

1           **Analysis of neurotransmitters in *Daphnia magna* affected by neuroactive**  
2           **pharmaceuticals using liquid chromatography-high resolution mass**  
3           **spectrometry**

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23  
24          **Abstract**

25          Neurotransmission plays an essential role during the central nervous system (CNS)  
26          development. A few number of studies about the neurotransmitters changes by the effect of  
27          pharmaceuticals in aquatic organisms have been reported in the last years. There are evidences  
28          that the aquatic ecotoxicological model organism *Daphnia magna* shares with vertebrates several  
29          of the neurotransmitters targeted by antidepressant and other neuro-active drugs. So, a method  
30          based on liquid chromatography coupled to high-resolution mass spectrometry (LC-HRMS) has  
31          been applied for the first time to *Daphnia magna* and validated to study the levels of 41  
32          neurotransmitters under the effect of four different neuro-active pharmaceuticals (sertraline,  
33          venlafaxine, duloxetine and fluoxetine). In addition, the performance of LC-HRMS was studied in  
34          terms of linearity, sensitivity, intra- and inter-day precision, and overall robustness. The developed  
35          analytical method using LC-HRMS is a new tool for neurotoxicology research using the *Daphnia*  
36          *magna* model. The results show a general increasing or decreasing in the concentrations of the  
37          neurotransmitters after the exposure of the pharmaceuticals studied respect to controls.

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39

## 40 **Capsule**

41 This is the first study that analyse the changes in the neurotransmitter profiles using  
42 liquid chromatography coupled to high resolution mass spectrometry.

43 **Keywords:** *Daphnia magna*; LC-Orbitrap; pharmaceuticals; neurotransmitters.

44

## 45 **INTRODUCTION**

46 Neurotransmitters are endogenous chemicals that enable neurotransmission. It is a  
47 type of chemical messenger which transmits signals across a chemical synapse, such  
48 as a neuromuscular junction, from one neuron (nerve cell) to another "target" neuron,  
49 muscle cell, or gland cell (Lodish et al., 2000). In this line, neurotransmitters plays an  
50 essential role during the central nervous system (CNS) development (Horzmann and  
51 Freeman, 2016; Kurian et al., 2011). Thus, there is a growing interest to investigate the  
52 effect of neuro-active contaminants on neurotransmitter dysregulation and how such  
53 effect alter the structure and function of the CNS in different biological models (Faria et  
54 al., 2018; Gómez-Canela et al., 2018; Ren et al., 2015). Of particular interest are  
55 antidepressant drugs belonging to selective serotonin and serotonin-norepinephrine  
56 reuptake inhibitors (SSRI, SNRI), which act increasing the levels of serotonin and  
57 norepinephrine in the CNS, and are broadly used in the first world (HAINER et al., 2006).  
58 Consequently chemical residues of SSRIs such as fluoxetine and sertraline and of SNRIs  
59 such as venlafaxine and duloxetine are conspicuous of most surface waters (Coelho et  
60 al., 2019; Reis-Santos et al., 2018).

61 The crustacean *Daphnia magna* (*D. magna*) may be a suitable non-vertebrate model  
62 to study the toxicological consequences of contaminant mediated disruptive effects on  
63 neurotransmitters. *D. magna* is one of the organisms most used in Aquatic toxicology  
64 (Barata et al., 1998; Miner et al., 2012). *D. magna* is a small planktonic invertebrate  
65 crustacean that can be found in lakes and shallow ponds rich in organic matter sediment  
66 distributed throughout the northern hemisphere. Therefore, it is an important herbivore  
67 and at the same time the prey for a huge number of predators (fishes, amphibians, and  
68 invertebrates). Moreover, their genome has been recently coded, being quite similar to  
69 the vertebrate genome (Colbourne et al., 2005). *D. magna* has a high number of  
70 neurotransmitters in common with many vertebrates (Dircksen et al., 2011; McCool et

71 al., 2011) that are affected by the action of neuroactive pharmaceuticals (Rivetti et al.,  
72 2016). Indeed immunohistological studies have reported that the SSRI fluoxetine  
73 increases the activity of serotonin in the *D. magna* brain (Campos et al., 2016). In this  
74 study we selected four pharmaceuticals that share similar modes of action and are widely  
75 used worldwide to treat depressive disorders (Abreu et al., 2016; Fong and Ford, 2014;  
76 Schultz et al., 2010).

