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Molecular impacts of dietary exposure to nanoplastics combined with arsenic in Canadian oysters (*Crassostrea virginica*) and bioaccumulation comparison with Caribbean oysters (*Isognomon alatus*)

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PII: S0045-6535(21)00801-8

DOI: <https://doi.org/10.1016/j.chemosphere.2021.130331>

Reference: CHEM 130331

To appear in: *ECSN*

Received Date: 21 January 2021

Revised Date: 12 March 2021

Accepted Date: 16 March 2021

Please cite this article as: Lebordais, M., Gutierrez-Villagomez, J.M., Gigault, J., Baudrimont, M., Langlois, V.S., Molecular impacts of dietary exposure to nanoplastics combined with arsenic in Canadian oysters (*Crassostrea virginica*) and bioaccumulation comparison with Caribbean oysters (*Isognomon alatus*), *Chemosphere*, <https://doi.org/10.1016/j.chemosphere.2021.130331>.

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Credit Author Statement

Marc Lebordais: methodology, investigation, formal analysis, writing - original draft. **Juan Manuel G. Villagomez:** methodology, investigation, writing - review & editing. **Julien Gigault:** funding acquisition, resources, writing - review & editing. **Magalie Baudrimont:** funding acquisition, conceptualization, supervision, writing - review & editing. **Valerie S. Langlois:** funding acquisition, project administration, conceptualization, resources, supervision, writing - review & editing.

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1 **Molecular impacts of dietary exposure to nanoplastics combined with**
2 **arsenic in Canadian oysters (*Crassostrea virginica*) and bioaccumulation**
3 **comparison with Caribbean oysters (*Isognomon alatus*).**

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1 **Graphical Abstract**

2

48 h exposure

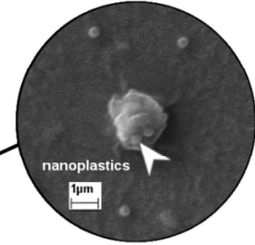


Nanoplastics
+ microalgae



1 week exposure

Crassostrea virginica



Tisochrysis lutea

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1 Abstract

2

3 Despite the urge need to address the possible impact of plastic debris, up to now,
4 little is known about the translocation of nanoplastics through the trophic web. Plus,
5 due to their surface reactivity, nanoplastics could sorb and thus increase metals
6 bioavailability to aquatic filter-feeding organisms (e.g., bivalves). In this study we
7 investigated the dietary exposure route on the oyster *Crassostrea virginica* through
8 microalgae themselves exposed to three nanoplastic dispersions (PSL, PSC and
9 NPG) at reportedly environmental concentrations combined or not with arsenic.

10 Interactive effects of nanoplastics on arsenic bioaccumulation were studied, along
11 with the expression of key genes in gills and visceral mass. The investigated gene
12 functions were endocytosis (*cltc*), oxidative stress (*gapdh*, *sod3*, *cat*), mitochondrial
13 metabolism (*12S*), cell cycle regulation (*gadd45*, *p53*), apoptosis (*bax*, *bcl-2*),
14 detoxification (*cyp1A*, *mdr*, *mt*), and energy storage (*vit*). Results showcased that
15 nanoplastic treatments combined with arsenic triggered synergetic effects on gene
16 expressions. Relative mRNA level of *12S* significantly increased at 10 and 100 $\mu\text{g L}^{-1}$
17 for NPG combined with arsenic and for PSC combined with arsenic. Relative mRNA
18 level of *bax* increased for PSL combined with arsenic and for PSC combined with
19 arsenic at 10 and 100 $\mu\text{g L}^{-1}$ respectively. We also observed that relative arsenic
20 bioaccumulation was significantly higher in *Crassostrea virginica* gills compared to
21 *Isognomon alatus*'. These results are the first comparative molecular effects of
22 nanoplastics alone and combined with arsenic investigated in farmed *C. virginica*
23 oysters. Together with *I. alatus* results we thus shed light on species different
24 sensitivity.

25

26 **Keywords** : nanoparticles, toxicity, bivalves, gene expression, bioaccumulation,
27 Scanning Electron Microscopy

28 1. Introduction

29 Plastic contamination is of global concern (UNEP, 2001; Bank and Hasson, 2019)
30 and approximately 10% of the annual worldwide plastic production ends up in the
31 oceans (Mattsson *et al.*, 2019). Current estimates indicate that there are 30 million
32 tons of plastic in the oceans, and according to projections by 2025, there will be 220
33 million tons (Wright and Kelly, 2017). Plastics have become ubiquitous in aquatic
34 ecosystems to the point where they reached the Arctic polar circle (Lusher *et al.*,
35 2015) and the Mariana trench (Koelmans *et al.*, 2015; Chiba *et al.*, 2018). Plastic
36 waste is persistent in the environment (Carpenter and Smith, 1972) and can last
37 hundreds of years without being completely degraded (Barnes *et al.*, 2009; Briand
38 2014).

39 Physical abrasion, chemical, and photo-oxidations (e.g., waves, salt, UV; Gigault
40 *et al.*, 2018a; Magrí *et al.*, 2018) can lead to plastic fragmentations resulting in the
41 formation of microplastics (MPs) and nanoplastics (NPs). MPs are synthesized or
42 broken plastic pieces smaller than 5 mm (Arthur *et al.*, 2009), while NPs are the
43 colloidal fraction of the plastic debris (Lambert and Wagner, 2016; Gigault *et al.*,
44 2018b). Primary NPs refer to those coming from industrial synthesis and secondary
45 NPs are those resulting from environmental degradation (Koelmans *et al.*, 2015).
46 Both, MPs and NPs, including primary and secondary, are emerging contaminants
47 (Richardson and Kimura, 2020) due to the limited knowledge regarding their
48 hazardous structural properties and long-term fate in living beings. The toxicity of
49 MPs and NPs raises concern globally since most marine species ingest plastic
50 regularly (Gall and Thompson, 2015; Lusher *et al.*, 2017).

51 Recent evidences showed that organisms could excrete most of the ingested
52 MPs; therefore, minimal translocation into the circulatory system and tissues is
53 expected (Lusher *et al.*, 2013; Sussarellu *et al.*, 2016; Santana *et al.* 2018). However,
54 as plastics age, they lose their physical integrity and their additives (e.g., plasticizers,
55 UV-filters, flame-retardants, metals) and become less chemically stable (Lambert
56 and Wagner, 2016; Wright and Kelly, 2017). Also, weathered and nanofragmentated
57 plastics present highly reactive surface leading to adsorb numerous contaminants
58 including metals and metalloids (Ashton *et al.*, 2010; Tien and Chen, 2013; Rochman
59 *et al.*, 2014; Davranche *et al.*, 2019). For example, El Hadri *et al.* (2020a) revealed

60 that arsenic (As) was one of the most abundant metal adsorbed on plastic debris
61 sampled on Guadeloupean beaches (a French island in the Caribbean Sea).

62 Despite recent improvement of NP technical analyses, data on NPs' fate and
63 toxicity is limited (Quik *et al.*, 2011; Ter Halle *et al.*, 2017; Nguyen *et al.*, 2019). They
64 are the least understood plastic fraction, potentially the most toxic to biologic
65 systems by tissue bioaccumulation and vectoring chemical contamination (Nel *et al.*,
66 2006; Mattsson *et al.*, 2015; Chae and An, 2017). Riverine systems are the major
67 input of plastic debris to the open ocean (Mayer and Wells, 2012; Lasareva *et al.*,
68 2017). Therefore, since estuaries link rivers to oceans they are the key interface
69 controlling the fate, transport and accumulation of plastics debris, more specifically
70 for NPs (Galgani *et al.*, 2000; Browne *et al.*, 2010; Sadri and Thompson 2014).
71 Indeed, the change in estuarine systems of ionic strength and natural organic matter
72 (NOM) will control the aggregation of NPs. Aggregation is the main parameter
73 affecting NPs final reactivity in regard to other contaminants. In such systems,
74 mangroves are a specific category where these parameters are intensified (Bouillon,
75 2011). Moreover, mangroves are essential ecosystems weakened by anthropogenic
76 pressure, thus they are relatively used as proxy for worldwide environmental issues
77 (Mitra, 2013; Carugati *et al.*, 2018; Kulkarni *et al.*, 2018).

78 Bivalves are a class of marine and freshwater molluscs. They include oysters,
79 clams, mussels, and most of them are filter feeders (Hancock *et al.*, 2008;
80 Baudrimont *et al.*, 2019) also they have been used historically as biomonitoring
81 species to assess exposure to metals (Aguirre-Rubí *et al.*, 2017; Kulkarni *et al.*,
82 2018). However, recently, plastic exposure to aquaculture bivalves raised concerns
83 about NPs transfer within the food web (Baun *et al.*, 2008; Bouwmeester *et al.*, 2015;
84 Bank and Hansson 2019). Therefore we conducted a trophic exposure using the
85 Eastern oyster (*Crassostrea virginica*) to assess the potentially toxic effects of three
86 distinct NPs and in combination with As. The oyster *C. virginica* is one of the most
87 extensively farmed oyster species in the world. Due to its habitat tolerances ranging
88 from the Gulf of Mexico to Canada's eastern bays (Ozbay *et al.*, 2014), this species
89 is highly commercially valuable and is also commonly used in toxicity assays
90 (McCarthy *et al.*, 2013; Ward *et al.*, 2019; Smith *et al.*, 2020). We used three NP
91 models following a gradient of environmental relevancy : (i) carboxylated polystyrene
92 nanoparticles of latex (PSL) free of additives as it can induce toxicity (Pikuda *et al.*,

93 2019), (ii) the crushed pristine polystyrene nanoparticles (PSC) that was considered
94 more relevant in size and shape distribution, and finally (iii) nanoplastics
95 environmentally weathered from microplastic debris collected in Guadeloupe. We
96 analyzed *C. virginica* As bioaccumulation and gene expressions in gills and visceral
97 mass tissues after NPs trophic exposure. Then, the As bioaccumulation factor was
98 compared between *C. virginica* and widely caught flat tree oysters (*Isognomon*
99 *alatus*) that had been previously studied (Lebordais *et al.*, *submitted*). The tropical *I.*
100 *alatus* oyster is native to Caribbean mangroves that are exposed to the North
101 Atlantic gyre where plastic debris concentrate (7th expedition continent, Baudrimont
102 *et al.*, 2019).

