

1 **Modulation of neuro-dopamine homeostasis in juvenile female Atlantic cod (*Gadus morhua*)**  
2 **exposed to polycyclic aromatic hydrocarbons and perfluoroalkyl substances**

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6 Essa A. Khan<sup>1</sup>, Luisa B. Bertotto<sup>2</sup>, Karina Dale<sup>3</sup>, Roger Lille-Langøy<sup>3</sup>, Fekadu Yadetie<sup>3</sup>, Odd André  
7 Karlsen<sup>3</sup>, Anders Goksøyr<sup>3</sup>, Daniel Schlenk<sup>2</sup>, Augustine Arukwe<sup>1\*</sup>

8  
9 <sup>1</sup>Department of Biology, Norwegian University of Science and Technology (NTNU),  
10 Høgskoleringen 5, N-7491 Trondheim, Norway

11 <sup>2</sup> Department of Environmental Sciences, University of California-Riverside, California, USA

12 <sup>3</sup> Department of Biological Sciences, University of Bergen, N-5020 Bergen, Norway

13  
14  
15  
16 **\*Corresponding author:**

17 E-mail: [augustine.arukwe@ntnu.no](mailto:augustine.arukwe@ntnu.no)

27 **Abstract**

28 The dopaminergic effect of PAH and PFAS mixtures, prepared based on environmental levels has  
29 been studied in juvenile female Atlantic cod (*Gadus morhua*). Benzo[a]pyrene, dibenzothiophene,  
30 fluorene, naphthalene, phenanthrene and pyrene were used to prepare a PAH mixture, while PFNA,  
31 PFOA, PFOS and PFTrA were used to prepare PFAS mixture. Cod were injected intraperitoneally  
32 twice, with either a low (1x) or high dose (20x) of each compound mixture or various combinations.  
33 After two week of exposure, levels of plasma 17 $\beta$ -estradiol (E2) were significantly high in high  
34 PAH/high PFAS treated groups. Dopamine: metabolite ratios (DOPAC/dopamine and  
35 HVA+DOPAC/dopamine) in brain homogenate changes with the levels of E2 in plasma except for  
36 high PAH/low PFAS and low PAH/high PFAS treated groups. In general, *th* mRNA levels inversely  
37 correlated with dopamine: metabolite ratios and *gnrh2* mRNA levels. Respective decreases and  
38 increases of *dr1* and *dr2a* after exposure to the high PAH dose were observed. Whereas, high PFAS  
39 exposure decreased both *drs*, leading to high plasma E2 concentrations. Other investigated endpoints  
40 suggest that these compounds at different doses and combinations have different toxicity threshold  
41 and mode of actions. These effects indicate potential alternations in feedback signalling processes  
42 within dopaminergic pathway by these contaminants.

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44 **Keywords:** PAH; PFAS; Dopaminergic pathway; Estrogenic pathway; Feedback control; Atlantic  
45 cod.

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## 54 **1. Introduction**

55 Polycyclic aromatic hydrocarbons (PAHs) and perfluoroalkyl substances (PFASs) are among the  
56 most common xenobiotics found in the environment.<sup>1,2</sup> PAHs are generally categorized into  
57 petrogenic and pyrogenic hydrocarbon groups, depending on whether they are produced by the  
58 incomplete combustion of petroleum or by irreversible temperature-mediated change of chemical  
59 composition.<sup>3</sup> PFASs have been used in a wide variety of products, including fire-fighting foams, ink,  
60 paper coating and textile, and as water repellents.<sup>4,5</sup> Both PAHs and PFASs are found in the aquatic  
61 environment and may threaten marine organisms.<sup>6,7</sup> Concentrations of PAHs as high as 10 µg/g dry  
62 weight of sediment have been previously reported in the Seine estuary, Normandy, France.<sup>8</sup> PFAS  
63 concentrations have been observed at ng/L and ng/g levels in surface waters and sediments  
64 respectively.<sup>9-14</sup> Co-occurrence of both PAHs and PFASs at toxic levels has also been reported in  
65 tidal flats and costal ecosystems of the Ariake Sea, Japan.<sup>7,15</sup>

66 The concentrations of these compounds in the aquatic environment might not significantly  
67 affect survival, but may severely alter reproductive capacity and endocrinology in fish. PAHs have  
68 been shown to disrupt the endocrine system through binding and activating the aryl hydrocarbon  
69 receptor (AhR) and produce anti-estrogenic responses.<sup>16-18</sup> PAHs may also weakly bind to the  
70 estrogen receptor (ER) and have estrogenic properties.<sup>17</sup> PFASs have also been reported to possess  
71 estrogenic properties. For example, *in vivo* and *in vitro* exposure to PFOS produced an up-regulation  
72 of vitellogenin (*vtg*) mRNA expression in fish liver.<sup>19,20</sup> In contrast, other studies did not find  
73 significant changes in Vtg levels<sup>21</sup> or reported downregulation of *vtg* mRNA expression.<sup>22</sup> Further  
74 investigation is required to better understand the mechanism of action of PAHs, PFASs, and the  
75 combination of both compound classes on the reproductive system of teleosts.

76 Biosynthesis of estrogen is regulated through the hypothalamus-pituitary-gonadal (HPG)  
77 axis.<sup>23</sup> The hypothalamus produces gonadotropin-releasing hormone (GnRH) that controls the release  
78 of gonadotropins (GtHs): follicle-stimulating hormone (FSH) and luteinizing hormone (LH) from the  
79 pituitary gland. Gonadotropins induce oocyte development, maturation, and production of 17β-  
80 estradiol (E2) in gonads. Among different feedback mechanisms, the dopaminergic system plays an

81 essential role in controlling GnRH and GtH releases. High levels of E2 activate dopaminergic neurons  
82 generally to reduce the production of these hormones.<sup>23</sup> However, increases in the production of GtHs  
83 may also occur.<sup>24,25</sup> Dopamine can also regulate brain aromatase (Cyp19b), an enzyme that catalyzes  
84 the production of estrogen in the brain. Unlike GnRH, brain aromatase responds differently to  
85 dopaminergic agonists.<sup>26</sup> Fontaine et al.<sup>27</sup> showed that co-exposure of female fish to the dopamine  
86 receptor (Dr2a) antagonist domperidone and a GnRH agonist, resulted in an increased expression of  
87 GtH mRNA, suggesting that removal of dopamine inhibitory effects allow the hypothalamus to  
88 produce GnRH and consequently modulate plasma E2 levels.

89 The production of dopamine is initiated by tyrosine hydroxylase (TH), the rate-limiting  
90 enzyme that converts tyrosine to 3,4-dihydroxy-L-phenylalanine (L-DOPA) which is subsequently  
91 metabolized by DOPA decarboxylase (DDC) to dopamine.<sup>28</sup> Once released in the synaptic cleft,  
92 dopamine binds to two dopamine receptor (DR) families DR1 and DR2, which activate specific G  
93 proteins. For example, DR1 is coupled to  $G\alpha_s$  and activates adenylyl cyclase (AC) which increases  
94 the concentrations of cyclic adenosine monophosphate (cAMP) and calcium ions. Conversely, DR2  
95 is coupled to  $G\alpha_{i/o}$  and inhibits AC. In excess, dopamine is reabsorbed through the dopamine active  
96 transporter (Dat) in presynaptic neurons and further catabolized into 3,4-dihydroxyphenyl acetic acid  
97 (DOPAC) and homovanillic acid (HVA) by monoamine oxidase (MAO) and catechol-O-  
98 methyltransferase, respectively.<sup>28,29</sup> Exposure of mice to endocrine disruptive chemicals (EDCs),  
99 such as bisphenol A (BPA) and 2,4-dichlorophenoxyacetic acid, altered the expression of DRs and  
100 DAT, modulating dopamine synthesis, release and turnover in mice and rats.<sup>30,31</sup> Similar results were  
101 observed in zebrafish and rainbow trout where bifenthrin altered E2 concentrations and dopaminergic  
102 systems.<sup>32</sup>

103 Despite reports showing estrogenic responses of PAHs and PFASs, either individually or in  
104 combination, there is limited data concerning the effects of these compounds on dopaminergic  
105 signaling pathways. Therefore, the aim of this study was to investigate changes in dopaminergic  
106 signaling and endocrine function after *in vivo* exposure of Atlantic cod (*Gadus morhua*) to a low dose  
107 chosen based on environmental levels (low) and a higher (20 x low) dose (high) of PAHs and PFASs

108 alone or in combination. In the North Atlantic, Atlantic cod is major fisheries species, and is important  
109 in costal as well as oceanic ecosystems. It has been used as an indicator as well as model organism in  
110 environmental monitoring and toxicological studies respectively.<sup>33-35</sup> Therefore, cod is a valuable tool  
111 for ecotoxicological studies and risk assessment.

112

## 113 **2. Materials and methods**

### 114 2.1. Chemicals and reagents

115 Direct-zol<sup>TM</sup> RNA isolation and MiniPrep kit from Zymo Research Corporation (Irvine, CA, USA),  
116 iTaq SYBR Green Supermix with ROX and iScript cDNA synthesis Kit from Bio-Rad Laboratories  
117 (Hercules, CA, USA). 17 $\beta$ -estradiol (E2) enzyme immunoassay (EIA) kits (Cat. No. 582251 and  
118 582701) purchased from Cayman chemical company (Ann Arbor, MI, USA).

119

### 120 2.2. Animals

121 Juvenile Atlantic cod (*G. morhua*), approximately 5 months old were obtained from  
122 Havbruksstasjonen in Tromsø AS (Tromsø, Norway) and reared at Industrilaboratoriet in Bergen  
123 (ILAB, Bergen, Norway) in 500 L tanks supplied with seawater at 8 to 10 °C, 34 ppt salinity. The  
124 cod were held at a 12:12 h light/dark cycle and fed with a commercial marine diet (Amber Neptune,  
125 Skretting, Stavanger, Norway). At the start of the exposure the cod were approximately 18 months  
126 with average bodyweight of 172  $\pm$  34 g. The experimental setup was approved by the Norwegian  
127 Food Safety Authorities (FOTS # 11730/17/18948) and performed accordingly.

128

### 129 2.3. Exposure and sampling

130 Cod were exposed for two weeks and were injected intraperitoneally once per week (day 0 and day  
131 7) with two different doses: low (1x) and high (20x) dose of PAH and PFAS (Table 1), individually  
132 and in various combinations consisting of the following groups; vehicle control, low PAH, low PFAS,  
133 high PAH, high PFAS, low PAH/low PFAS, high PAH/low PFAS, low PAH/high PFAS and high  
134 PAH/high PFAS (Fig. 1). The stock solutions were prepared in a 1:1 (v/v) mixture of rapeseed oil

135 (Eldorado rapsolje) and PBS and injected at 1 mL/100 g fish. Control cod were injected with solvent  
136 vehicle (1:1 of oil and PBS). The PAH concentrations (1x) were chosen based on PAH levels detected  
137 in Atlantic cod from Tampen and Egersund in the monitoring report of the Institute of Marine  
138 Research (IMR) from 2012.<sup>36</sup> The PFAS concentrations (1x) were chosen based on reported values  
139 in cod samples from the Nordic environment and northern Norwegian mainland.<sup>37, 38</sup> Following two  
140 weeks of exposure, the cod were euthanized and tissue samples were collected and frozen in liquid  
141 nitrogen before being transferred to -80 °C for downstream analyses.

