

1 **Environmental doses of perfluorooctanoic acid change the**  
2 **expression of genes in target tissues of common carp**

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15 **Running Head:** Environmental PFOA changes gene expression in target tissues

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25 **Abbreviations:** PFOA, perfluorooctanoic acid; GST, glutathione S-transferase; RT-qPCR, real-time  
26 quantitative PCR; LOD, limit of detection; HPG, hypothalamic-pituitary-gonadal.

## 28 **Abstract**

29 We aimed to evaluate the effects of environmental doses of perfluorooctanoic acid (PFOA) on  
30 bioconcentration and gene expression in common carp (*Cyprinus carpio*). Adult male and female carp  
31 were exposed to environmental (200 ng/L) and experimental (2 mg/L) doses of PFOA for 56 d. Carp  
32 exposed to 200 ng/L had levels of PFOA below the level of detection in all tissue samples analyzed,  
33 whereas variable concentrations were measurable in various tissues from carp exposed to 2 mg/L. The  
34 expression level of the glutathione S-transferase (GST) gene, coding for a detoxifying enzyme,  
35 increased in a PFOA dose- dependent manner in liver tissues from 200 ng/L to 2 mg/L exposure ( $p <$   
36  $0.05$ ). The expression levels of CYP19A gene, coding for the enzyme that converts testosterone into  
37 estrogen, were altered in gonadal tissues from male and female carp exposed to either 200 ng/L or 2  
38 mg/L; expression increased in male gonads and decreased in female gonads. Unexpectedly, the  
39 expression levels of CYP19A in male and female gonads from carp exposed to 200 ng/L or 2 mg/L  
40 were similar ( $p > 0.05$ ). Therefore, even though environmental doses of PFOA did not accumulate in  
41 tissues of the common carp, they did affect the gene expression levels of GST in the liver and  
42 CYP19A in the gonads. These observations raise concerns that exposure to environmental doses of  
43 PFOA may affect gene expression in animals and possibly in humans, with important health  
44 consequences.

45

## 46 **1. Introduction**

47 Perfluorooctanoic acid (PFOA) is a synthetic molecule of perfluorinated compounds [1]. It is widely  
48 used as a surfactant in industrial manufacturing of lubricants, medical equipment, paper and textile  
49 coatings, upholstery, shiny food packaging, and fire-fighting foams. Perfluorooctanoic acid is highly  
50 soluble in water and is nonbiodegradable. Various concentrations of PFOA have been detected in soil,  
51 air, and aquatic environments worldwide, indicating that PFOA is a persistent synthetic pollutant [2].  
52 Water is the major environmental reservoir of PFOA, as well as the main source of exposure for  
53 animals and humans [3]. Rivers and lakes with direct industrial emissions contain ng/L to mg/L  
54 concentrations of PFOA, whereas ocean levels are in the range of 10 to 100 pg/L [2].

55 Concentrations of PFOA, ranging from ng/L to mg/L, have been detected also in urban  
56 wastewater and in bottled water for human consumption [4].  
57 There is growing concern that environmental accumulation of PFOA is a risk factor for animal and  
58 human health. Biomonitoring studies have revealed the presence of PFOA in wildlife tissues  
59 [5,6] and in blood/serum of various human populations [7], suggesting that PFOA is a  
60 bioaccumulative compound. Bioaccumulation and toxic effects on the liver, kidneys, and gonads, as  
61 well as carcinogenic activity in the liver, have been revealed in fish and mammals exposed to PFOA  
62 [8–18]. Moreover, changes in expression of genes, such as the glutathione S-transferase (GST),  
63 which encodes a detoxifying enzyme, and CYP19A, which encodes the aromatase enzyme involved in  
64 steroidogenesis, also were reported in the liver and gonads [19,20]. However, most of these studies  
65 have been performed with experimental doses of PFOA that are higher than those detected in the  
66 natural environment of animals, and much higher than those to which humans are usually ex-  
67 posed [9,12,13,21]. These findings have raised the question of whether environmental concentrations  
68 of PFOA affect the health of living beings.  
69 Therefore, in the present study, we investigated the effects of environmental concentrations of PFOA  
70 on gene expression levels in target tissues of common carp, *Cyprinus carpio*. Adult male and female  
71 carp were exposed for 56 d to the environmental dose of 200 ng/L of PFOA, which is an intermediate  
72 concentration in the range of those measured at the upstream reaches of the Po River, between Turin  
73 and Milan, in northern Italy [22,23]. Bioaccumulation of PFOA in tissues and mRNA expression  
74 levels of GST and CYP19A genes were analyzed in 2 PFOA target tissues: the liver and gonads. The  
75 results were compared with those obtained from carp exposed to the experimental dose of 2 mg/L of  
76 PFOA.