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104 2. Materials and methods

105 2.1 Preparation of nanoplastic dispersions and size characterisation

106 In this study, we used three different NP dispersions : (i) spherical carboxylated
107 polystyrene nanoparticles of latex (PSL) synthesized according to a previous
108 protocol (Pessoni *et al.*, 2019) free of additives; (ii) the crushed polystyrene
109 nanoparticles (PSC) that were in-laboratory nanofragmented from polystyrene
110 pristine pellets by ball-milling; (iii) and a mixture of nanoplastics produced from
111 microplastic debris environmentally weathered and collected from a beach in
112 Guadeloupe (NPG). The PSC and NPG plastics pellets were degraded with 99%
113 ethanol in a blade grinder to get a primary powder and later fragmented using a
114 planetary ball mill (El Hadri *et al.*, 2020b). The resultant powder was then dried by
115 lyophilization to remove ethanol, then suspended in deionized water (MilliQ, Millipore,
116 18 MΩcm) and filtered on cellulose acetate filters (5-6 μm pore size, VWR).

117 To assess NP hydrodynamic diameters and population distributions, dynamic
118 light scattering (DLS) was used. Hydrodynamic diameters of PSC and NPG
119 dispersions were measured using non-invasive backscatter optics at 25 °C on a
120 Zetasizer nano zs instrument (Malvern Panalytical®). The intensity fluctuations by the
121 time were processed as a correlation function. A cumulants algorithm was used to fit
122 this function to obtain a size distribution (z-average) and the polydispersity index
123 (PDI). The z-averages were 692.4 ± 68.55 nm (0.441 PDI) and 1071 ± 30.65 nm
124 (0.755 PDI) for PSC and NPG, respectively. The calibrated z-average of PSL was
125 390 ± 20 nm (0,002 PDI). The three NPs were suspended in deionized water. To
126 measure NP mass concentrations in the three NP dispersions, analyses were
127 performed by total organic carbon (TOC) on a Shimadzu® instrument with a
128 detection limit at 0.05 mg.L^{-1} . Spike recovery was performed (yield = 92%) to validate
129 the analysis accuracy.

130

131 2.2 Scanning electron microscopy observation

132 To observe the potential interaction between NPs and microalgae surface, we
133 exposed the microalgae *Tisochrysis lutea* (formerly known as *Isochrysis galbana*;
134 Bendif *et al.*, 2014) for 48 h to PSL dispersions at 0 (control), 10, 100, 1000, and

135 5000 $\mu\text{g L}^{-1}$ and observed them using Scanning Electron Microscopy (SEM). For this,
136 *T. lutea* was purchased from Bigelow National Center for Marine Algae and
137 Microbiota (Maine, USA). The micro algae were aliquoted into previously cleaned
138 glass vials (one per treatment, $n = 1$) and acclimatized for one week at room
139 temperature under 24:24 artificial light. All microalgae solutions yielded the same
140 concentration (2.1×10^6 cells/mL) measured on a Coulter Multisizer II particle
141 counter (Fortin *et al.*, 2000) before starting the exposure. Right after the 48 h
142 exposure, *T. lutea* solutions were fixed in 3% glutaraldehyde (Fisher Scientific[®],
143 United Kingdom) for 1 h (Bergami *et al.*, 2017). Samples were not previously
144 centrifuged as we suspected PSL precipitation could cover microalgae cell walls
145 without adsorption interactions. Micro volumes of fixed *T. lutea* were thus dried at
146 room temperature for 30 min in a laminar hood. Later, the samples were gold
147 sprayed and observed under SEM (Carl Zeiss EVO[®] 50) at 7 kV. All the glassware
148 used in the experiments was cleaned using an acid bath of 3% nitric acid, rinsed with
149 distilled water then with 70% ethanol and dried under a fume hood.

150

151 2.3 Microalgae and oyster cultures

152 Marine microalgae species *Chaetoceros calcitrans* and *Tisochrysis lutea* were
153 obtained from Fisheries & Oceans (Tracadie-Sheila, New Brunswick, Canada).
154 *Chaetoceros calcitrans* and *T. lutea* were cultured in glass balloons with F/2 medium
155 (Guillard, 1975) at 26‰ salinity, and acclimatized at room temperature under 24:24
156 artificial light of 73 $\mu\text{mol/m}^2/\text{s}$.

157 Individuals of *C. virginica* were acquired at the beginning of fall before the
158 hibernation stage from New Brunswick's raised oysters (Canada). They were
159 selected to have a diameter range from 6.4 to 7.6 cm, meaning all oysters were
160 between 3 to 4 years old. Once brought to the laboratory, they were individually
161 brushed to remove external parasites and placed in 25 L tanks (39 oysters per tank).
162 Tanks contained reconstituted seawater (Instant Ocean[®]) at 30‰ salinity,
163 oxygenated, and filtered by an aquarium filter pump. Similar to *C. virginica* farming
164 conditions, they were acclimatized at 20 °C (Ward *et al.*, 2019) with aquarium
165 heaters and under 12:12 (light:dark cycles) for nine weeks. During acclimation, the
166 oysters were fed twice a week with a mixt *T. lutea* and *C. calcitrans* (2.5×10^6 cells/L

167 and 1.7×10^6 cells/L, per tank respectively). The algae *C. calcitrans* was used only
168 during oysters acclimation for nutritional purposes (Gonzalez Araya *et al.*, 2012).

169

170 2.4 Trophic exposure

171 The microalgae *T. lutea* was used as NPs vector for *C. virginica* dietary exposition to
172 PSL, PSC, and NPG. To that end, the three NP dispersions were separately added
173 into *T. lutea* solutions for 48 h at nominal concentrations of 10 and 100 $\mu\text{g L}^{-1}$ NPs,
174 presumed to be environmentally low (Lenz *et al.*, 2016; Besseling *et al.*, 2014). This
175 exposure time was chosen according to previous experiments and proven to be
176 effective (Lebordais *et al.*, *submitted*). Before NP dosing, the microalgae
177 concentrations were assessed by a Coulter Multisizer II particle counter (Fortin *et al.*,
178 2000) and brought to 4.7×10^6 cells/mL. NP-*T. lutea* solutions were under the same
179 abiotic conditions as during the acclimation phase. All the *C. virginica* individuals
180 were fed every two days with 2.24×10^3 cells/oyster/L of *T. lutea*. The oysters
181 belonging to NP treatments were fed with NP-*T. lutea* at the same microalgae
182 concentration.

183 To study the complexity of NP interactions with environmental metallic
184 contaminants, we combined each NP treatment with As. Arsenate (pentoxide arsenic)
185 is the most abundant inorganic form of As found in oxygenated marine waters and
186 also the least toxic one (Francesconi and Edmonds, 1996; Neff, 1997; Zhang *et al.*,
187 2013). Therefore, a solution of dissolved arsenate (hereafter referred to As treatment)
188 was prepared to use during the exposure (US EPA, 2001). On day one, the oysters
189 treated with As were exposed to a nominal concentration of 1 mg L^{-1} in water (Zhang
190 *et al.*, 2015). The As concentration was chosen based on previous studies (Langston,
191 1984; Zhang *et al.*, 2015) to compensate the oysters' natural chronic exposure.
192 Water samples were collected one day out of two and the total As was measured. If
193 needed, the As level was adjusted to keep the 1 mg L^{-1} concentration throughout the
194 experiment.

195 The experimental design included twelve NP treatments, a negative control
196 (reconstituted seawater) and a positive control (As at 1 mg L^{-1}). Thus, the experiment
197 encompassed three single-NP treatments (NPG, PSC, PSL) at both 10 and 100 μg
198 L^{-1} and three combined-NP treatments (also at 10 and 100 $\mu\text{g L}^{-1}$ for each NPs) with

199 1 mg L⁻¹ of As. One litre jars were filled up to 500 mL with reconstituted seawater
 200 and parafilm-covered to lower evaporative loss. Air distribution pumps were set up
 201 for water oxygenation. To avoid plastics contact, silicone tubing with glass pipette
 202 tips were used in each jar. Salinity, temperature, and light were the same as during
 203 acclimation. All treatments were conducted in five independent replicates with one
 204 oyster per glass jar. Oysters were dissected at the end of the one-week diet
 205 exposure. Biometric data are reported in supplementary file **Fig. S1**. Shell length
 206 was assessed on individual oyster pictures by ImageJ software. Whole-body were
 207 manually dried with paper then weighed (fresh weight). Gills and visceral mass
 208 tissues were quickly collected and stored at -80 °C for As dosage and subsequent
 209 molecular assays. The condition index (CI; Lucas and Beninger, 1985) was
 210 calculated by the following equation.

211

212 Equation 1 :

213

$$CI = \frac{\text{leftover tissues} * \text{weight}}{\text{shells weight}} \times 100 \quad * \text{leftover tissues: whole body excluding gills and visceral mass}$$

214

215 2.5 Arsenic quantification in water by ICP-OES and in oyster tissues by ICP-MS

216 Total As_(water) concentration was monitored throughout the one-week exposure.
 217 Water samples of 10 mL per jar were collected for control and As treated oysters.
 218 Then, the water samples were acidified with 3% nitric acid for inductively coupled
 219 plasma optical emission spectrometry (ICP-OES) analysis with a detection limit for
 220 As at 0.005 mg L⁻¹.

221 Total As_(tissues) concentration was measured in gills and visceral mass for *C.*
 222 *virginica*. First dried at 50 °C for 48 h, tissues were then weighed before digestion.
 223 Tissue samples were acidified with 70% nitric acid (3 mL per sample) and heated at
 224 100 °C for 3 h. After dilution of the digestates with deionized water (1:36), the final
 225 volume was 7 mL per sample. Concentrations of total As_(tissues) were measured using
 226 an inductively coupled plasma mass spectrometry (ICP-MS) with a detection limit for
 227 As at 0.02 µg L⁻¹. A standard curve of an As reference solution (SCP Science[®],
 228 Multi-Element Std) was systematically analyzed with the samples to control the

229 measured concentrations. The single spike recovery method was performed (yield =
230 101.71%) to validate the analysis accuracy (Wolle and Conklin, 2018).

231 2.6 Comparison of As bioaccumulation factor

232 In previous work, wild-caught *Isognomon alatus* oysters native to the Caribbean Sea
233 were collected from Guadeloupean mangroves (Lebordais *et al.*, *submitted*).
234 *Isognomon alatus* underwent similar acclimation and exposure conditions with
235 *Crassostrea virginica*. These consistent parameters allow us to compare the As
236 uptake for each oyster species. Therefore, the potential differences in the As
237 bioaccumulation between *I. alatus* and *C. virginica* should be mainly due to their
238 physiological and natural background differences (Moreira *et al.*, 2018). To compare
239 As bioaccumulation levels between both oyster species, we normalized the
240 concentration from exposed individuals to controls and presented it for gills and
241 visceral mass.