142

#### 143 2.4. Quantitative (real-time) PCR

144 Total RNA was extracted from brain tissues using the Direct-zol™ RNA kit, following the  
145 manufacturer's protocol. Quality of RNA was confirmed by formaldehyde agarose gel electrophoresis  
146 and spectrophotometric analysis. cDNA was generated by following the instruction of the iScript  
147 cDNA synthesis kit (Bio-Rad) and transcripts were amplified using Mx3000P real-time PCR machine  
148 (Stratagene, La Jolla, CA), details are presented in section 1.1 of the supplementary information (SI).

149

#### 150 2.5. Steroid hormone analysis

151 Enzyme immunoassay (EIA) was used to measure the concentration of 17β-estradiol (E2) in plasma  
152 using EIA kit (Cayman Chemical Company, Ann Arbor, MI, USA). Detailed description of hormone  
153 extraction and quantification are presented in section 1.2 of SI.

154

#### 155 2.6. Ultra-Performance Liquid chromatography-mass spectrometry (UPLC-MS/MS)

156 To measure dopamine and its metabolites, samples were prepared and ran, following the protocol of  
157 Bertotto et al.<sup>39</sup> Detailed procedure is presented in section 1.3 of SI.

158

#### 159 2.7. Statistics

160 Statistical analysis was performed on RStudio (version 1.1.456), the statistical difference between  
161 control and exposure groups were determined through one-way ANOVA (and a post-hoc Dunnett's

162 test). To investigate interaction of compounds (and corresponding to mixture exposure), a two-way  
163 analysis of variance (ANOVA), followed by Dunnett's post-hoc test was performed on a linear model  
164 with the significance level set at  $p \leq 0.05$ . Details of the performed calculations and statistical tests  
165 can be found in section 5 of SI. Relationship between biological parameters and their response to  
166 chemical exposures were visualized in a principle component analysis (PCA). The first two principle  
167 components and their factor scores were summarized in a biplot using XLSTAT. Data Analysis and  
168 Statistical Solution for Microsoft Excel. Addinsoft, Paris, France (2017).

169

### 170 **3. Results**

#### 171 3.1. Effects of PAH and PFAS on plasma E2

172 The plasma concentrations of E2 showed apparent dose-specific effects for both PAH and PFAS  
173 exposure groups (Fig. 2). The low PAH dose, increased E2 levels, while no change was observed in  
174 high dose exposure group. For PFAS, a non-significant increase in E2 levels was observed in the high  
175 dose exposure group, but, a decrease was noted in the low dose treatment (Fig. 2). Combined exposure  
176 to low PAH/low PFAS, low PAH/high PFAS and high PAH/high PFAS showed an increasing trend,  
177 significantly so at the later value (Fig. 2).

178

#### 179 3.2. Effects of PAH and PFAS on brain dopamine pathways

180 The concentrations of brain dopamine showed a decreasing trend except to high PAH/high PFAS  
181 exposure. Cod exposed to low PAH, high PFAS and low PAH/low PFAS showed a significant  
182 decrease in dopamine levels (Table 2). One of the major dopamine metabolite, DOPAC was  
183 significantly decreased in cod exposed to high PAH dose. A decreasing trend (non-significant) of  
184 DOPAC was observed in other exposure groups, except for cod exposed to low PAH/low PFAS, high  
185 PAH/low PFAS and high PAH/high PFAS, only the latter showed a significant increase (Table 2).  
186 The second most abundant dopamine metabolite (HVA) did not show any significant change after  
187 exposure to PAH and PFAS, singly, at different doses and their various combinations (Table 2). In



188 addition, a significant increase in DOPAC-dopamine and DOPAC+HVA-dopamine ratios were  
189 observed in cod exposed to low PAH/low PFAS and high PAH/low PFAS (Table 2).

190 In this study, an interactive effect of PAH and PFAS was investigated on the level of dopamine  
191 and its metabolites as being represented in Fig. S5 and Table S3. PAH and PFAS at low dose produced  
192 a strong interactive effect on the levels of DOPAC. In single exposure, there is a mild decrease in  
193 levels of DOPAC, however in the mixture, the level was similar to the control group (Fig. S5D and  
194 Table S3). High PAH with low and high dose of PFAS in a mixture also produced an interactive  
195 effect on DOPAC levels (Fig. S5E and F and Table S3). For HVA, significant interaction was  
196 observed between high PAH and high PFAS in a mixture (Fig. S5C and Table S3). Exposure in the  
197 mixture of low PAH/high PFAS and high PAH/high PFAS showed strong interactive effects on the  
198 levels of dopamine. High PFAS and Low PAH single exposure significantly decreased dopamine  
199 level. However in the mixture, the level was similar to the control group (Fig. S5A and B and Table  
200 S3).

201

### 202 3.3. Effects on brain dopaminergic and estrogenic signaling

203 Significant changes were observed in the expression of dopaminergic (*dat*, *dr1*, and *dr2a*) and  
204 estrogenic (*esr1*, *esrrb* and *cyp19a1b*) signaling genes in cod brain following exposure to PAH and  
205 PFAS, given alone or in combination (Fig. S3 and S4). The high PAH dose decreased *dr1* mRNA  
206 and increased of *dr2a* transcripts. On the other hand, high PFAS decreased both *drs*. Whereas, the  
207 combined exposure scenarios did not change the expression of *dr2a* mRNA. (Fig. S3C and D).  
208 Exposures in the mixture of low PAH/high PFAS and high PAH/high PFAS produced interactive  
209 effects on the expression of *dr1*. In an individual exposure, high PFAS significantly decreased the  
210 expression of *dr1* than both doses of PAH. However, when they occurred in a mixture, the expression  
211 was similar to the control group (Fig. S6G and H and Table S4). In contrast, expression of *dat* was  
212 significantly increased after exposure to high PAH/high PFAS (Fig. 3B and D).

213 The expression of *esr1* was significantly decreased after exposure to high PAH and high PFAS  
214 doses, while combined exposures, including high PAH/low PFAS and low PAH/high PFAS,

215 significantly reduced *esr1* expression (Fig. S4A). Both exposures, in the mixtures of low PAH/high  
216 PFAS and high PAH/high PFAS produced interactive effects, but the former showed a significant  
217 decrease compared to control (Fig. S6A and B and Table S4). Expression of *esrrb* mRNA was  
218 significantly decreased after low PAH, high PFAS and high PAH/low PFAS treatments (Fig. S4B).  
219 However, other combined exposures including low PAH/low PFAS, low PAH/high PFAS and high  
220 PAH/high PFAS showed significant interactive effect on *esrrb* expression (Fig. S6C, D and E and  
221 Table S4). Brain aromatase (*cyp19a1b*) gene expression was also significantly decreased in fish  
222 exposed to both low and high PAH dose and combined exposure of low PAH/high PFAS (Fig. S4C).

223

#### 224 3.4. Principle component analysis (PCA)

225 The relationship between levels of all analysed observation and different exposure groups showed  
226 that the first two factor score (F1 and F2) could accounted 69.12 % of the total variance in the dataset  
227 (Fig. 3). Respectively, F1 and F2 covered 50.5 % and 18.59 % of the total variability, showing that  
228 low PAH, high PFAS and combination exposure including, low PAH/low PFAS, high PAH/low  
229 PFAS and high PAH/high PFAS has a strong positive relationship with dopamine: metabolite ratios  
230 (Fig. 3 and Table S2). Despite of their strong positive association with dopamine: metabolite ratio,  
231 only high PAH/high PFAS exposure showed significant increase in E2 level. The expression levels  
232 of *th*, *dr1*, *esr1*, *esrrb* and *cyp19a1b* mRNA showing a negative relationship with all exposure groups  
233 and strong particularly to those that has highest dopamine: metabolite ratios. F1 and F2, showed  
234 strong negative relationship between *th* (-1.886, 0.022), *dr1* (-1.703, 0.911), *esrrb* (-2.306, 0.139)  
235 mRNA expression and low PAH (0.955, 0.061), high PFAS (0.886, -0.408), low PAH/low PFAS  
236 (0.879, -0.342) and high PAH/low PFAS (0.788, -0.282) exposure groups. In general, *th* showed a  
237 decreasing trend and was negatively correlated to the expression of brain *gnrh2* with F1 value of -  
238 1.886 and 0.142 respectively (Pearson's correlation coefficient,  $r = -0.54$ ) (Fig. 3, Table S2 and S2.2).

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241

#### 242 4. Discussion

243 Several *in vivo* and *in vitro* studies have shown that PFASs and PAHs are respective agonist and  
244 antagonists of ER and subsequently affect ER-mediated responses in vertebrate species.<sup>40,41</sup> To better  
245 understand the underlying processes, alternative pathways by which PAH and PFAS may affect  
246 cellular E2 levels and E2-mediated downstream responses should be investigated. For example, PFOS  
247 and PFOA, despite their weak affinities to ER, have been shown to increase E2 production through  
248 modification of steroidogenic pathways.<sup>42,43</sup> In addition, Cyp pathways contribute to estrogenic  
249 properties displayed by B[a]P, where hydroxylated metabolites such as 9-OH-B[a]P displayed  
250 estrogenic properties or may undergo subsequent metabolic steps to form more estrogenic species.<sup>44</sup>  
251 Nevertheless, there are few studies examining the effects of PAH and PFAS on E2 clearance,  
252 biosynthesis or physiological control mechanisms, including feedback processes. Consequently,  
253 additional biochemical processes that regulate neuronal and gonadal E2 homeostasis may be involved  
254 in previously reported estrogenic and anti-estrogenic responses of PAH and/or PFAS.

255         Herein, the low PAH dose and high PFAS dose produced non-significant increases of plasma  
256 E2 levels. Previous studies have demonstrated an association between liver ER expression and plasma  
257 E2 concentrations, in contrast to brain ER expression patterns after exposure to a xenoestrogen  
258 (nonylphenol, NP), showing that *er* isotype expression in various tissues paralleled other  
259 xenoestrogen biomarkers (such as Vtg) in liver or plasma samples of Atlantic salmon (*Salmo salar*).<sup>45</sup>  
260 In addition, the authors reported differential expression pattern of *er* isotypes in liver and brain of  
261 NP-exposed fish.<sup>45</sup> Regression analysis of brain and liver *er-α* and *er-β* transcripts and *vtg* expression  
262 levels showed a linear relationship between liver *er-α* and *vtg* mRNA, whereas brain *er-β* had limited  
263 linearity with liver *vtg*. Unlike liver *er-α*, both brain and liver *er-β* showed a non-linear relationship  
264 with *cyp19* isotypes in the brain.<sup>45</sup> Despite the high plasma E2 levels measured at low PAH dose,  
265 there was a decrease in the levels of both brain *esr1* and *esrrb* mRNA. In addition, we observed that  
266 the high PFAS dose produced elevated plasma E2 concentrations, while the brain showed decreased  
267 levels of both *esr1* and *esrrb* transcripts. The fact that brain *ers* have limited association with  
268 dopaminergic and estrogenic signaling, mixture exposure scenarios may help in understanding

269 interactions among chemicals. For example, single exposures of high PAH and PFAS significantly  
270 decreased the expression of *esr1*, compared to their mixture, suggesting antagonistic interactions  
271 among these chemicals at the tested doses. On the other hand, exposure to low PAH/high PFAS  
272 significantly reduced *esr1* mRNA. Similar antagonistic and chemical masking effects were observed  
273 with *esrrb* by low PAH/low PFAS, low PAH/high PFAS and high PAH/high PFAS mixtures,  
274 suggesting a complex interaction mode by these chemicals.