77

## 78 **2. Materials and methods**

### 79 *2.1. Fish exposure test and tissue collection*

80 Thirty 2-yr-old carp were purchased from a local fish farm and acclimated according to Test No. 305  
81 of the Organisation for Economic Co-operation and Development (1996). The captive maintenance

82 procedures and research protocols were approved by the University of Ferrara Institutional Animal  
83 Care and Use Committee and by the Italian Ministry of Health (Law 116 of 27 January 1992) and the  
84 European Union (Directive 86/609/EEC of the European Union Parliament and of the Council of  
85 24 November 1986). Four weeks before starting the experiment, the carp were equally divided into 3  
86 different test tanks (10 fish per tank) and left to adapt to experimental conditions. The carp were fed  
87 with a commercial pellet food (Tetra Pond Pellets Mini) 3 times per week at 2% of their total body  
88 weight. Waste and uneaten food were removed regularly. The carp had an average total length of  
89  $19.3 \pm 2.5$  cm and weight of  $104.8 \pm 27.8$  g (mean  $\pm$  standard deviation [SD]). A flow-through  
90 exposure test was conducted for 56 d by means of a system that continuously delivered PFOA, diluted  
91 to the test concentrations, to the test tanks. The tanks were 120-L glass aquaria filled with tap water: a  
92 continuous supply of fresh water was provided at a flow-through rate of 500 mL/min. Exposures were  
93 carried out in 2 different tanks containing 200 ng/L of PFOA or 2 mg/L of PFOA; a third tank was  
94 used as negative control. These concentrations were selected on the basis of environmental reports  
95 [22,23] and experimental data from the literature [12,24,25]. The stock solutions were prepared by  
96 dissolving the PFOA (chemical purity 96%; Sigma-Aldrich) in distilled water and then delivering it  
97 into the 2 treatment tanks by a peristaltic pump at a flow rate of 0.42 mL/min. At time 0 of the  
98 exposure, an initial volume of the stock solution was added to the treatments tanks to immediately  
99 achieve the desired exposure concentration. The actual concentration of PFOA in the test water from  
100 each test tank was measured by liquid chromatography–mass spectrometry (LC–MS) 3 times during  
101 the exposure period (Table 1). The results of these analyses indicated that PFOA concentration was  
102 maintained at > 80% of each nominal concentration. The water temperature, pH, and oxygen  
103 saturation were recorded 3 times a week and kept at 10 to 15 8C, 6.70 to 8.00, and above 80%,  
104 respectively. At the end of the 56-d exposure, the fish were anesthetized with MS-222, killed by a  
105 deep cut through the neck, dissected, and sexed. The sex ratio was 1:1. Blood (0.5 mL), liver, and  
106 gonad (1 g from each carp) were flash-frozen and stored at  $-20$  8C until they were analyzed for  
107 PFOA.