242 The bioaccumulation factor (BAF) of As was calculated for *C. virginica* and *I.*
243 *alatus* oysters using Equation 2 (n = 4 per treatment and species). Wet weights were
244 calculated from the measured dry weights with 0.2 and 0.1 corrections factor for gills
245 and visceral mass, respectively (Klinck *et al.*, 1992; Choi *et al.*, 1993; Kobayashi *et*
246 *al.*, 1997). The As concentrations measured in exposed oysters were normalized
247 with natural As background from control oysters. These relative As concentrations in
248 gills and visceral mass were then averaged to estimate oyster's whole body content.
249 As concentrations in water were measured for each oyster jars.

250

251 Equation 2 :
252

$$\text{BAF} = \frac{[\text{As}]_{\text{wet tissue}}}{[\text{As}]_{\text{water}}} \quad \begin{array}{l} [\text{As}]_{\text{wet tissue}} : \text{in } \mu\text{g/kg} \\ [\text{As}]_{\text{water}} : \text{in } \mu\text{g/L} \end{array}$$

253

254 2.7 RNA extraction and cDNA synthesis

255 A similar amount of tissues were individually homogenized with one stainless steel
256 ball (5 mm) per sample in 600 μL of RNA lysis buffer. To that end, a Retsch[®] mixer
257 mill MM 400 (Fisher Scientific[®], Toronto, Canada) was used for 4 min at 20 Hz.

258 Oyster tissues are rich in fat and proteins and to separate these components, 500 μ L
259 of Phenol:Chloroform:Isoamyl Alcohol (25:24:1 v/v/v, Sigma-Aldrich[®], Oakville,
260 Canada) were added and vortexed before RNA extraction. This organic solvent is
261 indeed highly suitable for insoluble tissue extractions (Vicient and Delseny, 1999). To
262 separate the aqueous phase that contained the RNA, the samples were centrifuged
263 for 1 min at 13,000 rpm. From there, total RNA was extracted using the Quick-
264 RNA[™] Miniprep Kit (Zymo Research[®]) with on-column DNAase I treatment as
265 described in the manufacturer's protocol. The total RNA concentration was
266 measured using a NanoDrop-2000 spectrophotometer (Thermo Fisher Scientific[®]).
267 The nucleic acid purity was ensured by ratios 260/280 > 1.8 and the RNA integrity of
268 each sample was assessed on a 1% agarose gel. Generally, in eukaryotes the RNA
269 integrity is assessed by the presence of two defined bands representing the 28S and
270 18S ribosomal RNA (rRNA) (Gutierrez-Villagomez *et al.*, 2019). In the RNA integrity
271 analysis, a single band with no smear was observed in the gels. This is possible due
272 to a "hidden break" in the 28S rRNA of *C. virginica* after heat denaturation.
273 Winnebeck *et al.* (2010) documented the 28S rRNA splits into two fragments that
274 overlap with the 18S rRNA during a gel electrophoresis.

275 All the samples were diluted to obtain the same RNA concentration (2000 ng in 8
276 μ L). Complementary DNA (cDNA) was prepared using Maxima[™] H Minus First
277 Strand cDNA Synthesis Kit with dsDNase (Thermo Fisher Scientific[®]). A first step of
278 DNA denaturation was conducted with dsDNase according to the supplier's
279 instructions. Then 4 μ L of mix including the reverse transcriptase enzyme, buffer and
280 primers were added with 6 μ L of water per sample. This second step was performed
281 on a Mastercycler Thermocycler Pro S (ThermoFisher, Ottawa, Canada) following
282 the supplier's protocol. All cDNA was synthesized at the same time for each tissue
283 sample including a no reverse transcriptase control (NRT) and a no template control
284 (NTC). Then the cDNA samples were stored at -20 °C.

285

286 2.8 *qPCR assays and validations*

287 Quantitative polymerase chain reaction (qPCR) was performed to measure relative
288 messenger RNA (mRNA) levels in the following 15 genes : clathrin heavy chain (*cltc*)
289 to evaluate endocytosis; catalase (*cat*), glyceraldehyde-3-phosphate-
290 deshydrogenase (*gapdh*) and superoxide dismutase Cu/Zn extracellular (*sod3*) to

291 assess oxidative stress; mitochondrial encoded 12S rRNA (12S) to measure
292 mitochondrial metabolism; growth arrest DNA damage (*gadd45*) and tumor protein
293 P53 (*p53*) to measure cell cycle regulation; bcl-2-associated X apoptosis regulator
294 (*bax*) and apoptosis regulator (*bcl-2*) to assess apoptosis; cytochrome P450 family 1
295 sub-family A1 (*cyp1A*), ATP binding cassette sub-family B1 (*mdr*) and
296 metallothionein (*mt*) to assess detoxification; and vitellogenin (*vit*) to evaluate the
297 energy storage. Elongation factor 1 alpha (*ef1a*) and ribosomal protein L7 (*rpl7*) were
298 included as reference genes (Lee and Nam, 2016). Specific primer sets were
299 designed for all genes with primer-BLAST on the NCBI platform (supplementary file
300 **Table S1**) and synthesized by Sigma-Aldrich[®]. The complete details concerning the
301 qPCR analysis is summarized in the Supplementary materials.

302

303 2.9 Statistical analyses

304 Analysis of outliers for biometric data, relative mRNA levels and As tissue
305 concentrations were performed using the ROUT method (Q = 1%) in GraphPad
306 Prism 8.0 (Motulsky and Brown, 2006). Raw data were then transformed by
307 commonly used functions (square root, double square root or decimal logarithm) to
308 satisfy parametric conditions in SigmaPlot 12.0. Normality was thus confirmed using
309 Shapiro-Wilk test and homoscedasticity was confirmed by Levene test. Transformed
310 data were compared using a two-way analysis of variance (ANOVA) for biometric
311 data and relative mRNA levels. As bioaccumulation transformed data were only
312 compared using a one-way ANOVA or a Student's t-test. For all data, the significant
313 differences were identified by a post-hoc Tukey HSD test on Prism 8.0. The
314 significance level was set at $\alpha = 0.05$.

315

316 3. Results and Discussion

317 3.1 Bottom-up complexity of nanoplastic dispersions

318 We selected monodispersed PSL nanoparticles free of additives to exclude
319 chemicals toxicity. Also, PSL were functionalized with carboxylated groups making
320 them stable and optimal for comparing results with literature in regard to impact
321 determination of NPs (Kim *et al.*, 2017; Thiagarajan *et al.*, 2019). Previous works
322 already described these PSL (Pessoni *et al.*, 2019; Lebordais *et al.*, *submitted*),
323 hence NPs characterization was focused on PSC and NPG. The PSC nanoparticles

324 were relevant NPs as they have been laboratory nanofragmented from large plastic
325 pellets (El Hadri *et al.*, 2020b). As a result, they were polydispersed and covering the
326 global colloidal size distribution (**Fig. 1 A-B**). This parameter is interesting since NPs
327 polydispersity is poorly addressed in ecotoxicological studies (Alimi *et al.*, 2018;
328 Bhagat *et al.*, 2020). Moreover, PSC had higher specific surface area and irregular
329 shapes increasing NP adsorption ability compared to PSL (Quik *et al.*, 2011;
330 Brennecke *et al.*, 2016). However, both PSL and PSC lacked natural aging and
331 exposure to contaminants. To go one step further in the environmental relevancy,
332 the collected Guadeloupean plastics have been naturally exposed to contaminants
333 and weathered by abiotic processes (such as UV light) which increase surface
334 oxidation (Holmes *et al.*, 2014; Andrady, 2017; Dawson *et al.*, 2018; El Hadri *et al.*,
335 2020b; Mao *et al.*, 2020). Therefore, NPG resulting from plastic debris was most
336 likely to reproduce the heterogeneity of NPs observed in the environment. As such,
337 NPG could include several plastic polymers (e.g., polyethylene, polypropylene,
338 polyvinyl chloride and polystyrene; Gigault *et al.*, 2016; Davranche *et al.*, 2020).

339 The NPG plastics mixture may also contain a wide variability of additives like
340 pigments (Frias *et al.*, 2010). Noteworthy, pigments absorbance have been
341 previously documented to bias the measures of light dispersion (Zook *et al.*, 2011;
342 Geißler *et al.*, 2015). As NPG are detected by dynamic light scattering, they present
343 a Brownian motion in aqueous system and therefore can be defined as NPs (Gigault
344 *et al.*, 2018b). Based on the intensity of light scattered, NPG z-average was $1071 \pm$
345 30.65 nm. However, it is well known in the colloidal field that for large size
346 distribution the presence of bigger particles contributes significantly to the intensity of
347 light scattered ($I_{\theta} \approx r^6$, r the particle radius) compared to smaller particles. Such high
348 polydispersity tend to mask the presence and characterization of lower size
349 distribution in the colloidal dispersions of materials using CONTIN algorithm. (**Fig. 1**
350 **C-D**). Moreover, NPG dispersions present irregular shapes, high surface oxidation
351 and specific surface area increasing their adsorption properties (El Hadri *et al.*,
352 2020a). Numerous contaminant and pollutant families can thus interact with
353 environmental NPs, but also change their bioavailability and toxicity (Alimi *et al.*,
354 2018; Bhagat *et al.*, 2020). Therefore adding environmentally weathered plastics
355 brings relevance to NP ecotoxicological studies.