275 Inconsistency in relationship between brain ERs expression and plasma E2 levels may suggest  
276 that other cellular pathways play important roles in regulating GnRH2 release in the brain or that  
277 neuronal ER expression does not parallel cellular E2 function and regulation. This speculation is  
278 supported by our observation that changes in *gnrh2* expression paralleled changes in plasma E2  
279 levels, with Pearson's correlation coefficient between dopamine metabolite ratio and *gnrh2*  
280 expression showing an  $r = 0.86$  and  $0.81$  (Section 2.2 SI). Modulation of dopamine and its metabolites  
281 after exposure to PAH and PFAS, singly or in combination, suggests a disruption in dopaminergic  
282 function, which may subsequently disrupt E2 synthesis and regulation through feedback  
283 mechanisms.<sup>23</sup> The high PAH treatment decreased the expression of *dr1* with a concomitant increase  
284 in *dr2a*. A decrease in *gnrh2* mRNA was observed in the brain. In rainbow trout (*O. mykiss*), binding  
285 of dopamine to Dr2a inhibited the production of GnRH-stimulated gonadotropin release from the  
286 pituitary.<sup>46-50</sup> Elsewhere in goldfish (*Carassius auratus*), the release of GnRH-activated LH was  
287 blocked through Dr2a activation.<sup>51</sup> In contrast, dopaminergic signaling via Dr2a was reported in  
288 *Tilapia zillii*, where an increase of *dr2a* mRNA paralleled increase in plasma E2 levels<sup>47,48,52</sup>,  
289 suggesting the presence of ER responsive elements (ERE) in the promoter of *dr2a*.<sup>53</sup> In the present  
290 study, we observed a decrease in *dr2a* mRNA with a corresponding increase in plasma E2 for high  
291 PFAS and low PAH/ low PFAS treatment. Other combined exposures (high PAH/low PFAS, low  
292 PAH/high PFAS, and high PAH/high PFAS) did not produce any change in the expression of brain  
293 *dr2a* transcripts.

294 For dopaminergic signaling, we observed a reciprocal association between expression of *dr1*  
295 and *dr2a* in the high PAH exposure group. Combined exposure, especially high PAH/low PFAS also

296 decreased the expression of *dr1*. In previous studies using rat prefrontal cortex, reduced expression  
297 of *Dr1* was observed which paralleled an increase in DR1 protein after exposure to PFOS.<sup>54</sup> In  
298 contrast, Pereiro et al.<sup>55</sup> reported a respective decrease and increase of *Dr1* mRNA and protein  
299 expressions in rat hippocampus after exposure to PFOS. Potential inconsistencies in the expression  
300 of transcript and protein might be due to an inhibitory effect of microRNA (miR-142-3p) that post-  
301 transcriptionally regulates *Dr1*.<sup>56</sup> This assumption is supported by recent studies showing that other  
302 regulatory micro RNAs, such as miR-326 and miR-9, which also control *DR* expression, were  
303 inhibited after exposure to PFAS.<sup>57</sup> A significant interaction among chemicals also affected the  
304 expression of *dr1*. High PFAS significantly decreased *dr1* expression. However, combined with  
305 PAHs in the mixtures of low PAH/high PFAS and high PAH/high PFAS, PAHs masked the inhibitory  
306 effect of high PFAS, enhancing *dr1* expression which was similar to control.

307 Modulation of *dr* transcription may ultimately affect dopamine and its metabolites (HVA and  
308 DOPAC) in the brain. In humans, alteration of dopaminergic-signaling was assessed by measuring  
309 dopamine metabolites in plasma and urine.<sup>58</sup> Thus, the ratio of DOPAC:dopamine, as well as  
310 DOPAC+HVA:dopamine was used in the present study, to estimate the release and turnover of  
311 dopamine. The low PAH/low PFAS exposure significantly increased these ratios and paralleled an  
312 increase in plasma E2. Except for high PAH/low PFAS and low PAH/high PFAS, all exposure groups  
313 produced similar, but non-significant patterns between E2 and dopamine metabolite ratios. Contrary  
314 to individual chemicals, mixture exposures might have interactive effects at specific concentrations  
315 that regulate dopamine metabolism differently, thus violating their association with dopamine  
316 turnover. Previously, Bertotto et al.<sup>39,59</sup> used these ratios to determine dopamine turnover in zebrafish  
317 embryos and juveniles, showing that a low ratio of dopamine and its metabolites demonstrated a  
318 relationship with dopamine turnover.

319 In the HPG axis, feedback mechanisms regulate the endocrine physiology of vertebrates,  
320 including teleosts. Exposure to low PAH and high PFAS and combined low PAH/low PFAS triggered  
321 a mild increase in plasma E2 levels and subsequently decreased brain dopamine concentrations. On  
322 the other hand, high PAH and low PFAS exposures did not produce significant changes that

323 corresponded with plasma E2 levels. It should be noted that tyrosine hydroxylase (Th) plays an  
324 important role as the rate-limiting enzyme in the production of dopamine.<sup>28</sup> We observed a reciprocal  
325 association between dopamine metabolite ratios and *th* expression in fish exposed, either to a low and  
326 high PAH and PFAS singly or in their various combinations (Pearson's correlation coefficient,  $r = -$   
327 0.652 and -0.701, section 2.2 SI). For example, combined high PAH/low PFAS produced a decrease  
328 of *th* mRNA with a concomitant increase of both DOPAC:dopamine and DOPAC+HVA:dopamine  
329 levels. In contrast, there is a direct relationship between the expression of *th* and dopamine  
330 concentration in the brain. Elsewhere, Kumer and Vrana<sup>60</sup> reported that the expression of *th* is  
331 regulated through a negative feedback loop in dopaminergic signaling, where elevated concentrations  
332 of catecholamine down-regulated *th* mRNA expression, leading to a decrease in dopamine levels.

333 Locally produced brain E2 through aromatase (Cyp19b) activity plays an important role in  
334 reproductive- and neuroendocrine functions, and socio-sexual behavior in fish.<sup>61, 62</sup> The entire process  
335 involves multiple factors, including E2, dopamine and their receptors which regulate the expression  
336 of *cyp19a1b*.<sup>26</sup> Exposure of radial glial cells (RGCs) to a Dr1 agonist upregulated *cyp19a1b*  
337 expression through the phosphorylation of cyclic AMP response element binding protein (Creb).<sup>63</sup>  
338 This effect was shown to be enhanced using low E2 (100 nM) concentrations.<sup>63</sup> However, exposure  
339 to high E2 concentrations decreased *cyp19a1b* expression through a classical negative feedback  
340 mechanism.<sup>26</sup> As discussed above, the observed differences between dopamine receptor transcript  
341 and protein expression data and patterns of Cyp19b regulation reported by Xing et al.<sup>63</sup>, did not  
342 parallel our findings. These differences might be attributed to the direct exposure to E2 that may have  
343 suppressed the physiological feedback loop and should be investigated in more detail.

344 In conclusion, we have demonstrated that exposure to these compounds altered dopaminergic  
345 signaling, including the modulation of dopamine biosynthesis, catabolism and its receptor expression  
346 in the brain of juvenile female Atlantic cod. These changes may affect the HPG axis and apical  
347 endpoints such as reproduction or behavior. Overall, our findings contribute to the understanding of  
348 novel cellular pathways that control steroidogenesis after exposure to PAHs and PFASs, and in  
349 complex contaminant mixture scenarios.

350

351 **Conflict of interest**

352 The authors have no competing interests.

353

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361

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564 **Figure 1.** Three-level design of the study with two factors, PAH and PFAS. Each digit represents the  
565 dose (0=absent, 1=low, and 2=high dose) and values inside circle indicate the contribution of both  
566 factors in one exposure group.

567

568 **Figure 2.** Plasma concentration of 17 $\beta$ -estradiol (E2) in juvenile female Atlantic cod exposed to two  
569 different doses (low and high) of PAH and PFAS, given singly or in combinations (low PAH/low  
570 PFAS, high PAH/low PFAS, low PAH/high PFAS and high PAH/high PFAS). Data are presented as  
571 ng/mL of n=3-8  $\pm$  standard error of the mean (SEM). Groups marked with asterisks (\*) are  
572 significantly different compared with control (p<0.05).

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574 **Figure 3.** Principle Component Analysis (PCA) of measured biological parameter, two-  
575 dimensionally visualized on an x-y scatter plot, with a combined factorial score of 69.12%. The data  
576 included is plasma 17 $\beta$ -estradiol levels, brain dopamine metabolite ratios and expression of  
577 dopaminergic and estrogenic signalling gene in the brain of juvenile Atlantic cod following exposure  
578 to PAHs and PFASs at different doses and their various combinations.

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590 **Table 1:** Overview of exposure chemicals\* and their concentrations for *in vivo* exposure.

	Compound	CAS No.	µg/kg		% of total
			Low Dose (1x)	High Dose (20x)	
<b>PAH</b>	Naphthalene	50-32-8	12.64	252.8	31.6
	Phenanthrene	132-65-0	8.38	167.6	21.0
	Dibenzothiophene	86-73-7	0.58	11.6	1.4
	Pyrene	91-20-3	1.45	29.0	3.6
	BaP	85-01-8	1.93	38.5	4.8
	Fluorene	129-00-0	15.03	300.5	37.6
	<b>Total dose</b>			<b>40</b>	<b>800</b>
<b>PFAS</b>	PFOS	2795-39-3	25	500	48.3
	PFTTrA	375-95-1	16.95	339	32.8
	PFNA	335-67-1	5.925	118.5	11.5
	PFOA	72629-94-8	3.825	76.5	7.4
	<b>Total dose</b>			<b>51.7</b>	<b>1034</b>

591 \* All chemicals were purchased from Sigma Aldrich (St. Louis, MO, USA)

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607 **Table 2:** Concentration of dopamine, homovanillic acid (HVA) and 3,4-dihydroxyphenylacetic acid  
608 (DOPAC) as well as dopamine metabolite ratio (DOPAC/Da and DOPAC+HVA/Da) in the brain of  
609 female juvenile Atlantic cod exposed to different doses of PAH and PFAS, singly or in combination,  
610 analysed by UPLC-MS/MS.

Exposure group	pg/mg		ng/mg	Ratio	
	Dopamine	HVA	DOPAC	DOPAC/Da	DOPAC+HVA/ Da
Control	49.39 ± 4.39	182.53 ± 13.22	28.68 ± 1.56	0.615 ± 0.06	0.699 ± 0.08
Low PAH	<b>36.28 ± 1.46*</b>	178.60 ± 11.90	23.55 ± 1.57	0.669 ± 0.126	0.809 ± 0.12
High PAH	38.79 ± 3.24	198.61 ± 16.65	<b>18.23 ± 1.34*</b>	0.484 ± 0.06	0.638 ± 0.056
Low PFAS	46.56 ± 2.32	184.33 ± 35.59	24.33 ± 1.81	0.539 ± 0.041	0.629 ± 0.051
High PFAS	<b>33.93 ± 1.13*</b>	143.84 ± 5.65	28.79 ± 1.66	0.855 ± 0.06	0.991 ± 0.058
Low PAH/Low PFAS	<b>32.51 ± 3.59*</b>	133.59 ± 7.13	30.77 ± 1.09	<b>1.01 ± 0.106*</b>	<b>1.174 ± 0.139*</b>
High PAH/Low PFAS	38.80 ± 0.43	156.48 ± 13.29	36.80 ± 1.57	<b>0.965 ± 0.09*</b>	<b>1.076 ± 0.10*</b>
Low PAH/High PFAS	44.41 ± 3.27	199.74 ± 8.91	21.60 ± 1.58	0.512 ± 0.054	0.625 ± 0.063
High PAH/High PFAS	56.39 ± 3.43	258.15 ± 30.62	<b>43.52 ± 1.56*</b>	0.779 ± 0.065	0.868 ± 0.056

611

612 Concentration of both dopamine and HVA are given in pg/mg, while DOPAC is given in ng/mg. Each  
613 value represents the mean (n = 6-9 ± standard error of the mean [SEM]). Ratio data are presented in  
614 decimal of thousandth digit. Asterisk represent significant difference between control and exposure  
615 groups at p≤0.05.