108

109 2.2. Analysis of PFOA in tissues

110 Concentrations of PFOA in tissues were analyzed to determine the accumulation after 56 d of  
111 waterborne exposure. Perfluorooctanoic acid was extracted by use of an ion-pairing extraction  
112 procedure and measured by use of high-performance liquid chromatography (HPLC) with  
113 electrospray ionization tandem mass spectrometry, according to a widely used method [6,26–28].  
114 Approximately 1 g or 1 mL of sample was homogenized with 5 mL of Ultrapure water supplied by a  
115 Milli-Q system from Millipore. One milliliter of 0.5 M of tetrabutylammonium hydrogen sulphate  
116 (Sigma-Aldrich) solution and 2 mL of sodium carbonate buffer (Sigma-Aldrich; 0.25 M, pH 10) were  
117 added to 1 mL of the homogenate samples in a polypropylene tube and thoroughly mixed before  
118 extraction. Five milliliters of methyl tert-butyl ether (MTBE; Sigma-Aldrich) were added to the above  
119 mixture and shaken for 20 min. The organic and aqueous layers were separated by centrifugation, and  
120 an exact volume of MTBE (4 mL) was removed from the solution. The aqueous mixture was rinsed  
121 with MTBE and separated twice; both the rinses were combined in a second polypropylene tube.  
122 The solvent was evaporated under nitrogen and replaced with 0.5 mL of HPLC grade methanol  
123 (Sigma-Aldrich). This extract was passed through a nylon filter (0.2 mm; Supelco) into an HPLC  
124 vial. Extraction blanks were prepared using Milli-Q water. Analytes' separation was performed using  
125 a Finnigan Surveyor Plus HPLC System consisting of a quaternary pump, vacuum degasser, and  
126 autosampler. Chromatographic separation was achieved with a Betasil C18 column (50 x 2.1-mm i.d.,  
127 5  $\mu$ m) supplied by Thermo Electron. For quantitative determination, the HPLC system was interfaced  
128 to a Finnigan LTQ linear ion trap mass spectrometer (Thermo Electron) operating in negative  
129 electrospray mode. Primary and product ions monitored for PFOA determination were 498.8 >  
130 368.9. Ten microliters of each extract were injected in the LC-MS with 2mM LC-MS grade  
131 ammonium acetate (> 99%), as the mobile phase starting at 10% methanol (HPLC grade, Sigma-  
132 Aldrich). At a flow rate of 300 mL/min, the gradient increased to 95% methanol at 10 min before  
133 reverting to original conditions at 15 min. Column temperature was maintained at 25 °C. Standards  
134 for the 5-point calibration curve were prepared by progressive dilution with methanol from a neat  
135 standard purchased from Dr. Ehrenstorfer (Augsburg, Germany), and concentrations were evaluated

136 in comparison with this unex- tracted standard curve and were not corrected for the recoveries or for  
137 the purity of the standards (more than 98%). Individual stock solutions of the target analytes were  
138 prepared in methanol and stored in polypropylene bottles at –20 8C. Limit of detection (LOD),  
139 determined as 3 times the signal-to-noise ratio, was 0.4 ng/g wet weight. Chemicals and reagents  
140 were analytical grade, and Teflon-coated labware was avoided during the entire process of sampling,  
141 pretreatment, and analysis to minimize contamination of the samples. Data quality assurance and  
142 quality control protocols included matrix spikes, laboratory blanks, and continuing calibration  
143 verification. Blanks were analyzed with each set of 5 tissue samples as a check for possible laboratory  
144 contamination and interferences; recoveries, as- sessed using spiked matrix with a concentration of 5  
145 ng/g of the analyte, were more than 93% in blood, and more than 89% in solid tissues (gonad and  
146 liver).

147

### 148 *2.3. RNA extraction*

149 Total RNA was isolated from 20 to 30 mg of liver and gonadal tissues by use of an RNeasy Mini Kit  
150 (Qiagen) and treated with RNeasy-free DNase I (Qiagen) according to the manufacturer’s instructions  
151 [29]. Purity of RNA was determined by measuring the A260/A280 absorbance ratio using a  
152 NanoDrop 2000c (Thermo Fisher Scientific). Quality of RNA was checked by agarose gel (1.2%)  
153 electrophoresis.

154

### 155 *2.4. cDNA preparation*

156 Five hundred nanograms of RNA from each sample were reverse transcribed with a cDNA synthesis  
157 kit by use of random primers (ImProm-II Reverse Transcription System; Promega), adhering to the  
158 manufacturer’s instructions. The cDNA products were subsequently used as templates for real-time  
159 quantitative polymerase chain reaction (RT-qPCR) reactions.

