cause DNA breaks (Moloney and Cotter 2018). Moreover, many studies have shown that the inhibition of ROS by 528 529 the use of a scavenger leads to a lowered DNA damage response (Luo et al., 2013, He et al., 2014, 530 Chen et al., 2018). ROS have been shown to activate NF-κB, a transcription factor responsible for IL-8 531 regulation (Kurata et al., 1996, Pahl 1999, Oeckinghaus and Ghosh 2009). Since all these endpoints are interconnected, the systematic decrease of responses that was observed with mixtures at low 532 533 concentrations in our experiments appears consistent. The protective mechanisms behind this might 534 relate to changes in the absorption and/or the metabolism of toxins in the mixtures. For instance, OA was previously found to interact with regulatory nuclear receptors such as PXR (Fidler et al., 2012; 535 Ferron et al., 2016), which regulate the expression of some cytochrome P450 enzymes (Wang et al., 536 537 2012). PTX-2 is believed to interact with the AhR since it induces CYP1A protein in hepatic cells 538 (Alarcan et al., 2017). Moreover, OA, PTX-2 and SPX-1 have been shown to be metabolized through 539 cytochromes P450 (Guo et al., 2010, Hui et al., 2012, Kittler et al., 2014, Alarcan et al., 2017). 540 Therefore, at low concentrations, mixtures might induce CYP activity and efflux transporter 541 expression, resulting in higher detoxification/excretion of toxins and thus decreased toxic effects. In 542 favor of this hypothesis, the few oral *in vivo* studies conducted with phycotoxin mixtures showed a reduced uptake of toxins (Alarcan et al., 2018). As PTX-2, YTX and SPX-1 have different 543 mechanisms of action, we assume that the potential protective effect, as suggested by the mathematical 544 545 evaluation of our in vitro data, did not result from the disturbance of a single specific OA-dependent pathway. The strong synergistic effect observed for the cytotoxicity of the OA/PTX-2 mixture at 546 547 higher concentrations may imply cytoskeleton homeostasis. Indeed, even if no cytotoxicity following 548 treatment with PTX-2 has been observed, the toxin clearly disrupts the actin network (Alligham et al., 549 2007). OA has also been demonstrated to change the organization of the actin network (Fiorentini et 550 al., 1996; Vale and Botana 2008). Disruption of the cytoskeleton has been shown to be a starting event leading to an induction of pro-apoptotic effectors (e.g caspase-3 activation) (Yamazaki et al., 2000, 551 552 White et al., 2001, Martin and Leder 2001, Kulms et al., 2002). Therefore, the disruption of the actin 553 cytoskeleton by both, PTX-2 and OA, may explain the possible synergism observed with the mixture.
- Toxins of the OA group and of the PTX group share a common European regulatory limit but their behavior in mixtures is largely unknown. Results from this study suggest a possible synergistic cytotoxic effect at moderate and high concentrations in Caco-2 cells, a model mimicking the human intestinal barrier. Further research is needed to confirm the present findings and to judge on their relevance for humans exposed to phycotoxins via the diet.
- 559 **5.** Conclusions

- In this study, we assessed the *in vitro* effects of binary mixtures of lipophilic phycotoxins on human intestinal cells towards multiple toxicity endpoints. Mainly antagonistic effects were depicted at low concentrations and, in few cases, a strong synergism was detected at higher concentrations. These
- 563 findings need to be confirmed with additional experiments and completed with *in vivo* studies to bring
- 564 further evidence on the combined effects of lipophilic phycotoxins.
- 565 **Declarations of interest:** none

#### 566 Authorship Contributions

- 567 Participated in research design: Alarcan, Le Hégarat, Fessard, Hessel-Pras, Lampen
- 568 Conducted experiments: Alarcan, Barbé
- 569 Contributed new reagents or analytical tools:
- 570 Performed data analysis: Alarcan
- 571 Wrote or contributed to the writing of the manuscript: Alarcan, Le Hégarat, Kopp, Hessel-Pras,572 Braeuning, Lampen and Fessard

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