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PAH (Factor A)

0

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2

PFAS (Factor B)

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1

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0/0

1/0

2/0

0/1

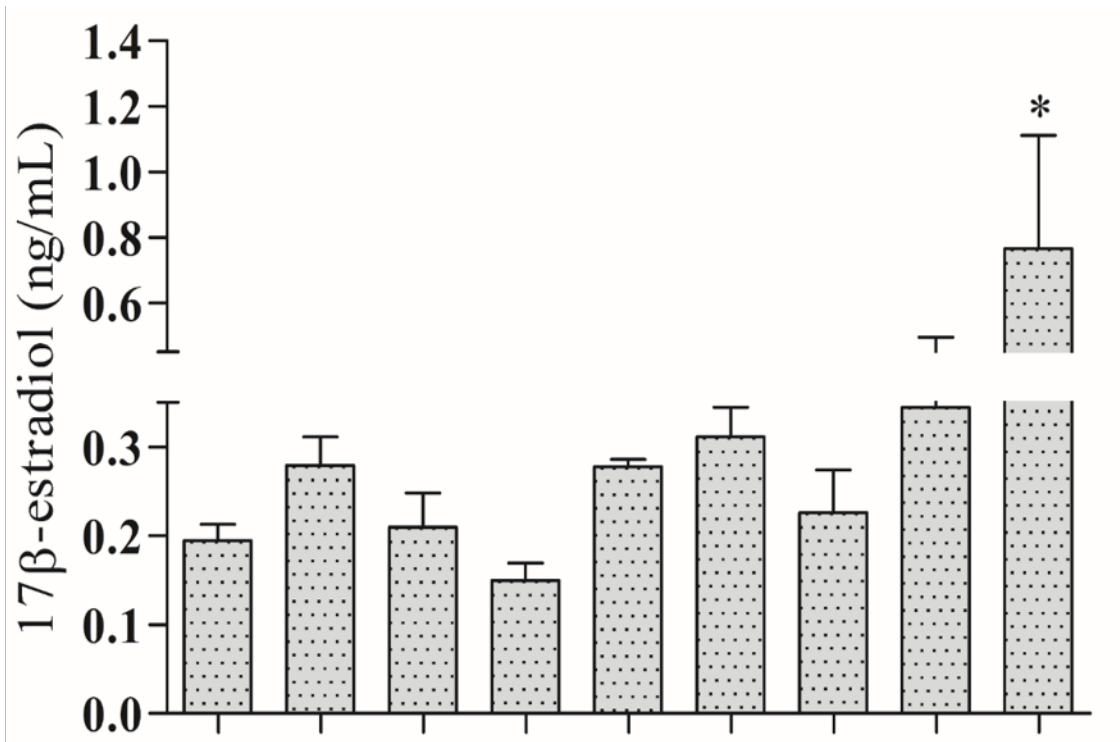
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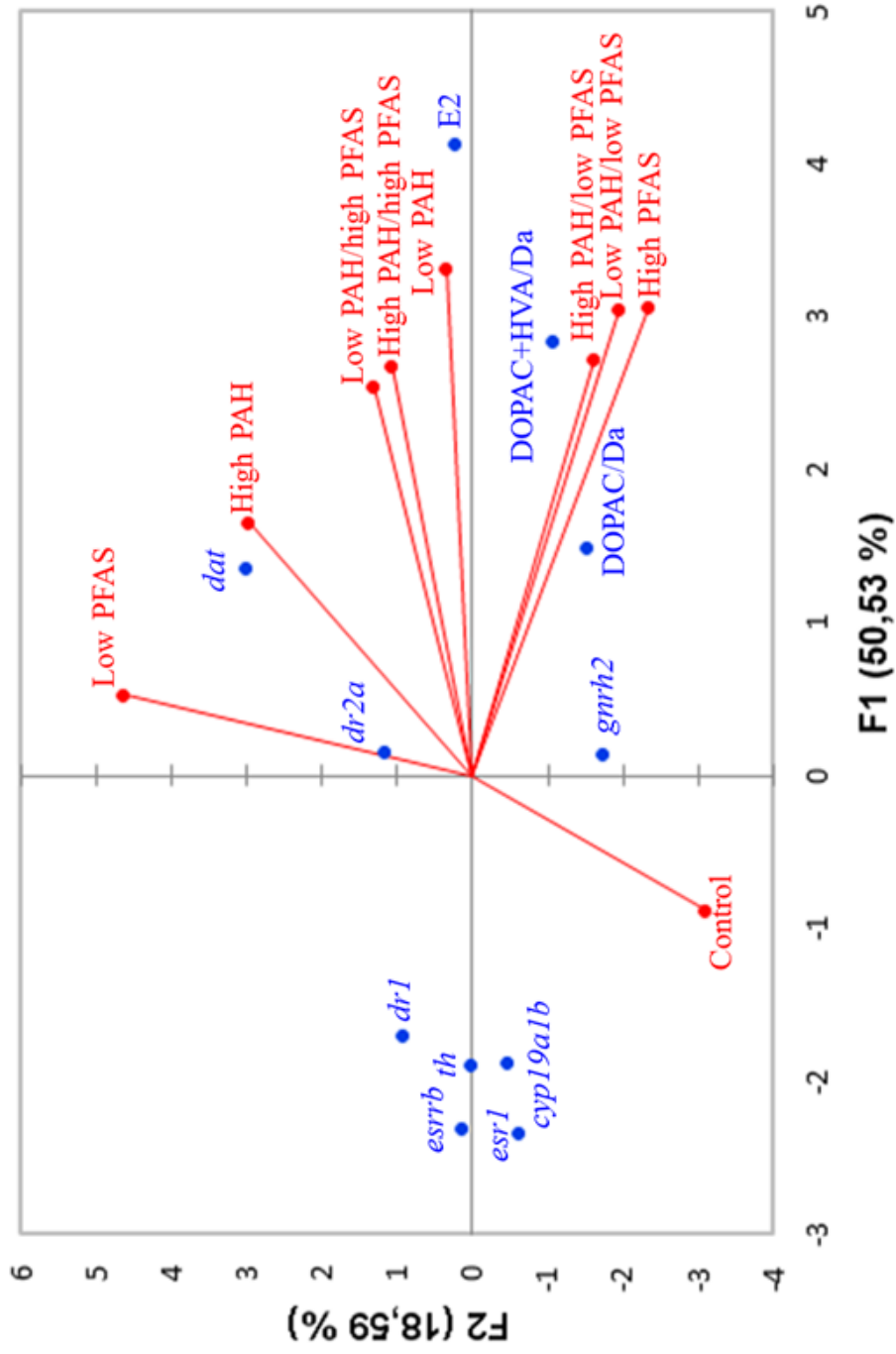
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PAH	Low	-	+	-	-	-	+	-	+	-
	High	-	-	+	-	-	-	+	-	+
PFAS	Low	-	-	-	+	-	+	+	-	-
	High	-	-	-	-	+	-	-	+	+



**Biplot (axes F1 and F2: 69,12 %)**



• Active variables • Active observations

1 **Supporting information (SI)**

2 **Neuro-dopamine homeostasis of juvenile female Atlantic cod (*Gadus morhua*) exposed**  
3 **to polycyclic aromatic hydrocarbons and perfluoroalkyl substances**

4  
5 Essa A. Khan<sup>1</sup>, Luisa B. Bertotto<sup>2</sup>, Karina Dale<sup>3</sup>, Roger Lille-Langøy<sup>3</sup>, Fekadu Yadetie<sup>3</sup>, Odd  
6 André Karlsen<sup>3</sup>, Anders Goksøyr<sup>3</sup>, Daniel Schlenk<sup>2</sup>, Augustine Arukwe<sup>1\*</sup>

7  
8 <sup>1</sup>Department of Biology, Norwegian University of Science and Technology (NTNU),  
9 Høgskoleringen 5, N-7491 Trondheim, Norway

10 <sup>2</sup> Department of Environmental Sciences, University of California-Riverside, Riverside,  
11 California, USA

12 <sup>3</sup> Department of Biological Sciences, University of Bergen, N-5020 Bergen, Norway

13  
14 **\*Corresponding author:**

15 Augustine Arukwe

16 Department of Biology

17 Norwegian University of Science and Technology (NTNU)

18 Høgskoleringen 5, N-7491 Trondheim, Norway

19 Telephone: +47 99552728

20 E-mail: [augustine.arukwe@ntnu.no](mailto:augustine.arukwe@ntnu.no)

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29 **1. Methodology**

30 **1.1. Biometric data**

31

32 **Table S1.** Biometric data of experimental fish used in the present study. Data are presented as  
33 mean (n = 6-9) ± standard error of the mean (SEM).

Exposure group	Biometric data		
	Fish weight (g)	Fish length (cm)	k-factor
Control	177.54 ± 5.56	26.84 ± 0.26	0.91 ± 0.02
Low PAH	167.27 ± 4.94	26.87 ± 0.21	0.85 ± 0.01
High PAH	174.18 ± 7.79	26.72 ± 0.36	0.90 ± 0.01
Low PFAS	180.81 ± 8.72	26.81 ± 0.32	0.92 ± 0.02
High PFAS	154.68 ± 6.27	25.93 ± 0.32	0.88 ± 0.03
Low PAH/Low PFAS	175.14 ± 7.07	26.81 ± 0.35	0.90 ± 0.02
High PAH/Low PFAS	155.90 ± 7.41	25.89 ± 0.35	0.88 ± 0.01
Low PAH/High PFAS	172.33 ± 7.53	26.59 ± 0.32	0.90 ± 0.01
High PAH/High PFAS	182.75 ± 7.43	27.13 ± 0.32	0.90 ± 0.01

34

35

36 **1.2. Quantitative (real-time) PCR**

37 Specific primer pair sequences (Table S2) for *th*, *dat*, *drd1*, *drd2a*, *er-a*, *esrrb*, *cyp19a1b* and  
38 *gnrh2* genes were amplified using the Mx3000P real-time PCR machine (Stratagene, La Jolla,  
39 CA). The primer pairs were tested by analyzing single amplified product of expected size for  
40 individual genes. A parallel control, lacking cDNA template was used to validate the specificity  
41 and target sequence amplification. PCR program includes an enzyme activation step at 95 °C  
42 (4 min) followed by 40 cycles of 95 °C (15 s), 60 °C (30 s) and 72 °C (15 s) and last step  
43 temperature profile include 95 °C (60 s), 65 °C (30 s) and 95 °C (30 s). Expression of each  
44 gene was determined by following the well-validated procedure of absolute quantification in  
45 our laboratory.<sup>1</sup> A known amount of plasmid cloned with an amplicon of interest used to

46 generate a standard curve. The pre-made standard plot of cycle threshold (Ct) versus log copy  
 47 number were used to quantify the expression of the target gene in unknown samples.

48  
 49 **Table S2:** Primer pair sequences with amplicon size and annealing temperature conditions for  
 50 genes quantified with real-time PCR.