77 The analysis of neurotransmitters requires a sample pretreatment, extraction  
78 method and an accurate and reliable instrumental method. Most of the analytical  
79 methods recently published for the analysis of neurotransmitters in aquatic organisms  
80 have been using liquid chromatography coupled to triple quadrupole (QqQ) mass  
81 spectrometers (LC-MS/MS) (Gómez-Canela et al., 2018; Santos-Fandila et al., 2013;  
82 Tufi et al., 2015). Triple quadrupole mass spectrometry is particularly fitted for the  
83 analysis of target compounds because its extraordinary sensitivity and selectivity  
84 (Gómez-Canela et al., 2013). The limitation of this technique is the possibility to have  
85 false positive due to the low resolution of the mass spectrometer. Therefore, the arrival  
86 of the high-resolution mass spectrometry (HRMS) got over the limitations commented  
87 before. Orbitrap, which is an ion trap mass analyzer, can operate in full-scan mode in a  
88 single run and the medium high resolution is around 100,000 FWHM (full width at half  
89 maximum) at  $m/z$  200 and high mass accuracy < 2 ppm using internal standard and  
90 within 5 ppm with external calibration (Hogenboom et al., 2009). Moreover, using HRMS  
91 a theoretically unlimited number of analytes can be extracted from the full scan data  
92 without any compromise regarding the resulting sensitivity. Furthermore, HRMS permits  
93 the elucidation of elemental composition of analytes based on exact masses and isotopic  
94 patterns (Kaufmann et al., 2010). Thus, the aim of this study has been to develop a  
95 multiresidue method for the analysis of 43 neurotransmitters and related metabolites in  
96 *D. magna* using LC-HRMS after the exposure to neuroactive pharmaceuticals.

97

## 98 **EXPERIMENTAL**

### 99 ***Chemicals and materials***

100 All neurotransmitter standards, including the isotope labelled compounds used for the  
101 optimization of the analytical method were purchased from Sigma-Aldrich (St. Louis, MO,  
102 USA) and Santa Cruz Biotechnology (Dallas, TX, USA). All target compounds and  
103 molecular formulas are shown in Table 1. Methanol (MeOH), water (H<sub>2</sub>O), and acetone  
104 HPLC grade were supplied from Merck (Darmstadt, Germany). Pharmaceuticals used

105 for the expositions (sertraline, venlafaxine, duloxetine and fluoxetine) were also  
106 purchased from Sigma-Aldrich (St. Louis, MO, USA). Acetonitrile (ACN) and formic acid  
107 (FA) were supplied by Fischer Scientific (Loughborough, UK). Ultra-pure water was  
108 obtained through Millipore Milli-Q purification system (Millipore, Bedford, MA, USA).  
109 Initially, all pure standards were prepared at 1000 ng  $\mu\text{L}^{-1}$  in MeOH or milliQ water  
110 depending on their solubility. Once prepared in silanized amber vials (2 mL), the  
111 standards were kept and at  $-80\text{ }^{\circ}\text{C}$  in the dark to prevent possible degradation.

112

### 113 ***Aquatic organism husbandry***

114 *D. magna* was kept in the laboratory at  $20\text{ }^{\circ}\text{C}$ , in ASTM medium water (reconstituted  
115 hard water formulation), adding  $4\text{ mL L}^{-1}$  of marine microalgae extract (as a medium  
116 conditioner). *D. magna* fed on the microalga *Chlorella vulgaris* at concentrations of  
117  $400,000\text{ cells mL}^{-1}$ . The hatchery for the production of neonates (juveniles under 24 h),  
118 were kept in tanks of 1.5 to 2.5 L, with a density of 5 females  $\text{L}^{-1}$ . The culture medium  
119 and the food were completely renewed 3 times per week.