356

357 3.2 Adsorption of nanoplastics on microalgae

358 The SEM observations highlighted that PSL was adsorbed on *T. lutea* surface in all
359 of the tested concentrations (**Fig. 2**). At environmentally realistic NP concentrations,
360 10 and 100 $\mu\text{g L}^{-1}$, we observed fewer adsorbed PSL per microalgae (**Fig. 2 B-C**)
361 compared to the higher concentrated treatments (**Fig. 2 D-E**). Noteworthy, PSL
362 aggregates consistently appeared starting at 1000 $\mu\text{g L}^{-1}$ (**Fig. 2 D**). No differences
363 were observed in the number of adsorbed PSL between 1000 and 5000 $\mu\text{g L}^{-1}$,
364 suggesting a saturation of the particles onto microalgae surfaces. Although these
365 results are representative of a short exposure (48 h) in estuarine-like conditions.
366 Indeed *T. lutea* solutions were homogenized twice a day to mimic dynamic
367 interactions and to avoid sedimentation. In general, we did not notice damage on the
368 microalgae surface. This result is contradictory with Wang *et al.* (2020) that observed
369 fragmented shapes and cracked surfaces onto marine microalgae exposed for 96 h
370 to 200 and 2000 $\mu\text{g L}^{-1}$ of PS NPs (70 nm) associated with molecular and
371 physiological hazards. Such difference can be potentially explained by the presence
372 of surfactant and additives generally used for commercially available PSL (Pikuda *et*
373 *al.*, 2018). Unlike PSC and NPG, PSL were a valuable NPs model easy to target by
374 SEM given their consistent spherical shape. Also, Pessoni *et al.* (2019) previously
375 demonstrated the PSL carboxylated surface functions should prevent them from
376 forming homoaggregates. Nonetheless, PSL most likely formed heteroaggregates in
377 the presence of NOM noticeable as clear dots in the SEM observations (**Fig. 2**). NP
378 aggregations had been observed to change their bioavailability, and thus, to either
379 increase or decrease NP toxicity (Corsi *et al.*, 2014; Zhang *et al.*, 2018). Surprisingly,
380 despite PSL heteroaggregations and their global negatively charged surface, the
381 SEM images showed that PSL was able to adsorb onto microalgae regardless of
382 natural organic matter's presence and for all the tested concentrations.

383

384 3.3 Oysters physiology and arsenic bioaccumulation

385 3.3.1 Biometric parameters and arsenic uptake

386 *Crassostrea virginica*'s biometric data are presented in **Fig. S1**. The registered
387 mortality during the exposure was minor (7%) and statistical differences were only
388 detected for fresh tissue weight (**Fig. S1 A**). Nonetheless, there were no significant

389 differences between the control and other treatments, between both concentrations
390 of the same NP treatment, nor between the As treatment and the NP + As
391 treatments (**Fig. S1 A**).

392 *Crassostrea virginica* oysters from control treatment had an average As
393 concentration of $1.6 \mu\text{g g}^{-1}$ (dry weight) in gills and $3.5 \mu\text{g g}^{-1}$ in visceral mass (**Fig. 3**).
394 As-exposed oysters (As and NPs + As treatments) displayed significantly different
395 levels of total As compared to the control treatment, around 5 fold-change for both *C.*
396 *virginica* tissues. In perspective with oysters' capacity to bioaccumulate As in several
397 orders of magnitude higher (Zhang *et al.*, 2015), our 1 mg L^{-1} treatment does not
398 suggest an excessive As exposure. Yet, all As-exposed oysters accumulated
399 statistically similar levels of total As respectively to each tissue. Thereby, the NP
400 treatments at $100 \mu\text{g L}^{-1}$ did not significantly influence the total As accumulation in *C.*
401 *virginica* after one-week exposure. We observed similar results for *I. alatus* oysters
402 that were exposed to NPG + As (Lebordais *et al.*, *submitted*). In both oyster
403 experiments with comparable NP and As levels, the presence of NPs did not affect
404 the total As bioaccumulation in any oyster tissues. It seems that NP exposures at
405 environmentally realistic concentrations were not sufficient to change the total As
406 bioaccumulation by adsorption. These observations could also witness the low
407 adsorption affinity of arsenate (an oxy-anion) towards the negatively charged
408 carboxylated functions onto the three NPs surface.

409 Indeed, these results are consistent with an environmental range of total
410 accumulated As (from 4.1 to $39 \mu\text{g g}^{-1}$, dry weight) in *C. virginica* individuals from
411 Gulf of Mexico coastal areas (Wilson *et al.*, 1992). To better assess and compare
412 oysters bioaccumulation, their native background should be considered. The *C.*
413 *virginica* oysters were farmed at St-Simon bay that is connected to Chaleur bay,
414 where controlled effluents from industrial activities have been discarded. The Human
415 Health Risk Assessment survey (Ministère de la Santé et du Mieux-être du Nouveau
416 Brunswick, 2005) has revealed a total As concentration up to 2 mg kg^{-1} (wet weight)
417 in local mussels. This background value is consistent with our measured total As
418 levels in *C. virginica* with an average of 0.7 mg kg^{-1} (wet weight) in gills and 1.4 mg
419 kg^{-1} in visceral mass for control. Of note, wet weight tissues have a limited relevance
420 to assess metals accumulation, yet we calculated *C. virginica* wet weights in order to
421 make the previous comparisons. These bioaccumulation comparisons between in-

laboratory and wild bivalves can be made since metals kinetic accumulation are well-known. Indeed non-essential metals (such as As for oysters, Rodney *et al.*, 2007; Ali and Khan, 2019) usually bioaccumulate with high rates reaching their peak during the first weeks post exposure followed by a plateau (Wallner-Kersanach *et al.*, 2000). It is thus expected that our exposed *C. virginica* would quickly bioaccumulate As up to comparative levels as those measured in wild *C. virginica* lifetime exposed to As.

3.3.2 Species bioaccumulation factor comparison

In a previous experiment, Caribbean flat oysters *I. alatus* from Guadeloupe mangrove swamps were also exposed in-laboratory to 1 mg L⁻¹ of As for one week (Lebordais *et al.*, *submitted*). *I. alatus* and *C. virginica* underwent identical experimental conditions regarding microalgae feeding, salinity, temperature, and light conditions. The relative As bioaccumulation yielded 2-fold change in *I. alatus* tissues compared to controls, while *C. virginica* yielded approximately 5- to 10-fold change in gills and visceral mass, respectively (**Fig. 4**). There was a significant difference between the gills of *I. alatus* and *C. virginica*. In most aquatic organisms, gills are known to be a transport organ for metallic contaminants (Kraemer *et al.*, 2005; Won *et al.*, 2016, Cao *et al.*; 2018). As gills respond rapidly to metal contamination in water, they have a short-term role in regulation (Langston, 1984; Strady *et al.*, 2011). Ultimately, they are more sensitive through waterborne contamination until metals move into storage organs, like the visceral mass (Soegianto *et al.*, 2013; Arini *et al.*, 2014). As expected, high As concentrations were measured in *I. alatus* oysters exposed in a laboratory setting (Lebordais *et al.*, *submitted*). Nonetheless, relative As levels suggested higher As uptakes for exposed *C. virginica* oysters (**Fig. 4**).

We then calculated the BAF using wet weight for each oyster species. Whole-body BAFs estimated from gills and visceral mass were 13.4 (\pm 6.1) for *I. alatus* and 57.7 (\pm 41.9) for *C. virginica*. These BAF values confirmed a higher As uptake for *C. virginica* individuals compared to *I. alatus* suggesting *I. alatus*' physiology is potentially adapted to grow into As-rich environments (Cherkasov *et al.*, 2010; Luo *et al.*, 2014). Regulation, biotransformation, and/or detoxification mechanisms may enable *I. alatus* to tolerate higher As concentrations as revealed by the major As

454 bioaccumulation baseline in the control treatment. Consequently, *C. virginica* is
455 presumably more vulnerable to As exposure than *I. alatus*. Indeed, New Brunswick's
456 nurseries and bays carried most likely minimal As availability to *C. virginica* (as
457 revealed by the low As bioaccumulation baseline in controls). Moreover, BAF
458 standard deviations show noticeable contrast for each species. This difference might
459 be related to oysters filtration rates. *Isognomon alatus* oysters showed consistent
460 BAF values between individuals, while *C. virginica* showed a higher BAF variability
461 between individuals. This high variability could be explained by a putative change in
462 filtration rates between *C. virginica* oysters trying to protect themselves from metal
463 contamination (Tran *et al.*, 2003; Pan and Wang, 2012; Freiras *et al.*, 2018).

464 The gills and visceral mass of exposed *I. alatus* accumulated 10 and 20 times
465 more As than *C. virginica* tissues, respectively (Lebordais *et al.*, *submitted*). However,
466 As bioaccumulation was already higher in *I. alatus* controls than in exposed *C.*
467 *virginica* oysters. This major difference could be due to the oyster's respective
468 habitats. The As concentration measured in Guadeloupean seawater was below the
469 limit of detection, and As concentration in upper St. Lawrence estuary (Quebec,
470 Canada) can be up to $1.5 \mu\text{g L}^{-1}$ for the highest salinity (Tremblay and Gobeil, 1990).
471 There is little variability of As levels in seawater, and according to Neff (1997) the
472 concentration in clean coastal and oceans ranges within $1\text{-}3 \mu\text{g L}^{-1}$. Thus, it can be
473 considered both species were in waters with naturally similar As concentrations.
474 Nonetheless, sediment plays a key role in As waterborne route for filter-feeding
475 bivalves (Langston, 1984; Zhang *et al.*, 2013; Maher *et al.*, 2018). Consequently, As
476 concentrations in Toucari bay (close to the Guadeloupe island) ranged from 27.8 to
477 40.9 mg kg^{-1} in submarine sediment (Johnson and Cronan, 2001); whereas, the
478 average As concentration in Chaleur bay sediment (New Brunswick, Canada) was
479 11 mg kg^{-1} (Parsons and Cranston, 2005). Also, As is accumulated by primary
480 producers like phytoplankton being the main As dietary route for marine consumer
481 organisms (Neff, 1997; Azizur Rahman *et al.*, 2012; Maher *et al.*, 2018) and warmer
482 water temperatures can facilitate total As uptake for bivalves (Ünlü and Fowler, 1979;
483 Gutierrez-Galindo *et al.*, 1994). As such, tropical *I. alatus* oysters grew in mangrove
484 swamps that are productive ecosystems with major sediment and phytoplankton
485 inputs (Saed *et al.*, 2004; Bouillon, 2011; Yap *et al.*, 2011). In contrast, subarctic *C.*
486 *virginica* oysters were farmed inside floating gears in less rich waters from a

487 Canada's Eastern bay. The age of the oysters should also be considered as it might
488 have influenced the As accumulation results. *C. virginica* oysters were three to four
489 years old, while *I. alatus* oysters were about six years old based on this oyster
490 growth rate in Jamaica mangroves (nearby Guadeloupe island, Siung, 1980). Overall,
491 As concentrations measured in both oyster controls witnessed their natural As
492 burden that is representative of their respective habitat.

493

494 3.4 Oysters gene responses after exposure

495 3.4.1 Relative gene expression in gills and visceral mass

496 We selected a set of genes of interest to target ecotoxicological endpoints gathered
497 in seven biological functions, such as endocytosis (*cltc*), oxidative stress (*cat*, *gapdh*,
498 *sod3*), mitochondrial metabolism (*12S*), cell cycle regulation (*gadd45*, *p53*),
499 apoptosis (*bax*, *bcl-2*), detoxification (*cyp1A*, *mdr*, *mt*), and energy storage (*vit*)
500 (**Table S1**). Of note, we did not detect *vit* in gills after PCR and agarose gel analysis;
501 thereby we only show its results for visceral mass. The **figures 5** and **6** show the
502 relative mRNA levels results for gills and visceral mass, respectively.