Target gene	Accession no.	Primer sequence*		Amplicon size (bp)	Annealing temp. (°C)
		Forward	Reverse		
<i>th</i>	ENSGMOG0	ACCAGTGGCTGGTT	GTGACCCAGAAGC	142	62
	0000017881	TGTT	TCATGTAT		
<i>dat</i>	ENSGMOG0	CTCCAAGCTATGGT	GCTATTCTATCGC	142	62
	0000006703	TCGTACAC	AGAACTTCCC		
<i>drd1</i>	ENSGMOG0	CTTCATCCTCAACT	GCGTTGAAGGCGT	147	62
	0000007704	GCATGGT	AGATGAT		
<i>drd2a</i>	ENSGMOG0	CCACCTCGCTGAAG	CTCATCCAGTTCC	152	62
	0000006531	GATAAG	AGGTCTTC		
<i>er-a</i>	ENSGMOG0	ATCTTCGCACAAGA	CCTTGAGACAGAC	142	62
	0000014898	CCTCATC	AAACTCCTC		
<i>esrrb</i>	ENSGMOG0	AAGCGGCAGGAGG	GGATGCTCCGCTT	146	62
	0000015180	AGAG	GAAGAA		
<i>cyp19a1b</i>	ENSGMOG0	CTGGAAGAAAGTG	CACAGATCCCCAC	145	51
	0000010165	AGGGCATATTT	GGTTCTC		
<i>gnrh2</i>	ENSGMOG0	TACCCTGGAGGAA	TGGCCAGGACATC	145	62
	0000009002	AGAGAGAG	CATAAAG		

51 \* Primer sequences in 5' to 3' direction.

52

### 53 1.3. Steroid hormone analysis

54 Steroid hormones were extracted from plasma with organic solvent. Plasma sample (100 µL)  
 55 was mixed thoroughly with diethyl ether (4:1 volume of plasma), and two phases were allowed  
 56 to separate by vortexing. The organic phase containing steroid hormones was transferred into  
 57 new glass tube whereas the frozen aqueous phase was extracted again. The combined extract  
 58 was allowed to evaporate at 30 °C, and the dry extract was reconstituted in 100 µL EIA buffer  
 59 by vortexing and stored at 80 °C until analysis. EIA for E2 was performed by following the  
 60 manufacturer's guideline. The plate was read using a Bio-Tek Synergy HT microplate reader  
 61 (Bio-Tek Instruments, Winooski, VT, USA) at 410 nm. A standard curve was made by a 4-  
 62 parameter logistic fit between log concentration and logit transformation of B/Bo (Bound  
 63 sample/maximum bound) and expressed as ng/mL.

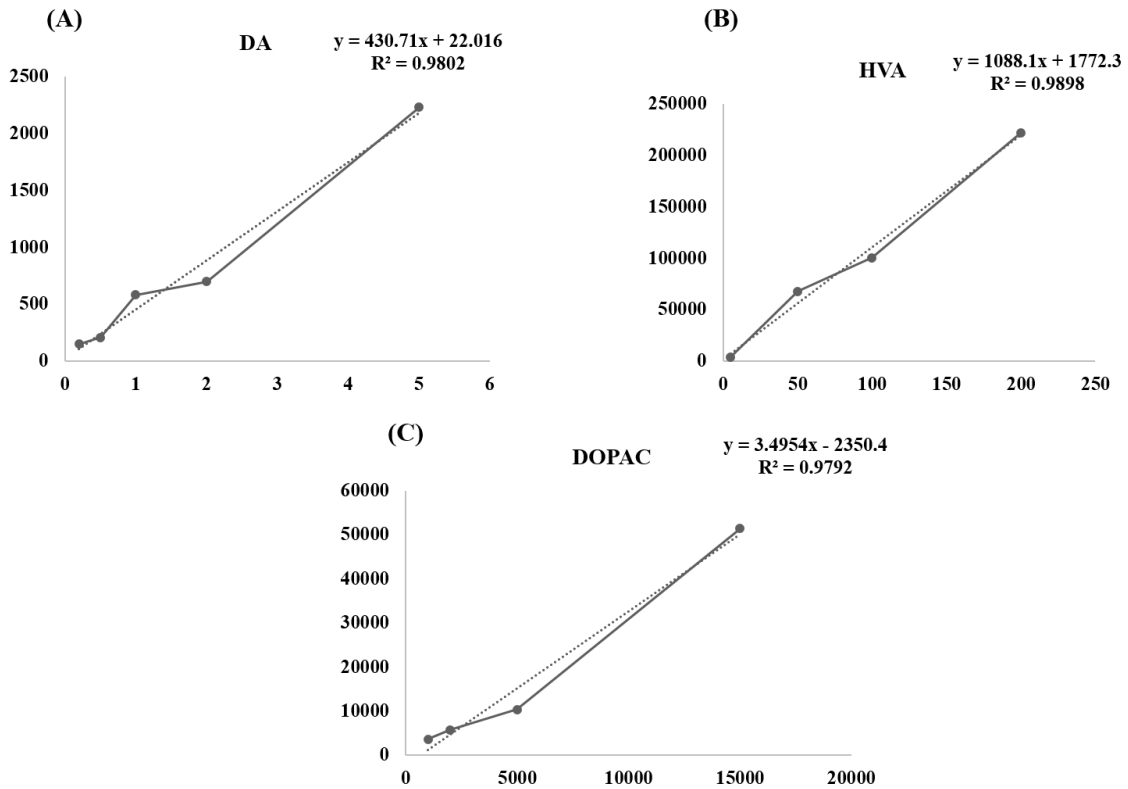
64

### 65 1.4. Ultra-performance Liquid chromatography-mass spectrometry

66 The brain was homogenized using 0.1 % ice-cold formic acid (LC-MS grade, Sigma-Aldrich)  
 67 with 1 ng DA-d4 and 2 ng of HVA-d5 per mg tissue as an internal standard. Following

68 centrifugation, the supernatant was subjected to solid-phase extraction with Starta X polymeric  
69 reverse-phase cartridges (33  $\mu\text{m}$ , 60 mg, 3 mL; Phenomenex) preconditioned with 0.1 % formic  
70 acid in different solvents. The order in which cartridge was conditioned includes 0.1 % formic  
71 acid in acetonitrile, followed by 0.1 % formic acid in methanol and 0.1 % formic acid in water.  
72 The analyte was eluted with 0.1 % formic acid in acetonitrile/methanol (1:1, v/v), evaporate  
73 the organic solvent with a stream of nitrogen gas and finally reconstituted in 0.1 % formic acid  
74 in water. The extract was ready for analysis through Waters ACQUITY ultra-performance  
75 liquid chromatography (UPLC) coupled with Micromass Triple Quadrupole mass spectrometer  
76 (qQq) equipped with an electrospray ionization (ESI) interface (Waters, Milford, MA). An  
77 injection volume of 5  $\mu\text{L}$  was passed through ACQUITY UPLC HSS T3, 2.1 mm  $\times$  100 mm,  
78 1.7  $\mu\text{m}$  column at the flow rate of 0.3 mL min<sup>-1</sup> with the gradient of two solvents; solvent A  
79 (0.1 % formic acid in DI water, 18 $\Omega$ ) and solvent B (0.1 % formic acid in an equal volume of  
80 acetonitrile/methanol). The gradient program starts from 95 % A and 5 % B for 1 min and  
81 ramped to 40 % A over the course of 1 min, it was further decreased to 10 % for 1.5 min then  
82 linearly decreased to 0 % for 0.5 min and finally ramped back to 95 % A involving intermediate  
83 step in which there is linearity increase to 10 % for 0.5 min and stayed at 95 % A, 0.5 min for  
84 equilibration. The specific instrument setting was as follows: source temperature 150 $^{\circ}\text{C}$ ,  
85 desolvation temperature 600  $^{\circ}\text{C}$ , capillary source voltage 3.00 kV, dwell time 0.028 s, cone gas  
86 150 L h<sup>-1</sup> and desolvation gas 1200 L h<sup>-1</sup>, the collision gas was 99.9% pure argon. Cone and  
87 collision voltage of 50 V and 20 V respectively were generated using IntelliStart software  
88 (Waters). Individual compound peaks were detected and integrated using TargetLynx XS  
89 software.

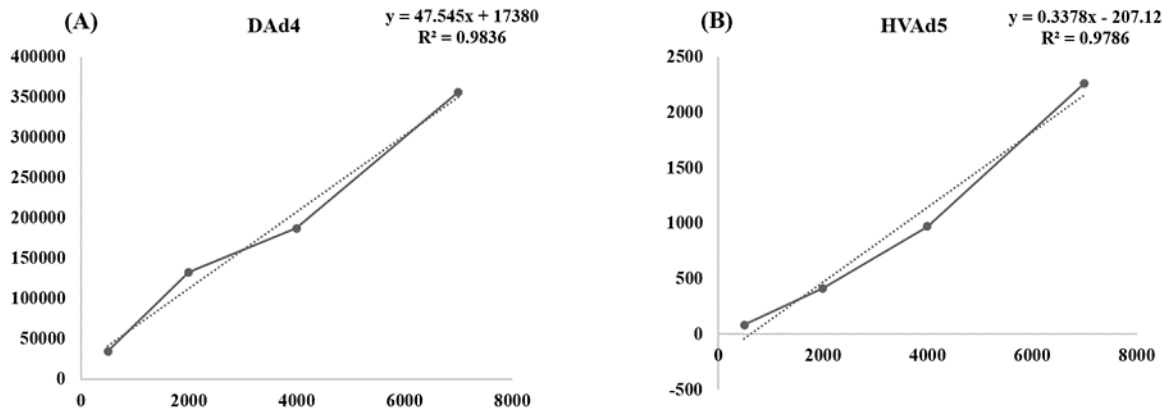
90 A linear calibration curve in a range of 0.2 - 5 ppb for dopamine, from 5 – 200 ppb for  
91 HVA and from 1 – 15 ppm for DOPAC (Sigma-Aldrich), purity >98 %) with  $r^2 > 0.97$  was  
92 used to quantify levels of these metabolites in the brain (Figure S1A, B and C). A deuterated  
93 derivative of dopamine-1,1,2,2-d<sub>4</sub> (DA-d<sub>4</sub>) and 4-hydroxy-3methoxyphenyl-d<sub>3</sub>-acetic-d<sub>2</sub> acid  
94 (HVA-d<sub>5</sub>) were purchased from Sigma-Aldrich and used as an internal standard for dopamine  
95 and HVA respectively (Figure S2A, B). The same HVA-d<sub>5</sub> standard was used for DOPAC.  
96 Recoveries were determined by spiking internal standard at a concentration of 4ppm of DA-d<sub>5</sub>  
97 and 8 ppm of HVA-d<sub>4</sub> per 100 mg of brain tissue.



98

99 **Figure S1:** Calibration curve of DA, HVA and DOPAC spiked at specific concentration range  
 100 through UPLC-MS/MS with  $r^2 > 0.97$ .

101



102

103 **Figure S2:** Calibration curve of deuterated derivative, DA-d5 and HVA-d4 spiked at specific  
 104 concentration range through UPLC-MS/MS with  $r^2 > 0.97$ .