160

### 161 *2.5. Real-time quantitative PCR*

162 Real-time quantitative PCR for CYP19A and GST expression analysis was carried out using SYBR  
163 Green JumpStart Taq ReadyMix (Sigma-Aldrich). The primers used for CYP19A expression [30]  
164 were 5'-GGTTTGCATCACTTCCACAA-3'- for 5'-AATACGGTCTGCCAGGTGTC-3'-rev [31].  
165 The size of the PCR product was 198 bp (position 546-762 of the *C. carpio* CYP19A mRNA. Gene  
166 Bank accession number DQ534411). The primers used for GST expression were 5'-  
167 CTATGTTAAGG- CATTGGGTCGCAAAC-3'- for 5'-ATCCACATAGCTCTTGA-  
168 GAGTTGGGAAGG-3'-rev [32]. The size of the PCR product was 338 bp (position 203-540 of the  
169 *C. carpio* Pi-class GST mRNA. Gene Bank accession number DQ411313). The glyceraldehyde 3-  
170 phosphate dehydrogenase gene was used as an internal control for gene expression analysis. The  
171 primers used were 5'-AGGGGCTCAGTATGTTGTGG-3'- for, and 5'-  
172 AGGAGGCATTGCTGACAACT-3'-rev [31]. The size of the PCR product was 185 bp (position  
173 278-422. Gene Bank accession number AJ870982.1). Cycling parameters for all reactions were 96  
174 8C for 10 min, 45 repeats at 95 8C for 30 s, and 62 8C for 1 min. The specificities of the PCR  
175 products were evaluated by melting curve profiles (temperature range from 60–95°C), with  
176 temperature increasing at a rate of 0.5 8C s<sup>-1</sup> and by 2% agarose gel electrophoresis. Data were  
177 analyzed by use of the 2-DDCt method [33]. Glutathione S-transferase mRNA and CYP19A  
178 mRNA were measured individually in each liver and gonadal tissue, respectively. Then fold-change  
179 values were averaged in each group. All RT-qPCR assays were performed 3 times (each with 3  
180 repeated measures) and results were averaged. To detect possible contamination, negative control  
181 reactions without cDNA templates were included.

182

### 183 2.6. Statistics

184 The statistical package Statistica Ver 7.1 (StatSoft) was used to analyze all data, with significance set  
185 at the 0.05 level. The mean, standard error, and range were determined for PFOA in tissues of each  
186 carp group. To allow inclusion of all samples in the statistical analyses, measured concentrations  
187 below the detection limit (0.4 ng g<sup>-1</sup> wet wt) were assigned a value of one-half the LOD. Shapiro–  
188 Wilk and Levene's tests were used to check the normality and the homogeneity of data variance. One-

189 way analysis of variance and the Student's t test were used to compare PFOA tissue concentrations  
190 among the treatment groups and between sexes.

191

### 192 **3. Results**

193 No mortality was observed during the acclimation or test in either control or tested fish. No carp  
194 showed signs of clinical disease and abnormality such as erratic swimming, incoordination of  
195 movements, respiratory distress, or hyporeflexia.

196

#### 197 *3.1. PFOA concentrations in tissue samples*

198 In all tissue samples from unexposed carp and from carp exposed to the environmental dose of 200  
199 ng/L of PFOA, the PFOA concentrations were below the LOD and significantly lower than those in  
200 carp exposed to the the experimental dose of 2 mg/L ( $p < 0.05$ ). Mean PFOA concentrations from  
201 carp exposed to the experimental dose of 2 mg/L ranged from 4 to 65 ng g<sup>-1</sup> wet weight (Table 2).  
202 Levels of PFOA were highest in blood, and then in liver and gonads. The mean PFOA concentration  
203 in blood was significantly higher than that in other tissues ( $p < 0.05$ ; Table 2). There were no  
204 significant differences in mean tissue PFOA concentrations between males and females ( $p > 0.05$ ;  
205 Table 2).

206

#### 207 *3.2. Expression level of GST mRNA in liver tissue*

208 Real-time quantitative PCR analysis of all liver tissues from carp (males and females together)  
209 indicated that mRNA expression of GST was significantly higher in the environmental dose of  
210 200 ng/L PFOA- and the experimental dose of 2 mg/L PFOA-treated groups than in unexposed carp  
211 (Figure 1). Specifically, the GST mRNA levels were 2.3-fold higher in livers from carp exposed to the  
212 environmental dose of 200 ng/L of PFOA and 5.4-fold higher in livers from carp exposed to the  
213 experimental dose of 2 mg/L of PFOA than in unexposed carp (Figure 1). There were no differences  
214 in liver GST expression level between males and females (data not shown).