503 The expression of *12S* significantly increased with PSL treatment at $10 \mu\text{g L}^{-1}$
504 compared to the control treatment, but it did not increase at $100 \mu\text{g L}^{-1}$ (**Fig. 5 A**).
505 The exposure to PSL + As at $10 \mu\text{g L}^{-1}$ did not affect significantly the expression of
506 *12S*, suggesting that the induction for PSL treatment at $10 \mu\text{g L}^{-1}$ was prevented by
507 the presence of As. Similar antagonist effects of As in mixture with PSL were
508 observed in *I. alatus* exposure (Lebordais *et al.*, *submitted*). Ribosomic RNA (rRNA)
509 *12S* is transcribed by the mitochondrial genome. Therefore, *12S* rRNA level is used
510 as a proxy to represent the number of mitochondrial copies in a given tissue (Al
511 kaddissi *et al.*, 2012; Arini *et al.*, 2015). Considering that gills are involved in
512 respiration, their mitochondrial activity can be more sensitive to contaminants like
513 metals (Akberali and Earnshaw, 1982). Significant differences in *12S* expression for
514 As treatment were expected in gills through waterborne route. Instead, the data
515 suggest that As was more available through the dietary route (**Fig. 6 C**). The
516 expression of *bax* significantly increased in the PSC + As treatment at $100 \mu\text{g L}^{-1}$
517 compared to control, As alone and PSC at $100 \mu\text{g L}^{-1}$ treatments (**Fig. 5 B**). Also, the
518 *bax* expression for PSL + As treatment significantly increased at $10 \mu\text{g L}^{-1}$, but it did

519 not change significantly at $100 \mu\text{g L}^{-1}$. In addition, PSL + As at $10 \mu\text{g L}^{-1}$ significantly
520 increased compared to control, As alone and PSL at $10 \mu\text{g L}^{-1}$ treatments. Overall,
521 the presence of As in PSC at $10 \mu\text{g L}^{-1}$ and PSL at $100 \mu\text{g L}^{-1}$ induced a synergetic
522 *bax* expression. *Bcl-2* gene family are central regulators of programmed cell death.
523 Indeed, the intrinsic pathway for apoptosis relies on upregulation of pro-apoptotic
524 genes like *bax* (Schuler *et al.*, 2000; Chipuk *et al.*, 2010). Seen *bax* regulation
525 integrates internal and external stress signals, its expression gives a relevant proxy
526 to assess contaminant toxicities at the cell level (Yanan 2012). For example, Cao *et*
527 *al.* (2018) observed a significant increase of *bax* relative mRNA level in *Crassostrea*
528 *gigas* gills and observed apoptosis at $10 \mu\text{g L}^{-1}$ of Cd. As such, the combination of As
529 treatment with PSC and PSL most likely induced apoptotic effects in *C. virginica* after
530 one week of exposure. The expression of *gapdh* significantly increased in PSC + As
531 treatment at $10 \mu\text{g L}^{-1}$ compared to control and PSC at $10 \mu\text{g L}^{-1}$ treatments (**Fig. 5**
532 **C**). The *gapdh* response can be associated to oxidative stress responses but no
533 further interpretation can be made since its protein is involved in the nucleus,
534 mitochondrial and cytosolic signal pathways (Sirover 2011, Tristan *et al.*, 2011).

535 The expression of *12S* significantly increased for NPG + As treatment at 10 and
536 $100 \mu\text{g L}^{-1}$ compared to control, As alone and NPG at 10 and $100 \mu\text{g L}^{-1}$ treatments
537 (**Fig. 6 C**). Similarly, the expression of *12S* significantly increased in PSC + As
538 treatment at 10 and $100 \mu\text{g L}^{-1}$ compared to control, to As alone, and PSC (at both
539 concentrations). Thus the presence of As in NPG + As and PSC + As induced a
540 synergetic *12S* expression. Also, PSL + As treatment significantly increased *12S*
541 expression at $100 \mu\text{g L}^{-1}$ compared to control, As alone and PSL + As at $10 \mu\text{g L}^{-1}$
542 treatments. Based on these gene expression profiles, As treatment alone did not
543 affect mitochondrial metabolism. Yet, in the presence of NPs the results could
544 suggest that As was more effective in impairing mitochondrial metabolism like
545 respiration. Indeed, metal toxicity can decrease oxygen consumption in mitochondria
546 (Sokolova *et al.*, 2005; Cherkasov *et al.*, 2010). We expected mitochondrial gene
547 response to be less sensitive to As exposure in visceral mass (Akberali and
548 Earnshaw, 1982). Nonetheless, the mRNA level of *12S* for NPs + As treatments was
549 significantly higher only in visceral mass. These results support stronger effects of
550 As combined with NPs and suggest that diet was the main driver for oysters
551 exposure to NPs. Therefore, the combined NPs + As treatments triggered a

552 synergetic effect that most likely increased the number of mitochondria. The
553 expression of *gadd45* for NPG + As treatment at $100 \mu\text{g L}^{-1}$ was not significantly
554 different from its control nor the As alone treatment. Yet, it significantly decreased
555 compared to NPG at $100 \mu\text{g L}^{-1}$. This indicates that the expression of *gadd45* for
556 NPG treatment at $100 \mu\text{g L}^{-1}$ was prevented in combination with As. The expression
557 of *p53* for NPG + As treatment at $100 \mu\text{g L}^{-1}$ did not significantly change compared to
558 the control and the As alone treatments, but its expression significantly increased
559 compared to NPG + As at $10 \mu\text{g L}^{-1}$ (**Fig. 6 D**). The expression of *p53* for PSL
560 treatment at $100 \mu\text{g L}^{-1}$ did not significantly change compared to the control, but its
561 expression significantly increased compared to the PSL + As at $100 \mu\text{g L}^{-1}$ and to
562 PSL at $10 \mu\text{g L}^{-1}$ treatments. Thereby, PSL treatment at $100 \mu\text{g L}^{-1}$ in combination
563 with As prevented *p53 gene* expression (**Fig. 6 E**). Overall, the exposure to As in
564 NPG + As treatment at $100 \mu\text{g L}^{-1}$ significantly changed *gadd45* and *p53* expressions
565 relatively to NPG treatment for the same concentration. The opposite *gadd45* and
566 *p53* responses to NPG + As treatment might be caused by several negative
567 feedback loops (Harris and Levine, 2005). Indeed *gadd45* regulation is (among
568 others) upon P53 transcription factor, hence its downregulation can be linked to P53
569 increase (Salvador *et al.*, 2013). The expression of *bcl-2* significantly increased in
570 the PSL treatment at $100 \mu\text{g L}^{-1}$ compared to control and to PSL at $10 \mu\text{g L}^{-1}$
571 treatments (**Fig. 6 G**). The *bcl-2* expression for PSL at $100 \mu\text{g L}^{-1}$ also significantly
572 increased compared to As treatment, but not to PSL + As at $100 \mu\text{g L}^{-1}$. Thereby, the
573 expression of *bcl-2* for PSL treatment at $100 \mu\text{g L}^{-1}$ was prevented in combination
574 with As. Despite being an anti-apoptotic gene (Aouacheria, 2005), *bcl-2* upregulation
575 has been associated with a significant decrease of red granulocyte percentage in
576 clams blood exposed to MPs (Tang *et al.*, 2019).

577

578 3.4.2 Combined effects of arsenic with nanoplastics

579 We observed specific gene responses for combined NPs + As treatments in both
580 tissues. A single antagonist effect was measured (**Fig. 5 A**), while consistent
581 synergetic effects were measured (**Fig. 5 B, 6 C**). Synergetic effects, according to
582 Bhagat *et al.* (2020) definition, were identified by genes upregulation for NPs + As
583 treatments significantly different from control, As and NP treatments alone. As such,

584 NPG + As and PSC + As synergetic effects on 12S expression revealed an increase
585 of rRNA level produced by mitochondria that could indicate the induction of
586 mitochondria number in visceral mass (**Fig. 6 C**). Yet, presumed toxicity on
587 mitochondrial metabolism (e.g., oxidative stress) cannot be associated with the
588 results since antioxidant gene responses of *cat*, *sod*, and *gapdh* were not consistent.
589 Interestingly, Freitas *et al.* (2018) measured a significant inhibition of electron
590 transport system activity, with significant inductions of superoxide dismutase and
591 catalase activities in clams exposed to combined multi-walled carbon nanotubes.
592 Synergetic or antagonist effects have been observed for several combined
593 contaminants (Kim *et al.*, 2017; Bhagat *et al.*, 2020) and are most likely related to
594 contaminant changes in bioavailability and/or speciation (Spurgeon *et al.*, 2020). In
595 our experiment *T. lutea* potentially biotransformed As, but only in NPs + As
596 treatments. These As speciation changes might have been reduced by the presence
597 of NPs. Indeed, *T. lutea* was very likely shortly exposed to As before being
598 consumed by *C. virginica*. Thus, As have been potentially metabolized by *T. lutea* as
599 documented for microalgae (Cullen and Reimer, 1989; Azizur Rahman *et al.*, 2012).
600 For instance, phytoplankton and bacteria are major producers of As methylated
601 forms (e.g., dimethylarsinous acid, monomethylarsonous acid; Wood, 1974;
602 Francesconi and Edmonds, 1996; Hellweger and Lall, 2004). However, in the
603 presence of NPs, adsorbed inorganic As is potentially not available for *T. lutea*
604 methylation. According to Farrell *et al.* (2011), adsorption can affect As speciation.
605 These observations matter for toxicity assessment using gene expression responses
606 since As methylated and inorganic forms have distinct molecular targets,
607 bioaccumulation rates and excretion pathways (Dixon, 1997; Petric *et al.*, 2001;
608 Hughes, 2002; Ng, 2005). Thereby As methylated forms present in As treatment may
609 triggered different mRNA level responses from inorganic As present in NPs + As
610 treatment. In this study, putative quantities of As methylated forms in As treatment
611 might be less effective since we observed no significant changes for any investigated
612 gene. Inorganic As forms are known to be more toxic than As methylated forms
613 (Sephar *et al.*, 1980; Neff, 1997; Petric *et al.*, 2001, Ng 2005). Therefore,
614 comparatively to As treatment, higher amounts of inorganic As available for *C.*
615 *virginica* could explain synergetic effects measured for NPs + As treatments.