105

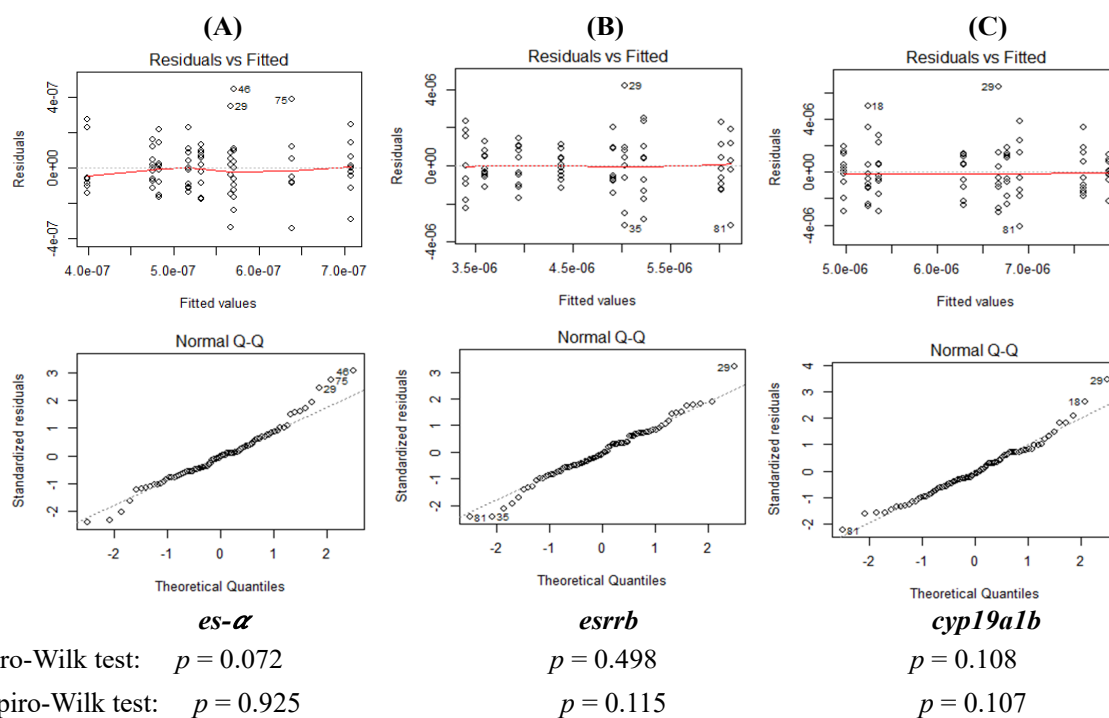
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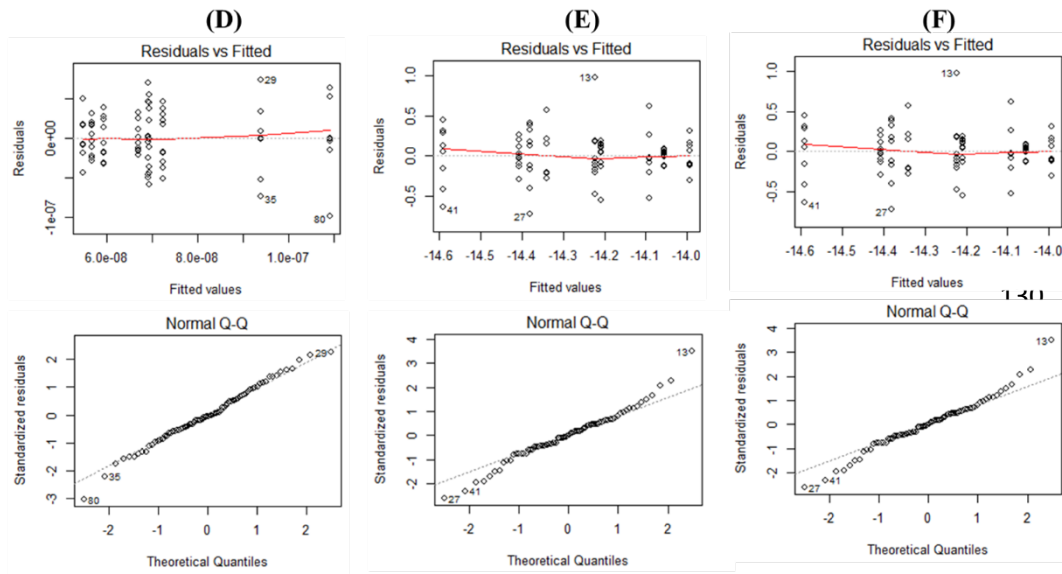
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109 **2. Statistical analysis**

110 To determine effect of both compounds, separately and their interactions (corresponding to the  
 111 mixtures), three-level factorial design, statistically analysed using R-Studio (version 1.1.456).  
 112 A one-way analysis of variance (ANOVA) followed by a Dunnett’s post-hoc test was  
 113 performed to test effect of all treatment groups separately, comparison against a single control  
 114 group (nine levels). A two-way analysis of variance (ANOVA) was performed on a linear  
 115 model to test an interaction effect of compounds on parameter of interest. To normalize the  
 116 residuals of model, data was transformed using natural logarithm and visually inspected using  
 117 quantile-quantile and histogram plots, as well as Shapiro-Wilk test. Some of the groups belong  
 118 to E2 and DOPAC data violated Shapiro-Wilk test, therefore applied Kolmogorov-Smirnov  
 119 test was applied to check normality (Figure S3). Homogeneity of variance was determined by  
 120 Levene’s test. The data belongs to E2, HVA and *drd2a*, violated Levene’s homogeneity test,  
 121 therefore Brown-Forsythe and Welch’s heteroscedastic F-test was applied. Gene expression  
 122 data of *drd1*, an outlier belonging to the Low PAH/Low PFAS treatment group was removed  
 123 from dataset before logarithm transformation. The dopamine catabolites, DOPAC, gene  
 124 expression data of *drd1* and E2 data were logarithmically transformed before analysis.



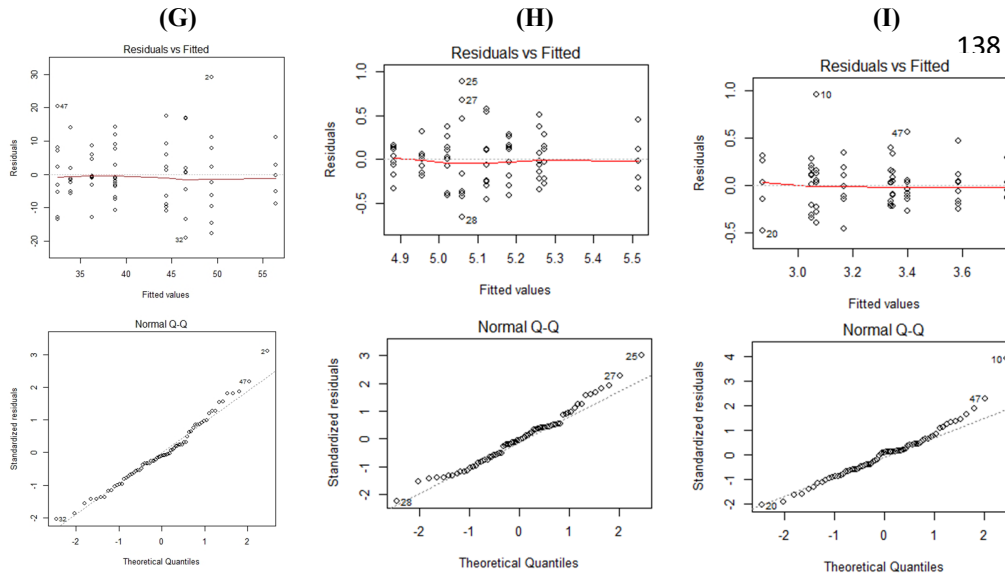


Shapiro-Wilk test:  $p = 0.925$

$p = 0.115$

$p = 0.107$

135  
136  
137



Shapiro-Wilk test:  $p = 0.271$

$p = 0.338$

$p = 0.014$   
147  
148

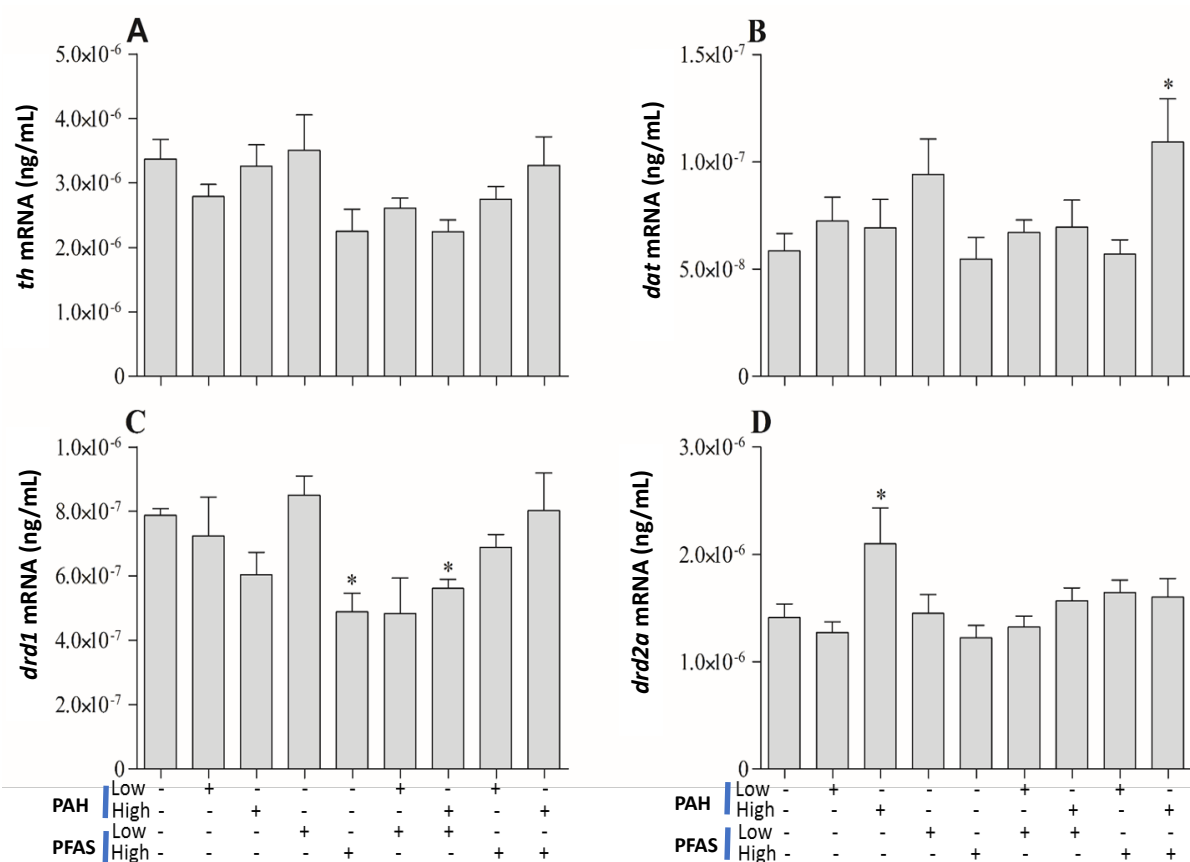
149 **Figure S3:** Normal distribution of gene expression (*er-a*, *esrrb*, *cyp19a1b*, *dat*, *drd1* and *drd2a*  
 150 in A, B, C, D, E and F panels respectively), dopamine and its metabolite data (dopamine, HVA  
 151 and DOPAC in G, H and I panels respectively) visually inspected using quantile-quantile plot  
 152 along with Shapiro-Wilk test,  $p$ -value.