215



216 *3.3. Expression level of CYP19A mRNA in gonads*

217 Expression levels of CYP19A mRNA in testes (n = 5) and ovaries (n = 5) from carp exposed to the  
218 environmental dose of 200 ng/L or the experimental dose of 2 mg/L of PFOA were analyzed with  
219 RT-qPCR and compared with corresponding unexposed carp (n = 5 males and n = 5 females). The  
220 expres- sion levels of CYP19A mRNA were 2.2-fold higher in testes from carp exposed to the  
221 environmental dose of 200 ng/L of PFOA and 3.4-fold higher in testes exposed to the experimen- tal  
222 dose of 2 mg/L of PFOA than in testes from unexposed carp (Figure 2). In ovaries, CYP19A  
223 expression was significantly downregulated by 3.7-fold in female carp exposed to the environmental  
224 dose of 200 ng/L of PFOA and by 3.4-fold in carp exposed to the experimental dose of 2 mg/L of  
225 PFOA compared with expression in ovaries from unexposed carp (Figure 3). Differences in CYP19A  
226 mRNA expression levels were no different between carp exposed to the environmental dose of 200  
227 ng/L and the experimental dose of 2 mg/L of PFOA ( $p > 0.05$ ).

228

229 **4. Discussion**

230 The present study reports—for the first time that we know of—the effects of environmental doses of  
231 PFOA in adult male and female carp. Perfluorooctanoic acid has been found in the aqueous  
232 environment worldwide at several ng/L concentrations as reported by Suja et al. [1] with reference to  
233 surface and tap water samples collected from different countries. The levels of PFOA measured in the  
234 control tank (Table 1) were similar to those reported in many surface waters [1], whereas the  
235 environmental dose of 200 ng/L used in the present study represents a high level of contamination  
236 reached in impacted rivers such as the Po River in Italy [22,23] or Cape Fear River (Wilmington, NC,  
237 USA) [34]. Thus PFOA effects and impact on fish biomarkers were herein evaluated under a field-  
238 relevant exposure scenario.

239 We investigated the accumulation of PFOA in carp tissues and changes in expression levels of the  
240 GST gene in the liver and the CYP19A gene in gonads, which are the main target tissues of PFOA.  
241 Our results showed that carp exposed for 56 d to the environmental dose of 200 ng/L of PFOA had  
242 levels of PFOA below the LOD in all tissue samples analyzed. Conversely, and in agreement with

243 previously reported data [17], concentrations of PFOA were detected in various tissues from carp  
244 exposed to the experimental dose of 2 mg/L of PFOA, with the highest levels in blood.

245 Perfluorooctanoic acid has low bioconcentration factors (BCFs) in fish (Organisation for Economic  
246 Co-operation and Development [35]); a BCF between 3.1 and 9.4 was reported for carp [36,37]. Thus  
247 it is not surprising that PFOA was not measurable in carp exposed to the lowest concentration  
248 (200 ng/L) and with the LOD of the method used in the present study.

249 Importantly, even though PFOA at environmental concentrations did not accumulate in the liver,  
250 liver tissues from carp exposed to the environmental dose of 200 ng/L of PFOA expressed GST  
251 mRNA levels significantly higher than those from unexposed carp. Moreover, GST mRNA levels in  
252 liver tissues from carp exposed to the experimental dose of 2 mg/L of PFOA were significantly higher  
253 than those from unexposed carp as well as those from carp exposed to the environmental dose of  
254 200 ng/L PFOA. Although only 2 PFOA concentrations were used in our experiments, this result  
255 suggests that the expression level of GST may be regulated in a PFOA dose-dependent manner.

256 Because GST expression is induced by increased levels of oxidant molecules [38], our data are  
257 consistent with PFOA acting as an inducer of reactive oxygen species in carp liver, as previously  
258 suggested in other animal models [39]. However, other studies of fish and mammals reported reduced  
259 expression of GST after administration/exposure to PFOA, indicating that PFOA can act differently in  
260 liver of some species.

261 In the present study, male gonads from carp exposed to the environmental dose of 200 ng/L of  
262 PFOA showed CYP19A mRNA levels significantly higher than those from unexposed male carp. In  
263 contrast, the expression levels of CYP19A were significantly lower in gonads from female carp  
264 exposed to the environmental dose of 200 ng/L of PFOA than those from unexposed female carp.