120 Under adequate conditions, *D. magna* reproduces asexually by parthenogenesis,  
121 having several litters throughout its life cycle. Once the sexual maturation is reached, the  
122 first litter of neonates is released (around the age of 9-10 days). From then, they continue  
123 to produce offspring, which are released from the incubator chamber of the female every  
124 3 days (approx.). The organisms used for toxicity tests were those produced from the  
125 third litter. Breeding tanks of females were kept for a maximum period of 30 days  
126 (although their longevity is longer), so they have staggered cultures, with a gap of 1 to 2  
127 weeks, to replace those that once they reach 30 days old were discarded.

128

### 129 ***Exposures and sample collection***

130 Chronic, toxicologically relevant concentrations of sertraline ( $0.1$  and  $1\text{ }\mu\text{g L}^{-1}$ ),  
131 venlafaxine ( $0.1$  and  $1\text{ }\mu\text{g L}^{-1}$ ), duloxetine ( $0.1$  and  $1\text{ }\mu\text{g L}^{-1}$ ) and fluoxetine ( $1\text{ }\mu\text{g L}^{-1}$ ) were  
132 studied in *D. magna*. At those concentrations, the studied species reveal reproductive  
133 changes (Campos et al., 2016; Rivetti et al., 2016). These drugs have been detected in  
134 the natural aquatic environment at low concentrations ( $\text{ng L}^{-1}$ ;  $\mu\text{g L}^{-1}$ ) (Schreiber and  
135 Szewzyk, 2008). Exposures were initiated with new born neonates (<24 h old) in groups  
136 of five individuals in 100 mL of ASTM medium plus food ( $5 \times 10^5$  cells of *Chlorella vulgaris*)  
137 and lasted four days. At the end of exposures animals were collected, pooled in a 2 mL

138 Eppendorf, frozen with N<sub>2</sub> and preserved at -80°C until processed for analysis. In addition  
139 the body length of least 10 animals from each treatment was measured. Body length  
140 measurements were performed from the top of the head to the base of the spine using  
141 a Nikon stereoscope microscope (SMZ 150, Nikon, Barcelona, Spain) and the ImageJ  
142 software (<http://rsb.info.nih.gov/ij/>) (Glazier, 1998).

### 143 **Sample extraction**

144 Pools of 5 exposed or control crustaceans (*D. magna*) were spiked with 500 ng of  
145 isotope labelled solution (Glycine-2-<sup>13</sup>C, L-Aspartic acid-<sup>15</sup>N, Serotonin-d<sub>4</sub>, Dopamine-  
146 1,1,2,2-d<sub>4</sub>, GABA-d<sub>6</sub>, Acetylcholine-d<sub>9</sub>, L-Glutamic-<sup>15</sup>N acid. Choline-1-<sup>13</sup>C, L-  
147 Phenylalanine-1-<sup>13</sup>C and Taurine-<sup>15</sup>N) as internal standard (internal standard mixture,  
148 ISM, in the next sections). Then, 500 µL of ACN (+0.1% formic acid + 0.02% ascorbic  
149 acid) were added and shaken using a vortex mixer. Three stainless steel beads (3 mm  
150 diameter) were placed in each sample and were homogenized using a bead mill  
151 homogenizer (TissueLyser LT, Qiagen) at 50 oscillations per min during 60 s. After that,  
152 samples were centrifuged 10 min at maximum speed at 4°C. The supernatant was  
153 transferred to a new centrifuge tube and evaporated to complete dryness under mild  
154 nitrogen. Then, the samples were resuspended in 100 µL of ACN:H<sub>2</sub>O (50:50). Finally,  
155 the supernatant was filtered using 0.20 µM PTFE filters (DISMIC -13 JP, Advantec®) and  
156 kept in amber chromatographic vials at -80 °C until LC-HRMS analysis. More details  
157 about the extraction protocol are included in a previous published paper about the  
158 characterization of neurotransmitter profiles in *D. magna* juveniles exposed to  
159 environmental concentrations of antidepressants, anxiolytic and antihypertensive drugs  
160 (Rivetti et al., 2019).