616 We revealed herein synergetic effects in *C. virginica* exposure in contrast with
617 the protective effects measured in *I. alatus* exposure, potentially driven by both
618 species different physiology and sensitivity to As. These gene expression differences
619 between both oyster species for As + NP treatments need to be considered in light of
620 their respective As bioaccumulation. Therefore, the gene expression results from
621 combined As + NP treatments highlighted the relevance of comparative species
622 studies. As such the natural background and biology differences (observed for As
623 bioaccumulation) led to contrasted effects between Canadian farmed oysters *C.*
624 *virginica* and Caribbean wild oysters *I. alatus*.

625

626 **4. Conclusion**

627 There is a current lack of relevance in the literature of NPs ecotoxicity that we
628 addressed by comparing the effects of three NP models with a gradient of
629 environmental relevancy. In regards to *C. virginica* gene responses, the most
630 effective NPs treatment combined with As was PSC, while PSL was the most
631 effective NPs treatment conducted alone. Yet, with NPG, all three NPs in presence
632 of As showed synergetic effects in both tissues of *C. virginica*. Our trophic exposure
633 shed light on significant synergetic effects measured in mitochondrial metabolism
634 and apoptosis related genes at low NP concentrations (10 and 100 $\mu\text{g L}^{-1}$). These
635 are novel results regarding As combination with environmentally representative NPs
636 (NPG). Overall, there were more significant gene responses to NP treatments in
637 visceral mass than in gills. These results suggest that NPs were more available or
638 effective by dietary route as we expected. Furthermore, our BAF results compared
639 between two oyster species revealed higher As uptake for Canadian farmed oysters
640 *C. virginica* than Caribbean wild oysters *I. alatus*. This comparative approach
641 provided valuable data that revealed the effects variability of combined As + NP
642 treatments between two oyster species. Therefore, future studies should target the
643 As speciation to better assess the As forms being adsorbed to NPs and the putative
644 role of microalgae. A longer exposure time at lower As concentration will also be
645 relevant to confirm the gene expressions and bioaccumulation data measured in this
646 study.

647

648 5. Acknowledgments

649 We acknowledge Bruno Grassl, Hind El Hadri and Stephanie Reynaud (IPREM) for
 650 providing the polystyrene nanoparticles; and Réjean Tremblay (UQAR) for providing
 651 microalgae solutions. The authors would like to thank Joy Gaubert (INRS) for help
 652 with the molecular assays and Jean-François Dutil (INRS) for help with arsenic
 653 analyses. The authors also appreciated the helpful comments from Patrice Couture
 654 and the unknown referees.

655

656 6. Funding source

657 This work was financially supported by the Agence Nationale de la Recherche (ANR)
 658 in the PEPSEA program (ANR-17-CE34-0008-05) to MB and JG, the Centre
 659 National de la Recherche (CNRS) in the EC2CO INIAME program to MB and JG,
 660 and the Canada Research Chair program (950-23223) to VSL.

661

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- 1047

1048 **Figures**

1049 **Figure 1.** Hydrodynamic diameters of NP particles measured by DLS using a
 1050 Zetasizer nano zs. Size distribution for PSC dispersion with z-average of $692.4 \pm$
 1051 68.55 nm (A) and the corresponding number of nanoparticles (B). Size distribution
 1052 for NPG dispersion with z-average of 1071 ± 30.65 nm (C) and the corresponding
 1053 number of nanoparticles (D). Each NPG batch is presented with a different shade of
 1054 grey.

1055 **Figure 2.** Scanning Electron Microscopy (X8000) observation of carboxylated
 1056 polystyrene nanospheres of latex (PSL) adsorbed on *T. lutea* exposed for 48 h to 0
 1057 $\mu\text{g L}^{-1}$ (A), $10 \mu\text{g L}^{-1}$ (B), $100 \mu\text{g L}^{-1}$ (C), $1000 \mu\text{g L}^{-1}$ (D) and $5000 \mu\text{g L}^{-1}$ (E). Arrows
 1058 indicate the presence of adsorbed PSL to microalgae. Of note, only $10 \mu\text{g L}^{-1}$ and
 1059 $100 \mu\text{g L}^{-1}$ were used for the oyster diet exposure.

1060 **Figure 3.** Arsenic bioaccumulation ($\mu\text{g/g}$, dry weight, mean + sd) in *C. virginica* gills
 1061 and visceral mass ($n = 4$), after one week of exposure to 1 mg L^{-1} As and/ or $100 \mu\text{g}$
 1062 L^{-1} NPs.

1063 **Figure 4.** Relative arsenic bioaccumulation ($\mu\text{g/g}$, dry weight, mean + sd) compared
 1064 between *I. alatus* and *C. virginica* in gills and visceral mass ($n = 4$), after one week of
 1065 exposure to 1 mg L^{-1} As. Asterisks show a significant difference ($p < 0.01$) assessed
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1067 **Figure 5.** Relative gene expressions in *C. virginica* gills after one-week exposure to
 1068 1 mg L^{-1} As combined or not with 10 and $100 \mu\text{g L}^{-1}$ NPs. mRNA levels are
 1069 presented for *12S* (A), *bax* (B), *gapdh* (C) and *mt* (D). All the values are presented
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 1071 denote significant differences ($p < 0.05$) among treatments assessed by two-way
 1072 ANOVA followed by Tukey post-hoc test. Bold, italic and capital letters are used for
 1073 PSL, PSC and NPG treatments respectively.

1074 **Figure 6.** Relative gene expressions in *C. virginica* visceral mass after one-week
 1075 exposure to 1 mg L^{-1} As combined or not with 10 and $100 \mu\text{g L}^{-1}$ NPs. mRNA levels
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1082 **Supplementary file titles**

1083 **Supplementary file Figure S1** – *C. virginica* biometric parameters.

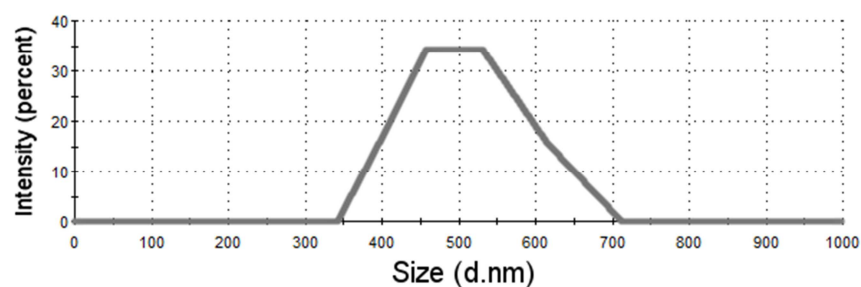
1084 **Supplementary file Table S1** – *C. virginica* targeted gene functions and optimized
1085 primer sets temperatures for qPCR.

1086 **Supplementary file Figure S2** – *C. virginica* relative gene expressions in gills.

1087 **Supplementary file Figure S3** – *C. virginica* relative gene expressions in visceral
1088 mass.