153



154 **3. Results**

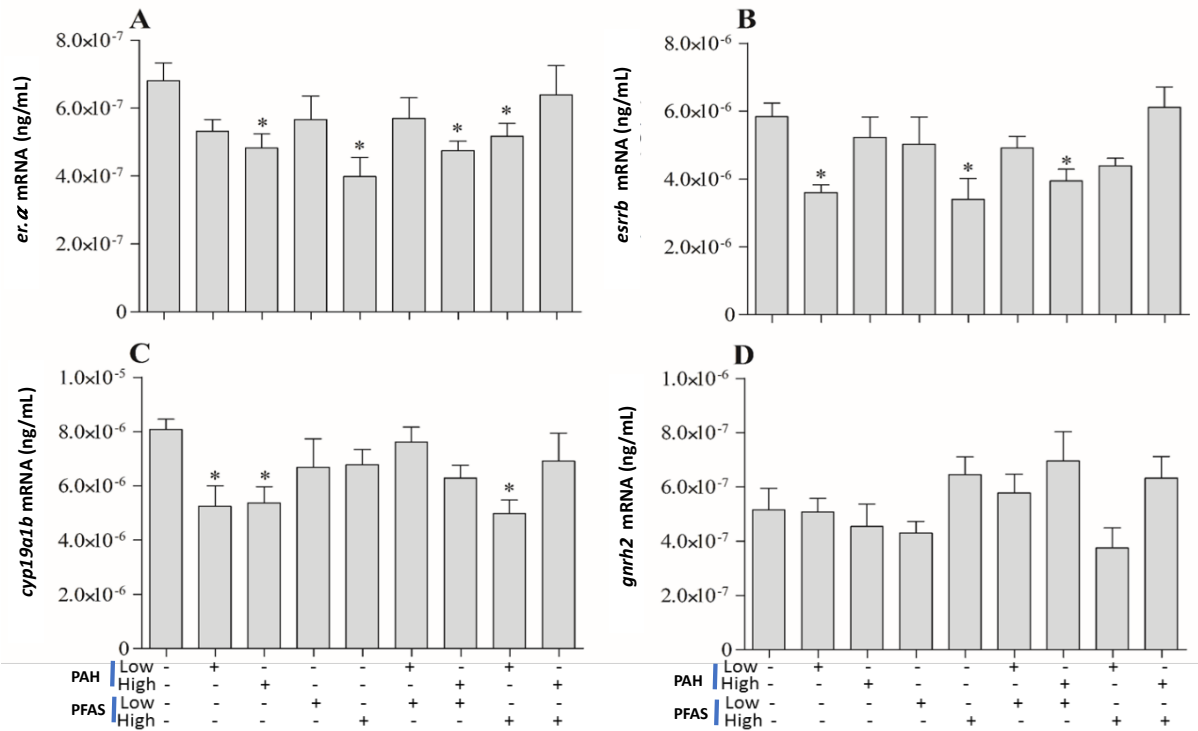
155 **3.1 Expression levels of genes measured in the present study.**



156

157 **Figure S4:** Transcriptional change of tyrosine hydroxylase 1 (*th*: **A**), dopamine active  
 158 transporter (*dat*: **B**), dopamine receptor (*drd1*: **C**) and (*drd2a*: **D**) in the brain of juvenile female  
 159 Atlantic cod exposed two different doses (low and high) of PAH and PFAS, given singly or in  
 160 various combinations (low PAH/low PFAS, high PAH/low PFAS, low PAH/high PFAS and  
 161 high PAH/high PFAS). Data are presented as mean (n = 6-9) ± standard error of the mean  
 162 (SEM). Groups marked with asterisks (\*) are significantly different, compared with control  
 163 (p<0.05).

164



165

166 **Figure S5:** Transcriptional change of estrogen receptors (*er-a*: **A**, *esrrb*: **B**), *cyp19a1b* (**C**) and  
 167 *gnrh2* (**D**) in the brain of juvenile female Atlantic cod exposed two different doses (low and  
 168 high) of PAH and PFAS, given singly or in combinations (low PAH/low PFAS, high PAH/low  
 169 PFAS, low PAH/high PFAS and high PAH/high PFAS). Data are presented as mean (n = 6-9)  
 170 ± standard error of the mean (SEM). Groups marked with asterisks (\*) are significantly  
 171 different, compared with control (p<0.05).

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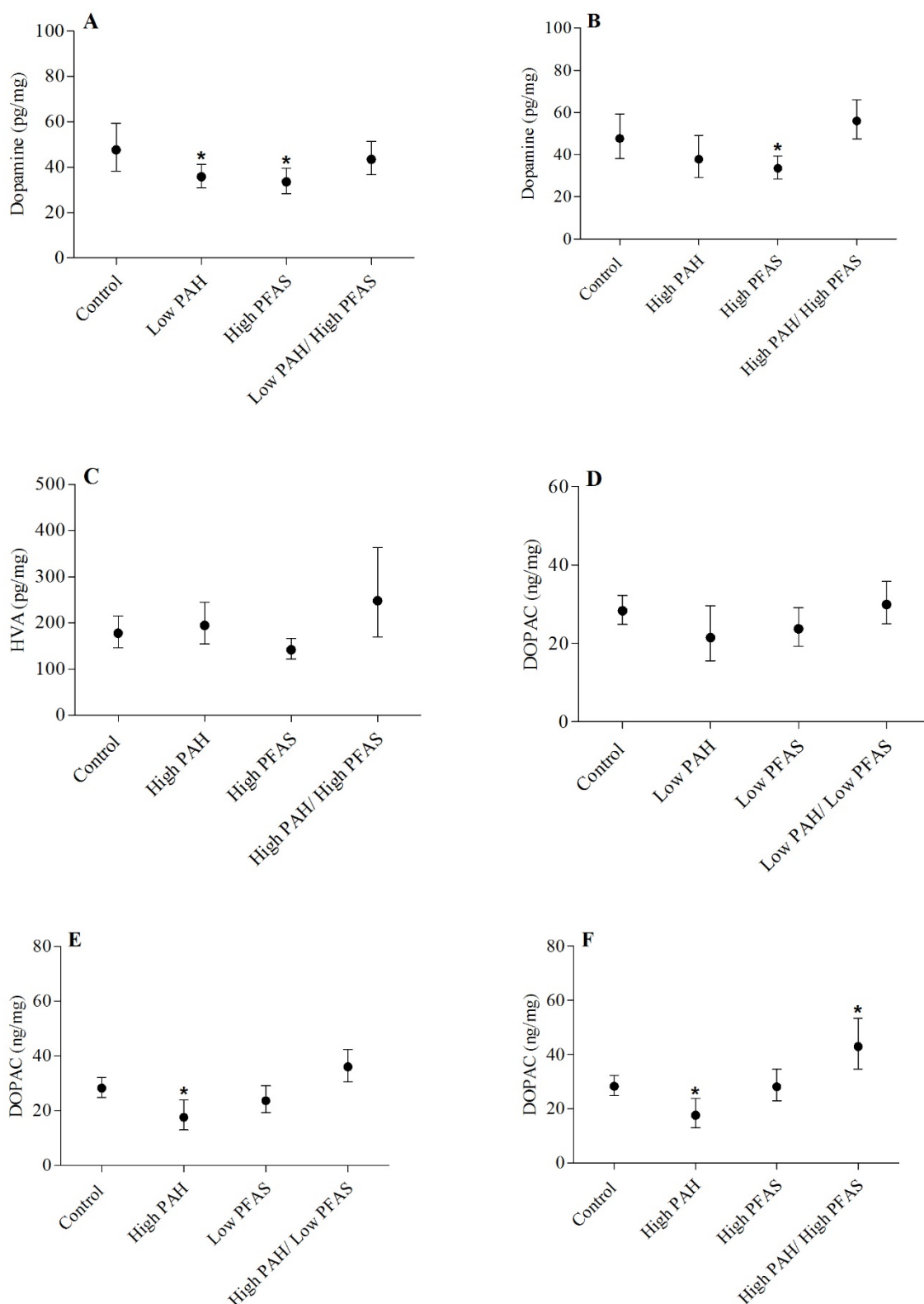
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185

186 **3.2 Significant interaction effects between variables investigated in the present study**



187  
 188 **Figure S6:** PAH and PFAS interaction in mixture significantly affect levels of dopamine ( $p =$   
 189  $0.001$  and  $8.3e-05$ ; **A** and **B**), HVA ( $p = 0.046$ ; **C**) and DOPAC ( $p = 0.005$ ,  $9.7e-06$  and  $4.2e-$   
 190  $05$ ; **D**, **E** and **F**). Data are presented as mean of concentration ( $n = 6-9$ ) and 95 % confidence  
 191 interval. Group marked with asterisks (\*) are significantly different, compared with control  
 192 ( $p \leq 0.05$ ).

193

194 **Table S3:** Significant interaction effect of PAH and PFAS on dopamine and its metabolite  
 195 (DOPAC and HVA). All metabolic data are presented in concentrations, dopamine and HVA  
 196 (pg/mg) and DOPAC (ng/mg) with 95% confidence interval (n = 6-9).