161

### 162 **LC-HRMS analysis**

163 An LC-Orbitrap-MS (Exactive) from Thermo Fischer Scientific (Bremen, Germany)  
164 equipped with a heated electrospray ionization (H-ESI) source was used. The system  
165 was equipped with a HTC PAL autosampler and a Surveyor MS Plus pump. Synergi  
166 Polar-RP 80 Å column (250 mm× 4.6 mm i.d., particle size 4 µM, Phenomenex, Torrance,  
167 USA) was chosen following a previous published paper about the analysis of  
168 neurotransmitters in zebrafish larvae using an LC-MS/MS instrument (Gómez-Canela et  
169 al., 2018). The mobile phase composition consisted of binary mixtures with 0.1% of  
170 formic acid in water (A) and 0.1% formic acid in MeOH (B). Gradient elution started at  
171 95% A and 5% B in the first 2 min and increased to 30% B in 5 min. Then, gradient

172 increased to 95% B in the next 13 min, and held for 5 min. Initial conditions to stabilize  
173 the system were attained in 5 min being 30 min the total run time. The flow rate was set  
174 at 600  $\mu\text{L min}^{-1}$  and 10  $\mu\text{L}$  was injected. Neurotransmitters were measured under both  
175 positive and negative electrospray ionization (ESI+/ESI-). In the case of the metabolites  
176 detected by positive and negative ionization, the mode with better MS signal intensity for  
177 each compounds was used. Full scan acquisition over a  $m/z$  50-600 was performed at  
178 70,000 FWHM and spray voltage at 3.00 kV, capillary voltage at 25 V, skimmer voltage  
179 at 16 V and tube lens voltage at 55 V were used. Sheath gas flow rate at 35 arbitrary  
180 units (au), auxiliary gas flow rate at 10 au and capillary temperature at 300  $^{\circ}\text{C}$  were  
181 chosen. High energy Collision Dissociation (HCD) voltage at 25 V was optimized to  
182 obtain the most intense fragments. All Orbitrap conditions are displayed in Table SI1  
183 (*Supplementary information*). The data were acquired and processed using the Thermo  
184 Xcalibur 2.1 software package. Solvent blanks did not contain any of the investigated  
185 metabolites, indicating no carryover effect in any of the LC-HRMS runs.

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### 189 **Quality assurance**

190 Table 2 displays the quality parameters obtained by LC-HRMS. The linearity of  
191 response was evaluated over a concentration range from 0.005 to 5  $\text{ng } \mu\text{L}^{-1}$ , using eight  
192 calibration points. Internal standard calibration was used to correct the analyte MS  
193 response and to ensure exact quantification performance. A mix of ten isotope labelled  
194 compounds, including L-aspartic acid- $^{15}\text{N}$ , glycine-2- $^{13}\text{C}$ , Serotonin- $\text{d}_4$ -hydrochloride,  
195 dopamine-1,1,2,2- $\text{d}_4$  hydrochloride, GABA- $\text{d}_6$ , acetylcholine- $\text{d}_9$ , L-glutamic- $^{15}\text{N}$  acid,  
196 choline chloride-1- $^{13}\text{C}$ , L-phenylalanine-1- $^{13}\text{C}$  and taurine- $^{15}\text{N}$ , were used as internal  
197 standard (IS) for extraction and analytical quality control. With LC-HRMS, good  
198 determination coefficients ( $R^2 > 0.99$ ) were obtained for all compounds with good linearity  
199 from 0.005 to 5  $\text{ng } \mu\text{L}^{-1}$  in most cases. Instrumental Detection Limit (IDL) was calculated  
200 as the concentration that yielded an S/N ratio higher than or equal to three using the  
201 lowest concentration standard solution at 2.5  $\text{ng/mL}$  and method detection limit (MDL)  
202 was calculated in the same way, using *D. magna* samples spiked at 200 ng. Intra-day  
203 precision was assessed by five consecutive injections of 1  $\text{ng } \mu\text{L}^{-1}$  standard solution, and  
204 inter-day precision was determined by measuring the same standard solution during four

205 different days. Solvent blanks did not contain any of the investigated analytes, indicating  
206 no carryover effect during the LC-HRMS runs.