265 It is well known that aromatase is the enzyme that converts testosterone into estrogen and plays a key  
266 role in the establishment and maintenance of sex in fishes. Almost absent in male gonads, CYP19A is  
267 highly expressed in female gonads. In addition, certain environmental contaminants, including  
268 PFOA, act as endocrine disruptors by modifying the expression levels of the CYP19A gene. Studies  
269 in different model organisms support the concept that PFOA mimics the activity of

270 estrogens, leading to the disruption of the hypothalamic-pituitary-gonadal (HPG) endo- crine axis  
271 [30,40]. Tilton et al. [13] reported that dietary exposure to PFOA significantly increased expression  
272 of the sensitive estrogen biomarker protein vitellogenin (Vtg) in rainbow trout, and induced an overall  
273 hepatic gene expression profile highly similar to that of 17 $\beta$ -estradiol. Liu et al. [11] stated that PFOA  
274 increased Vtg expression in cultured tilapia hepatocytes, and Wei et al. [12] showed that PFOA  
275 increased Vtg expression and caused disruption of gonad development in rare minnows (*Gobiocypris*  
276 *rarus*). Although Vtg expression was not investigated in the present study, our results confirm that  
277 PFOA acts as an endocrine disruptor in the gonads of carp by increasing the expression level of the  
278 CYP19A gene in testes and reducing it in ovaries. Most importantly, CYP19A expression levels in  
279 both male and female gonads did not differ significantly between carp exposed to the environmental  
280 dose of 200 ng/L PFOA and those exposed to the experimen- tal dose of 2 mg/L. These data suggest  
281 that the environmental dose of 200 ng/L of PFOA is nearly as effective as the experimental dose  
282 of 2 mg/L in causing endocrine disruption in the male and female gonads of carp. We may speculate  
283 that PFOA functions as physiologically active hormones that have biological effects at low doses. In  
284 this context, a few molecules of PFOA may be sufficient to saturate target receptors and elicit the  
285 downstream response.

286 Overall, the present study shows that PFOA at the environmental dose of 200 ng/L does not  
287 accumulate in carp tissues; nevertheless, it produces the same changes in gene expression as those at  
288 the experimental dose of 2 mg/L. Notably, whereas environmental doses of PFOA do not seem to  
289 affect the detoxifying activity of the liver (i.e., GST is efficiently expressed), they act as an endocrine  
290 disruptor in male and female gonads by dysregulating the expression levels of CYP19A. Given that  
291 many recent studies in animal models, as well as in human populations, have highlighted the adverse  
292 effects of PFOA in relation to its bioaccumulation, our data emphasize that environmental PFOA  
293 levels dysregulate the HPG axis, in the absence of measurable bioaccumulation. Therefore, in the case  
294 of the carp and fish in general that live in aquatic environments containing PFOA, the risk exists of  
295 permanently altering the HPG axis and leading ultimately to effects on the reproductive system.

296 Whether altered CYP19A expression in male and female carp exposed to the environmental dose of  
297 200 ng/L of PFOA can also affect fish reproductive potential (fecundity, hatching, growth, or  
298 development/sexual maturation) was not evaluated in the present study. It is known, however, that  
299 treatment with experimental doses of PFOA alters the sexual organs in male and female rare  
300 minnows, in the way that oocytes develop in testis and ovaries undergo degeneration [12]. Moreover,  
301 PFOA parental exposure of Japanese medaka resulted in adverse effects to offspring, leading to  
302 increased mortality in the F1 fish generation [41]. On the other hand, Hagenaaers et al. reported no  
303 effects on zebrafish reproduction after exposure at 0.1-, 0.5-, and 1-mg/L doses for 1 mo [39].  
304 Although fish reproductive endocrinology has some unique aspects, the basic structure and function  
305 of the HPG axis is well conserved across all vertebrates. Hence the results obtained with endocrine  
306 disruptor compounds in fish models might be extrapolatable to other species. Furthermore, because  
307 humans may be exposed to PFOA (although through pathways different from in fish), the  
308 causal relationships presented in the present study might be analogous to those described in humans  
309 [42]. Notably, in recent years, increases in endocrine disorders such as precocious puberty,  
310 menopause, and infertility have been suspected of being caused by exposure to endocrine disruptors  
311 [43–46]. Moreover, it is well known that endocrine disruptors that mimic natural hormones exert their  
312 biological effects at similarly low doses. Therefore, PFOA's estrogenic activity combined with its  
313 capability to induce biological effects in the range of environmental exposures should be seriously  
314 considered as a possible factor in the development of human diseases.

315 In conclusion, the present study reports that environmental concentrations of PFOA do not  
316 accumulate in tissues after 56 d of exposure but PFOA does act as an endocrine disruptor in male and  
317 female carp. Such evidence raises concerns that exposure to environmental doses of PFOA may affect  
318 gene expression in other animals and humans. Further studies are required to identify additional  
319 exposure pathways and determine if similar changes in gene expression occur in other animals and  
320 humans.