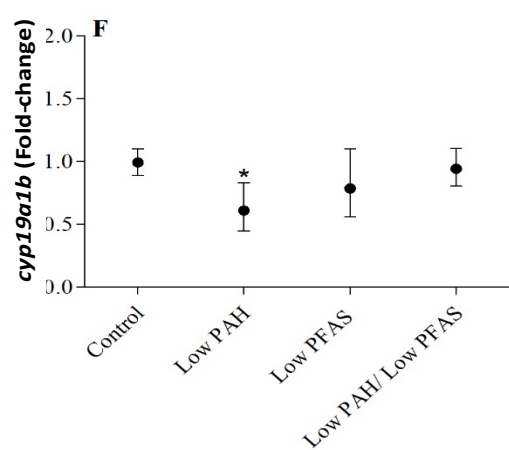
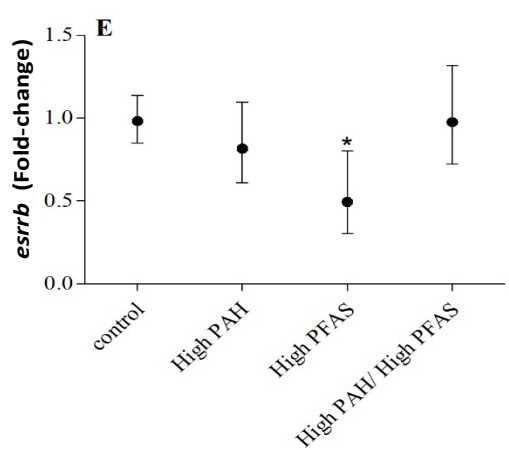
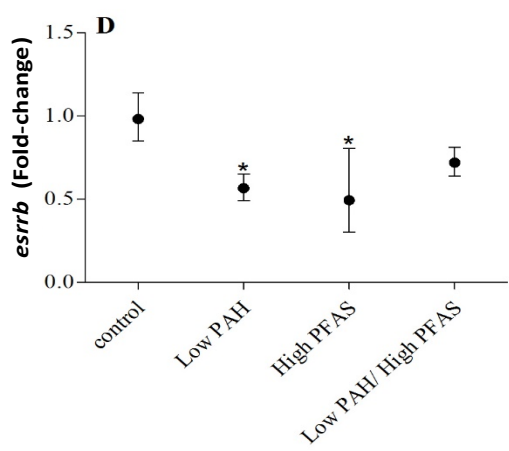
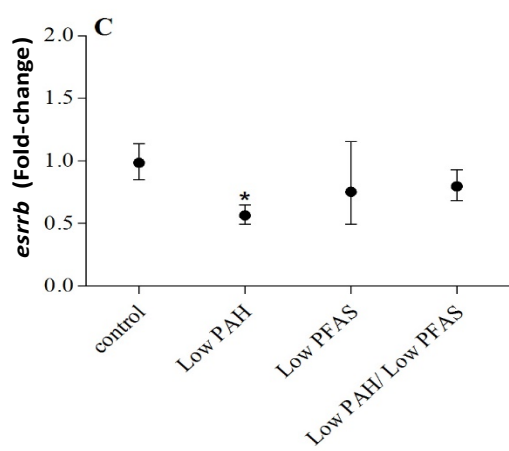
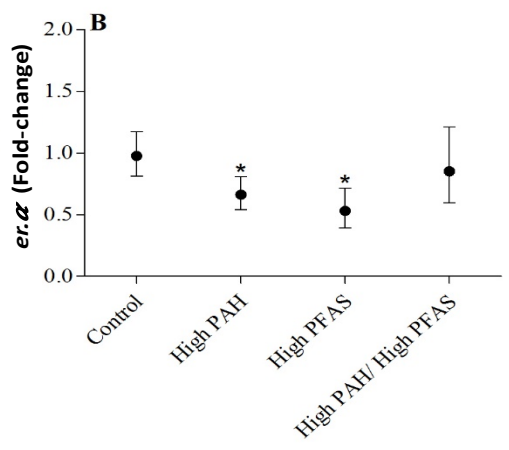
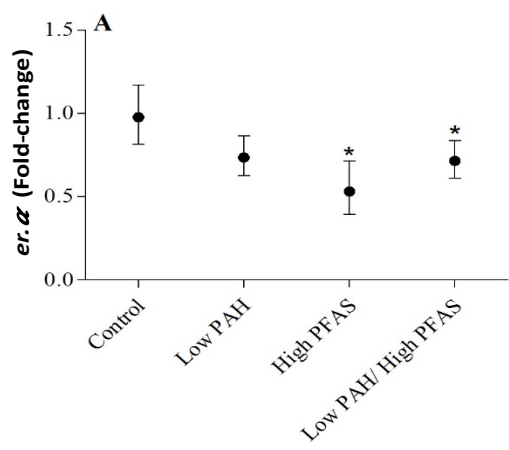
		<b>Conc.</b>	<b>Low 95% CI</b>	<b>High 95% CI</b>
<b>Dopamine</b>	Control	49.39	38.19	60.59
	( <i>p</i> = 0.001) Low PAH	36.28	31.52	41.05
	High PFAS	33.93	27.7	40.16
	Low PAH/ High PFAS	44.41	36.87	51.95
<i>(p</i> = 8.3e-05)	High PAH	38.79	28.46	49.12
	High PFAS	33.93	27.7	40.16
	High PAH/ High PFAS	56.39	46.84	65.94
<b>HVA</b>	Control	182.53	149.58	215.49
	( <i>p</i> = 0.046) High PAH	198.61	153.75	243.48
	High PFAS	143.84	119.63	168.04
	High PAH/ High PFAS	258.15	154.02	362.28
<b>DOPAC</b>	Control	28.68	24.72	32.65
	( <i>p</i> = 0.005) Low PAH	23.55	13.75	33.35
	Low PFAS	24.33	19.60	29.05
	Low PAH/ Low PFAS	30.77	24.20	37.34
<i>(p</i> = 9.7e-06)	High PAH	18.23	13.11	23.35
	Low PFAS	24.33	19.60	29.05
	High PAH/ Low PFAS	36.80	30.10	43.39
<i>(p</i> = 4.2e-05)	High PAH	18.23	13.11	23.35
	High PFAS	28.79	22.48	35.10
	High PAH/ High PFAS	43.52	33.43	53.61

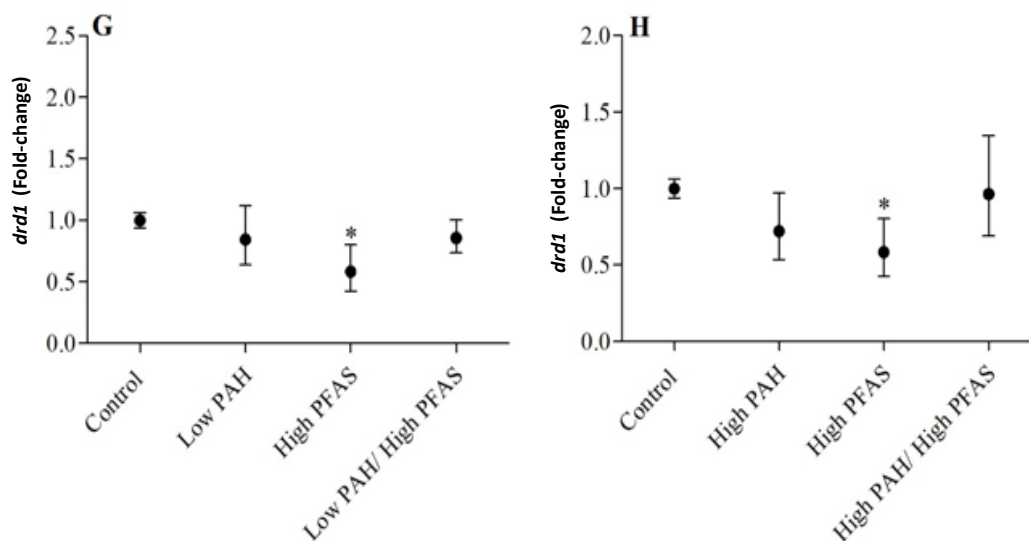
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203 **Figure S7:** PAH and PFAS interaction in mixture significantly affect levels of *er-α* ( $p = 0.004$   
 204 and  $4.7e-05$ ; **A** and **B**), *esrrb* ( $p = 0.015$ ,  $4.0e-04$  and  $6.0e-04$ ; **C**, **D** and **E**), *cyp19a1b* ( $p =$   
 205  $0.008$ ; **F**), *drd1* ( $p = 0.005$  and  $0.0001$ ; **G** and **H**) and *gnrh2* at High PAH/Low PFAS ( $p =$   
 206  $0.031$ ; not graphically presented). Data are presented as mean of fold change ( $n = 6-9$ ) and 95%  
 207 confidence interval. Group marked with asterisks (\*) are significantly different, compared with  
 208 control ( $p \leq 0.05$ ).

209

210 **Table S4:** Significant interaction effect of PAH and PFAS on gene expression of *er-α*, *esrrb*,  
 211 *cyp19a1b* and *drd1*. All expressions are presented relative to the control (set at 1) with 95%  
 212 confidence interval ( $n = 6-9$ ).

		Mean	Low 95 % CI	High 95 % CI
<i>er-α</i>	Control	1	0.83	1.16
	( $p = 0.0049$ ) Low PAH	0.73	0.64	0.86
	High PFAS	0.56	0.37	0.75
	Low PAH/ High PFAS	0.74	0.61	0.85
<i>er-α</i>	( $p = 4.7e-05$ ) High PAH	0.68	0.54	0.81
	High PFAS	0.56	0.37	0.75
	High PAH/ High PFAS	0.90	0.61	1.19
<i>esrrb</i>	Control	1	0.84	1.15
	( $p = 0.015$ ) Low PAH	0.57	0.49	0.65
	Low PFAS	0.83	0.52	1.14

	Low PAH/ Low PFAS	0.84	0.68	0.94
$(p = 4.0e-4)$	Low PAH	0.57	0.49	0.65
	High PFAS	0.56	0.32	0.80
	Low PAH/ High PFAS	0.72	0.64	0.81
$(p = 6.0e-4)$	High PAH	0.86	0.63	1.09
	High PFAS	0.56	0.32	0.80
	High PAH/ High PFAS	1.01	0.77	1.25
<b><i>cyp19a1b</i></b> $(p = 0.008)$	Control	1	0.89	1.10
	Low PAH	0.61	0.44	0.88
	Low PFAS	0.84	0.53	1.16
	Low PAH/ Low PFAS	0.98	0.80	1.12
<b><i>drd1</i></b> $(p = 0.005)$	Control	1	0.94	1.06
	Low PAH	0.90	0.57	1.26
	High PFAS	0.61	0.44	0.79
	Low PAH/ High PFAS	0.87	0.75	0.98
$(p = 0.0001)$	High PAH	0.76	0.56	0.96
	High PFAS	0.61	0.44	0.79
	High PAH/ High PFAS	1.02	0.65	1.38

213

214

215 **4. Principal component analysis**216 **4.1 Correlation matrix (Pearson)**

217 **Table S5:** Correlational matrix (Pearson) showing statistical interactions between studied  
218 variable. Values in bold fonts are significantly different, compared with control at  $\alpha \leq 0.05$ .

Observations	DOPAC/							
	Da	DOPAC+HVA/Da	<i>th</i>	<i>drd1</i>	<i>drd2a</i>	<i>er-<math>\alpha</math></i>	<i>esrrb</i>	<i>gnrh2</i>
DOPAC/ Da	<b>1</b>	<b>0.978</b>	<b>-0.652</b>	-0.620	-0.555	-0.150	-0.227	<b>0.856</b>
DOPAC+HVA/Da	<b>0.978</b>	<b>1</b>	<b>-0.701</b>	<b>-0.703</b>	-0.508	-0.277	-0.328	<b>0.811</b>
<i>th</i>	<b>-0.652</b>	<b>-0.701</b>	<b>1</b>	<b>0.798</b>	0.414	<b>0.706</b>	<b>0.802</b>	-0.543
<i>drd1</i>	-0.620	<b>-0.703</b>	<b>0.798</b>	<b>1</b>	0.127	<b>0.650</b>	0.509	-0.485
<i>drd2a</i>	-0.555	-0.508	0.414	0.127	<b>1</b>	-0.009	0.396	-0.367
<i>er-<math>\alpha</math></i>	-0.150	-0.277	<b>0.706</b>	<b>0.650</b>	-0.009	<b>1</b>	<b>0.827</b>	-0.205
<i>esrrb</i>	-0.227	-0.328	<b>0.802</b>	0.509	0.396	<b>0.827</b>	<b>1</b>	-0.159
<i>gnrh2</i>	<b>0.856</b>	<b>0.811</b>	-0.543	-0.485	-0.367	-0.205	-0.159	<b>1</b>

219

220

221 **Table S6:** Factor score of the variable and observation on the two factorial axes.

		<b>Axes</b>	
	<b>Descriptors</b>	<b>F1</b>	<b>F2</b>
<b>Variable</b>	Control	-0.256	-0.540
	Low PAH	0.955	0.061
	High PAH	0.479	0.521
	Low PFAS	0.153	0.809
	High PFAS	0.886	-0.408
	Low PAH/ Low PFAS	0.879	-0.342
	High PAH/ Low PFAS	0.788	-0.282
	Low PAH/ High PFAS	0.737	0.229
	High PAH/ High PFAS	0.771	0.189
<b>Observation</b>	DOPAC/Da	1.488	-1.541
	DOPAC + HVA/Da	2.835	-1.075
	17 $\beta$ -estradiol	4.136	0.231
	<i>th</i>	-1.886	0.022
	<i>drd1</i>	-1.703	0.911
	<i>drd2a</i>	0.162	1.159
	<i>dat</i>	1.352	2.990
	<i>er-<math>\alpha</math></i>	-2.341	-0.609
	<i>esrrb</i>	-2.306	0.139
	<i>cyp19a1b</i>	-1.879	-0.476
	<i>gnrh2</i>	0.142	-1.753

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230 **References**

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