207 On the other hand, recovery studies were performed with five replicates, using *D.*  
208 *magna* samples spiked at 200 ng with the neurotransmitter standard mixture and 500 ng  
209 of the ISM. Five replicates of a pool with five *D. magna* were analyzed first, subtracting  
210 the possible traces of target compounds. In addition, the matrix effect (ME) was  
211 assessed by comparing the peak area of analyte from the spiked zebrafish larvae with  
212 the peak area of the analyte from the standard solution at the same concentration in  
213 methanol (N = 5), following the equation:

$$214 \quad ME = \frac{(A - B)}{C} \times 100$$

215 where A is the peak area of each analyte from spiked *D. magna* samples; B is the peak  
216 area of each analyte from non-spiked *D. magna*; and C is the peak area of each analyte  
217 in methanol. ME values below 100% suggest signal suppression and ME values above  
218 100% suggest signal enhancement.

219

## 220 **RESULTS AND DISCUSSION**

### 221 ***Optimization of the LC-HRMS conditions***

222 As commented before, chromatographic conditions used were those reported in  
223 Gómez-Canela et al. study (Gómez-Canela et al., 2018). Optimization of the ionization  
224 source conditions was performed using LC-Orbitrap. Flow Injection Analysis (FIA)  
225 experiments were performed at a capillary voltage from 20 to 80 V, skimmer voltage from  
226 10 to 40 V, and tube lens voltage from 60 to 200 V. The best sensitivity (highest  
227 response) was obtained at a capillary voltage of 25 V, skimmer voltage of 16 V and tube  
228 lens voltage of 55 V that produced the most intense protonated molecule at high  
229 resolution (70,000 FWHM). Sheath gas flow rate was optimized from 35 to 55 au,  
230 auxiliary gas flow rate (from 2 to 20 au) and capillary temperature (>225 °C). These  
231 conditions were optimized taking into account the best results for all target compounds.  
232 HCD voltages were tested at 15, 25 and 45 eV, obtaining the most intense fragments at  
233 25 eV. Tables S11 displays the final Orbitrap ionization parameters. On the other hand,  
234 Synergy Polar-RP 80 Å column was used to obtain the best separation of all target  
235 compounds in a total run analysis of 30 minutes. Figure 1 shows the LC-HRMS extracted  
236 ion chromatogram of 43 target compounds from a full scan LC-Orbitrap-MS spectrum

237 using a 5 ppm extraction window. Compounds with the same exact mass could be  
238 identified by their different retention time ( $R_t$ ), were injected individually at the same  
239 conditions. Thus, octopamine and dopamine ( $[M+H]^+ = 154.0863$ ) eluted at 5.53 and 7.33  
240 min, respectively. Epinephrine and normetanephrine ( $[M+H]^+ = 184.0968$ ) at 5.59 and  
241 6.27 min; and betaine and valine ( $[M+H]^+ = 118.0863$ ) at 5.21 and 5.47 min, respectively.  
242 Therefore, one or two fragments were identified for each compound using HCD cell at  
243 25 eV, as indicated in Table 3. Directive 2002/657/CE, indicates the performance criteria  
244 and other requirements for mass spectrometric detection (Official Journal of the  
245 European Communities, 2002). To comply with this Directive, the identification criteria  
246 required that the accurate mass measurements of the molecular and the product ions  
247 should present an error  $<5$  ppm and when possible; the isotopic pattern should fit better  
248 than 90% to be accepted as a positive sample. To comply with the requirements of the  
249 Directive 2002/657/CE, a molecular and two fragment ions for  $R > 20,000$  earn 2 and 2.5  
250 IPs, respectively, which reinforces the robustness of the identification criteria ( $IPs \geq 4$ )  
251 for HRMS. In some cases, more than two HCD ions were formed, which can be used for  
252 identification criteria and provide  $2n$  identification points, being “n” the number of ions at  
253 high resolution.