321

322

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473

474 Table 1. Perfluorooctanoic acid nominal and measured test concentrations in the test tanks<sup>a</sup>.

Nominal test concentration	Measured test concentration <sup>b</sup>
2 mg/L	1.932 ± 0.023 mg/L
200 ng/L	182.53 ± 12.16 ng/L
0 ng/L (unexposed)	3.49 ± 0.55 ng/L

475

476 <sup>a</sup>Measures were taken at the beginning (day 1), middle (day 28), and end (day 56) of the exposure  
477 period.

478 <sup>b</sup>Values are mean ± standard deviation (SD) for n = 3.

479

480 Table 2. Perfluorooctanoic acid (PFOA) concentrations<sup>a</sup> in samples of carp exposed to 2 mg l<sup>-1</sup> for 56  
 481 d<sup>b</sup>.

Tissue/Organ	Range values (ng g <sup>-1</sup> )	Mean ± SE (ng g <sup>-1</sup> ) (all fish n = 10)	Mean ± SE (ng g <sup>-1</sup> ) (females n = 5)	Mean ± SE (ng g <sup>-1</sup> ) (males n = 5)
blood	21.8 - 97.8	64.9 ± 7.3A	66.3 ± 9.3A	63.2 ± 12.7A
liver	21.4 - 38.9	28.1 ± 1.6A	27.6 ± 2.6A	28.8 ± 1.9A
gonad	0.9 - 11.5	3.7 ± 0.9A	3.6 ± 1.6A	3.3 ± 5A

482 <sup>a</sup>ng g<sup>-1</sup> wet weight

483 <sup>b</sup>All the values were above the limit of detection (0.4 ng g<sup>-1</sup> wet wt). A indicates significant  
 484 differences in PFOA concentrations among diverse tissues/organs. No disparities between the sexes in  
 485 any tissue/organ were found using the Student's t test.

486 \*p < 0.05.

487 SE = standard error

488

489 **Figure legends**

490

491 **Figure 1.** Expression levels of glutathione S-transferase mRNA measured with real-time  
492 quantitative polymerase chain reaction in liver tissues from carp. The white bar represents  
493 unexposed carp; the gray bar indicates carp exposed to the environmental dose of 200 ng/L of  
494 perfluorooctanoic acid (PFOA); and the black bar denotes carp exposed to the experimental dose of 2  
495 mg/L of PFOA. Relative expression was calculated by use of the comparative threshold (DDCt)  
496 method. Data are expressed as a relative-fold change ( $2^{-DDCt}$ ) over the value of unexposed carp. Error  
497 bars represent standard error of mean for  $n = 10$ . The glyceraldehyde 3-phosphate dehydrogenase  
498 gene was used as internal control. All reactions were performed in triplicate and results were  
499 averaged.  $p < 0.0001$  versus carp exposed to the environmental dose of  
500 200 ng/L of PFOA and versus unexposed carp;  $^Dp < 0.05$  versus unexposed carp.

501

502 **Figure 2.** Expression levels of CYP19A mRNA measured with real-time quantitative polymerase  
503 chain reaction in testes tissues from male carp. The white bar depicts unexposed carp; the grayish bar  
504 identifies carp exposed to the environmental dose of 200 ng/L of perfluorooctanoic acid (PFOA); and  
505 the black bar represents carp exposed to the experimental dose of 2 mg/L of PFOA. Error bars are  
506  $\pm$  standard error for  $n = 5$ .  $p < 0.0001$  versus unexposed carp;  $^Dp < 0.05$  versus unexposed  
507 carp.

508

509

510 **Figure 3.** Expression levels of CYP19A mRNA measured with real-time quantitative polymerase  
511 chain reaction in ovarian tissues from female carp. The white bar shows unexposed carp; the grayish  
512 bar represents carp exposed to the environmental dose of 200 ng/L of  
513 perfluorooctanoic acid (PFOA); and the black bar denotes carp exposed to the experimental  
514 dose of 2 mg/L of PFOA. Error bars represent  $\pm$  standard error for  $n = 5$ .  $p < 0.0001$  versus  
515 unexposed carp;  $^Dp < 0.0001$  versus unexposed carp.