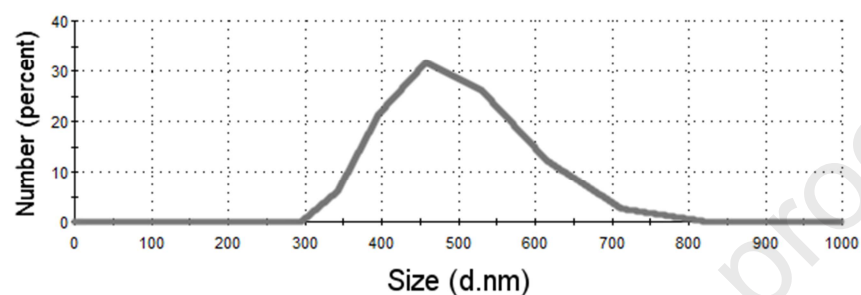
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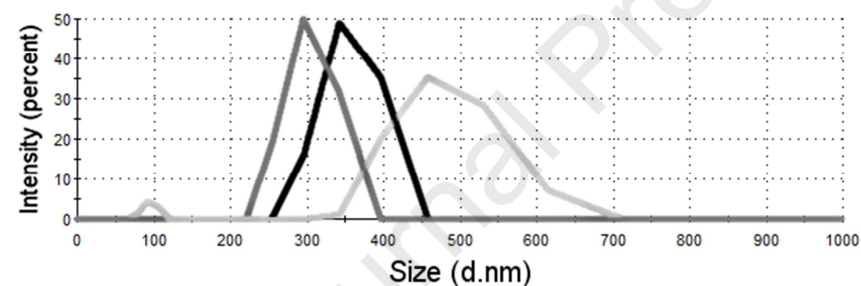
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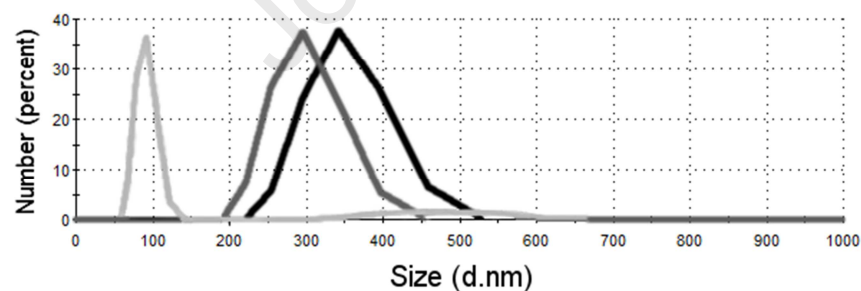
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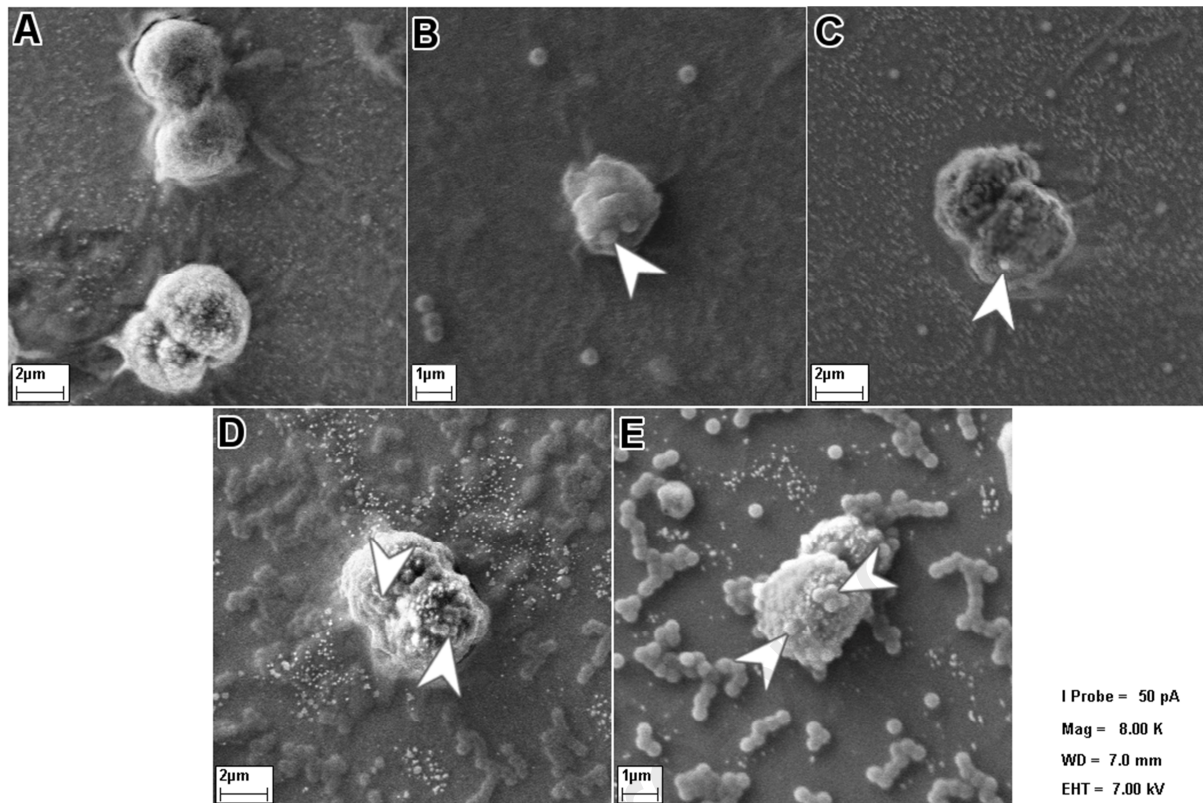
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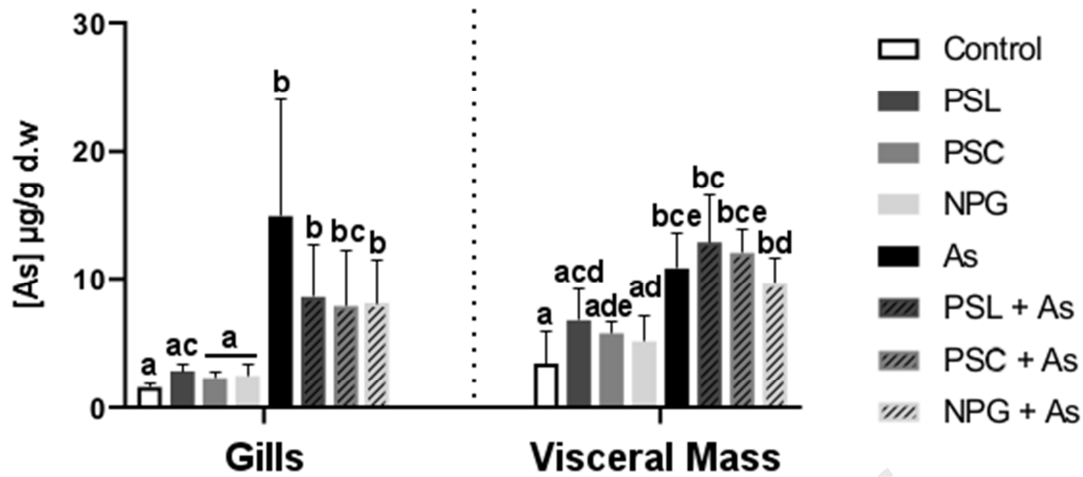
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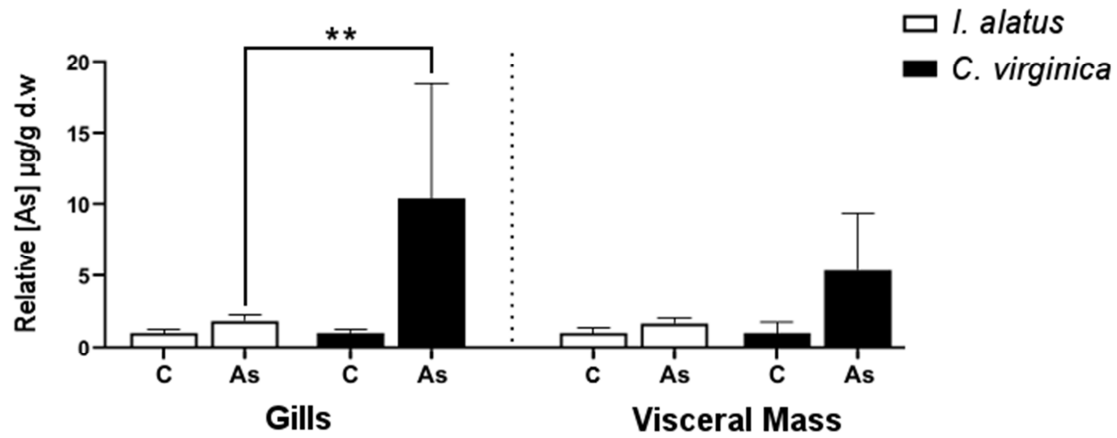
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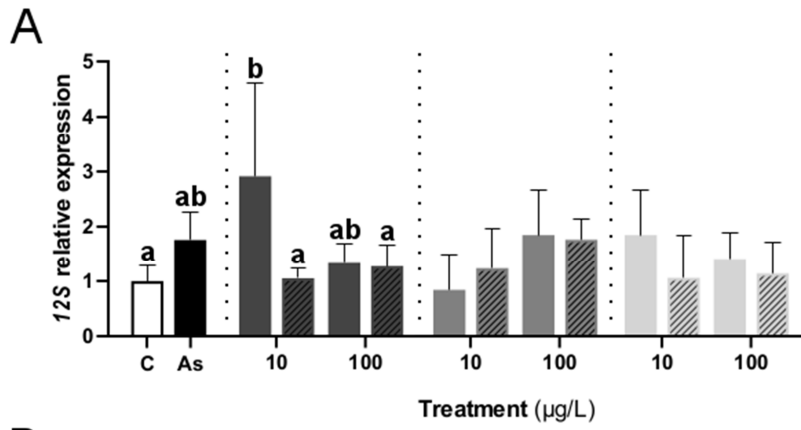
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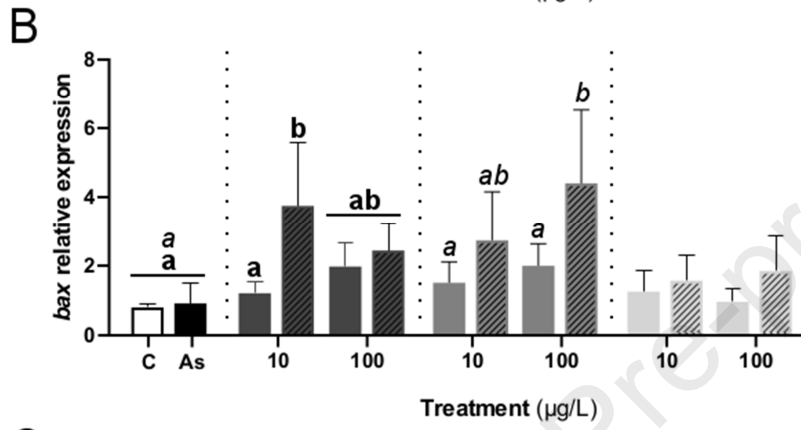
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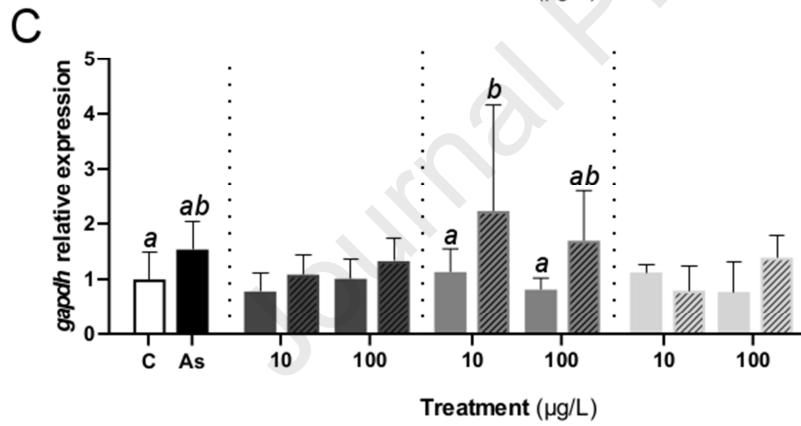
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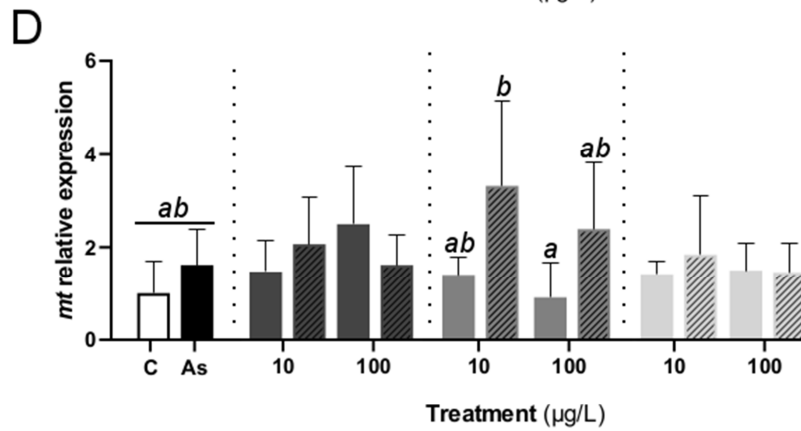
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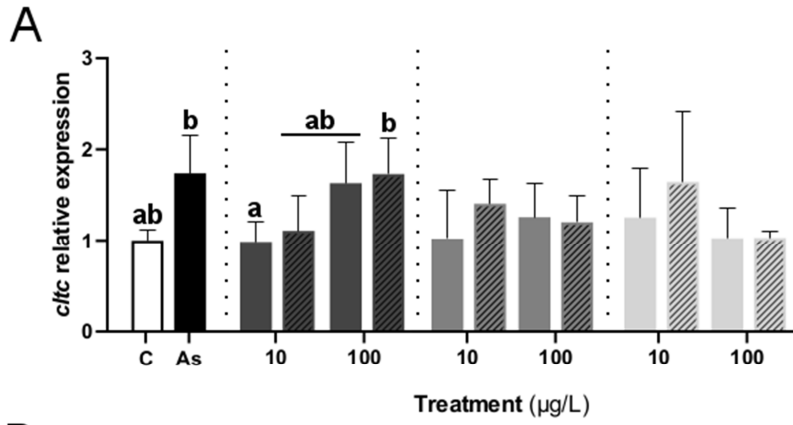
PSL PSL+As PSC PSC+As NPG NPG+As