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255

## 256 **Method validation**

257 Table 2 shows the performance parameters obtained by LC-Orbitrap-MS. In order to  
258 correct the MS responses and ensure exact quantification performance, internal  
259 standard calibration was used. Acceptable correlation coefficients ( $R^2$ ) were over 0.99  
260 for all target compounds in a range from 0.005 to 5 ng  $\mu\text{L}^{-1}$  in most of the cases. Linearity  
261 and response factors are displayed in Table 2. IDL values ranged between 0.2 pg  
262 (betaine) and 156.74 pg (CDP-Choline). On the other hand, MDL values of *D. magna*  
263 were from 0.03 pg daphnia<sup>-1</sup> (choline, valine and BW284c51) to 4.75 pg daphnia<sup>-1</sup> (L-  
264 DOPAC) (Table 2). Regarding intra- and inter-day precision, values differed significantly  
265 depending on the compound. Intra-day precision ( $n=5$ ) ranged from 0.29 to 26.26% and  
266 inter-day precision ( $n=5$ ) ranged from 1.3 to 40.9% at 1 ng  $\mu\text{L}^{-1}$  level, indicating a robust  
267 method response (Table 2).

268 Method performance was tested using five replicates of a pool with 5 daphnids at 100  
269 ng (low concentration) and 200 ng (high concentration) with the mix of the target  
270 neurotransmitters and 500 ng of the ISM in order to assess the efficiency of the extraction

271 method. Good recovery values (between 75±12%-124±8%) were obtained for all target  
272 compounds except for acetylcholine and guanidine, 55±9% and 67±3% respectively, and  
273 cytidine-5'-diphosphocholine that was not recovered. Matrix effect was also evaluated  
274 obtaining signal suppression for 8 neurotransmitters (acetylcholine, histidine, 5-hydroxy-  
275 L-tryptophan, octopamine, norepinephrine, taurine, L-arginine and guanosine). However,  
276 only phosphocholine, guanidine and phenylalanine had ME above 100% giving a signal  
277 enhancement. The rest of them had good matrix effect values in the range of 60±8% to  
278 111±13% (Table 2). Thus, quantification with the internal standard procedure seems  
279 adequate to avoid over- or underestimation of the calculated concentration.

280

### 281 ***Changes in the neurochemical profile of Daphnia magna***

282 The body length of the *Daphnia* individuals at the end of exposures was 2.22±0.39 mm  
283 (Mean±SD, N =83) and did not vary significantly across treatments (ANOVA results,  $F_{1,75}$   
284 =1.85,  $P>0.05$ ). Raw data (mean ± SE) is depicted in Table SI3 whereas significant and  
285 other important results relative to unexposed controls are shown in Figures 2-6. Results  
286 obtained indicated significant ( $P<0.05$ ) differences in 24 out of the 34 metabolites  
287 quantified across treatments (Table SI4). There is quite a lot of inconsistency between  
288 drugs acting similarly (Figures 2-6). For example, sertraline and fluoxetine are selective  
289 serotonin reuptake inhibitors that bind to the serotonin transporter and hence inhibit the  
290 re-uptake of serotonin, increasing its accumulation in the synaptic spaces. In humans,  
291 SSRI by increasing serotonin levels in the brain diminished depression disorders,  
292 whereas in *D. magna* it increases reproduction and carbohydrate metabolism (Campos  
293 et al., 2016; Jordão et al., 2016). Unfortunately, our analytical method did not allow  
294 detection of serotonin but instead two serotonergic metabolites: its precursor  
295 tryptophan and its major degradation product 5-HIAA. Sertraline and fluoxetine had  
296 opposed patterns for 5-HIAA increasing and decreasing it, respectively (Figure 2).  
297 Interestingly in rats similar opposed effects of sertraline and fluoxetine in the  
298 concentration of 5-HIAA have been reported (Harkin et al., 1995; Kudrin et al., 2010),  
299 which may be related to the fact that both compounds despite of having a similar primary  
300 mode show different bioaccumulation patterns and probably different secondary mode  
301 of action. In fish, sertraline bioaccumulated in neural and other tissues to a greater extent  
302 than fluoxetine (Chen et al., 2017; David et al., 2018), and there is reported evidence  
303 indication that sertraline is more toxic than fluoxetine in fish and other aquatic organisms  
304 (Alsop and Wilson, 2019; Sehonova et al., 2018).