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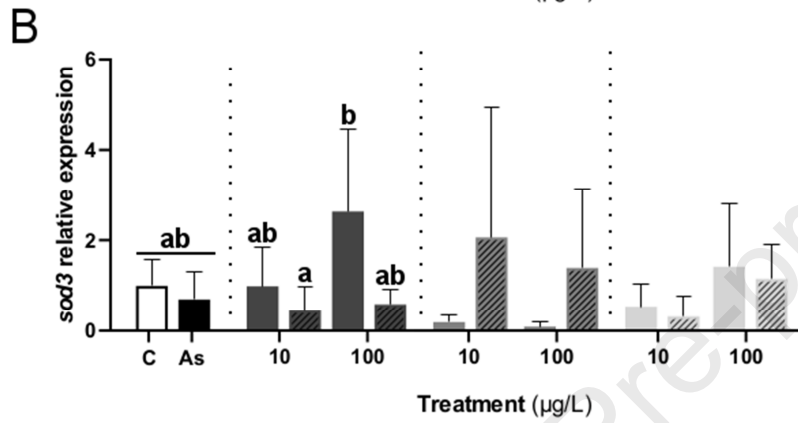
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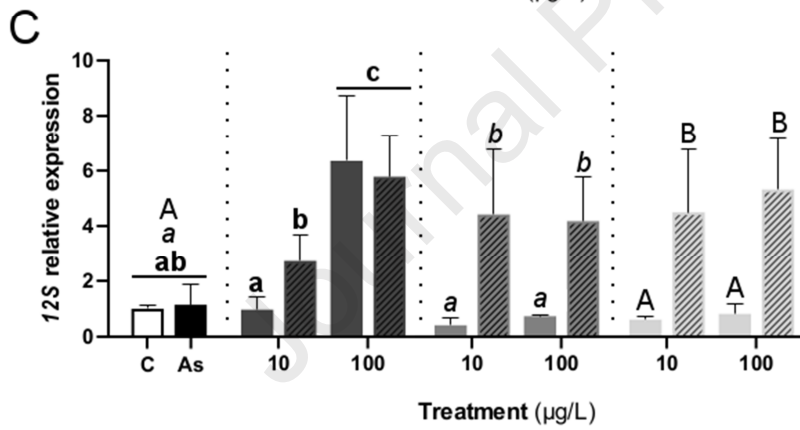
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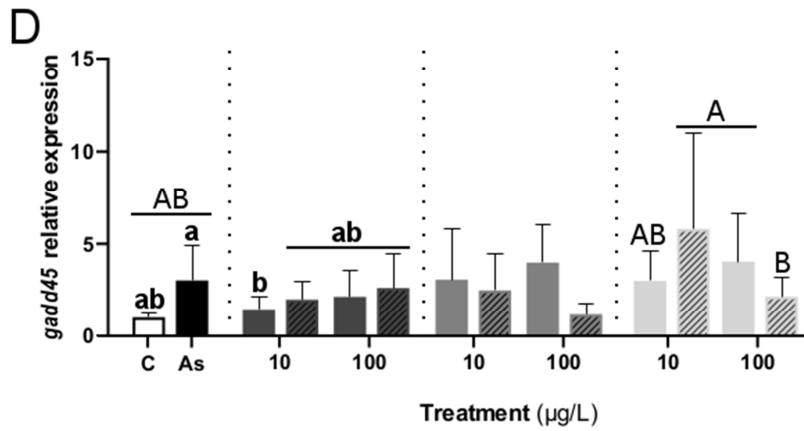
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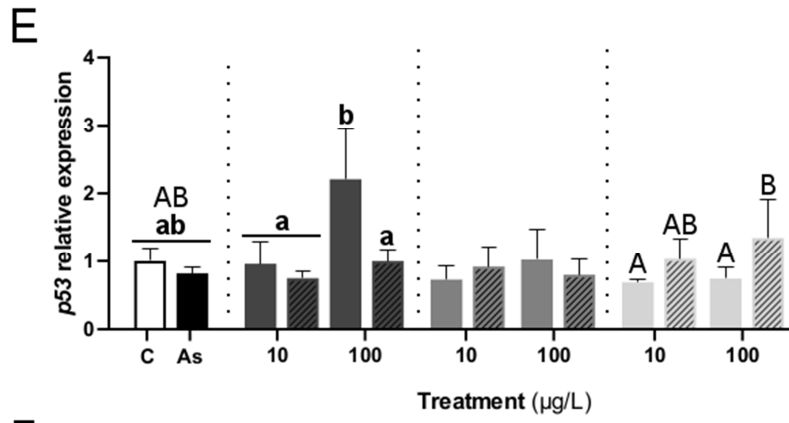
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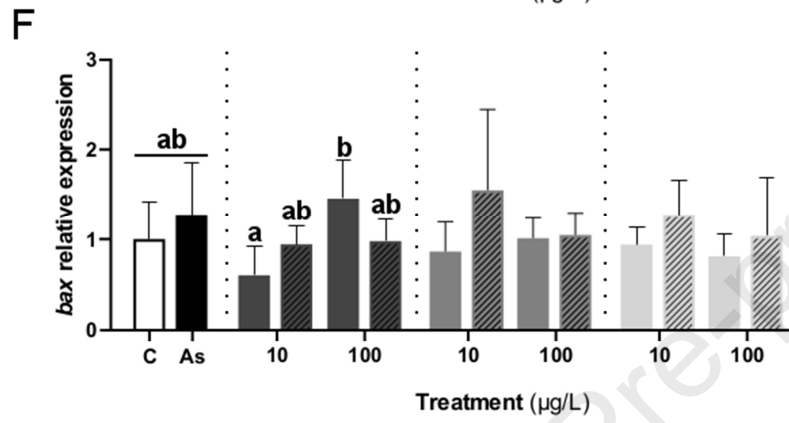
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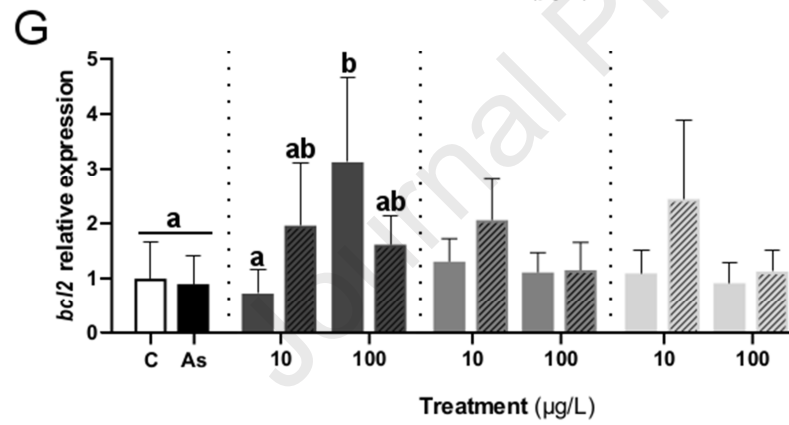
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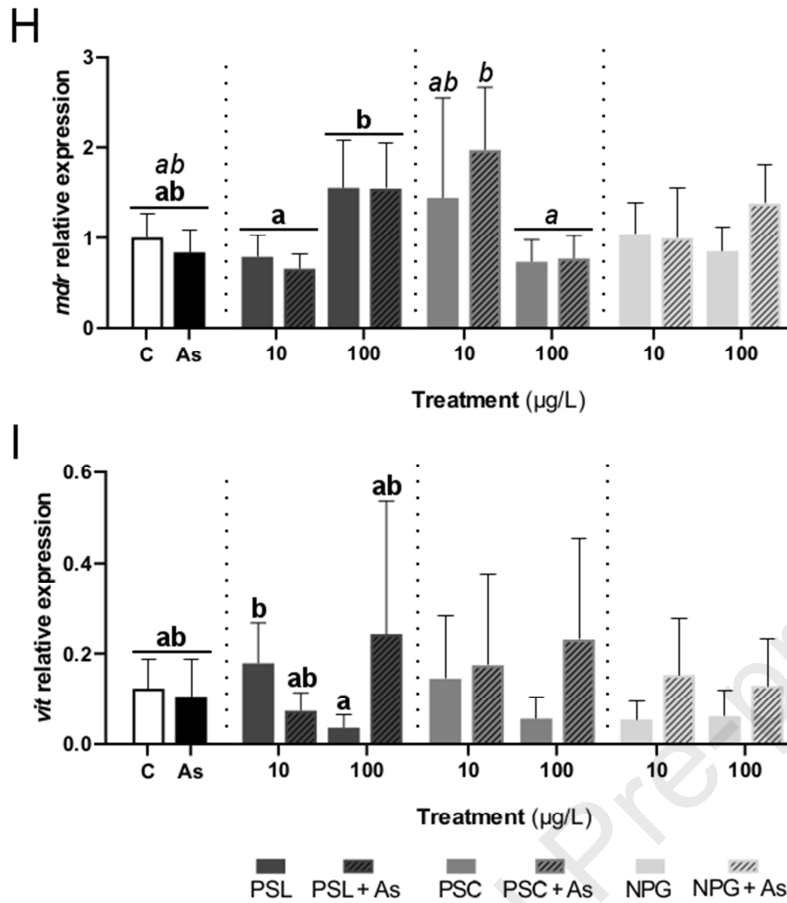
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 1149 respectively.

1 **Highlights**

2

3 - Nanoplastics are adsorbed on microalgae surface starting at $10 \mu\text{g L}^{-1}$
4 exposure.

5 - Nanoplastics triggered apoptotic and mitochondrial metabolism gene
6 responses.

7 - The combination of nanoplastics with arsenic induced synergetic effects.

8 - Arsenic bioaccumulation did not increase in the presence of nanoplastics.

Journal Pre-proof

Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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