305 Venlafaxine and duloxetine are antidepressants of the selective serotonin-  
306 norepinephrine reuptake inhibitor (SNRI) class. This means they increase the  
307 concentrations of the neurotransmitters serotonin and norepinephrine in the body and  
308 the brain. Unfortunately, norepinephrine could not be detected in *Daphnia* by the method  
309 used. Nevertheless, our results showed quite similar concentration patterns for 5-HIAA  
310 and epinephrine among the studied SNRI (Figure 2). None of them changed significantly  
311 the levels of 5-HIAA, although both compounds reduced it and only venlafaxine  
312 increased epinephrine probably since the concentrations used (100 and 1000 ng/L) were  
313 too low for duloxetine to affect its target neurotransmitter (Figure 2). These results are in  
314 line with reported studies in mammalian models (Chappell et al., 2014; Fuller et al., 1994;  
315 Vasudev et al., 2016).

316 The observed effects on the remaining neurotransmitter pathways are quite variable,  
317 which is a consequence that the studied drugs only affected them as secondary and  
318 unknown mechanisms of action in *Daphnia*. As expected, none of the studied  
319 compounds affected ACh, which is the target of organophosphorus and neonicotinoid  
320 pesticides in *Daphnia* (Figure 3). However, the concentrations of ACh precursors such  
321 as choline, betaine, CHOP and GPCh varied across the studied compounds (Figure 3).  
322 Choline and betaine decreased upon exposure to most compounds but fluoxetine, and  
323 CHOP and GPCh increased at low concentrations of venlafaxine and GPCh,  
324 respectively. Guanidine increased upon exposure to fluoxetine. It is difficult to provide a  
325 conclusive argument to the disparity of responses observed for the cholinergic related  
326 metabolites since many of them are also precursors of lipids and the studied drugs in  
327 general increased reproduction (Campos et al., 2012b). *D. magna* females provision  
328 large quantities of lipids to their eggs thus it is possible that by doing so lipid precursors  
329 such as choline, CHOP and GPCh diminished (Jordão et al., 2015).

330 GABA is the principal neurotransmitter with inhibitory effects in the central nervous  
331 system. GABA was enhanced upon exposure to low levels of sertraline but not at high  
332 levels neither by any other tested chemicals (Figure 4). Although these results are quite  
333 inconsistent there is reported information indication that SSRI can enhance GABA  
334 (Bhagwagar et al., 2004).

335 Tyrosine and phenylalanine were enhanced by fluoxetine, and dopamine  
336 concentrations decreased upon exposure to sertraline (Figure 4). Tyrosine and  
337 phenylalanine are amino acid involved in many metabolic pathways, thus it is difficult to  
338 establish a mechanistic functional relationship for fluoxetine. Reported studies in  
339 *Daphnia*, however, found that fluoxetine increased catabolic activity (Campos et al.,  
340 2012a), which may favour the accumulation of these amino acid as by-products of  
341 degradation of proteins. Effect of sertraline on dopamine contradict reported studies in

342 rats that found the opposite pattern (Kitaichi et al., 2010). Glutamatergic and glycinergic  
343 metabolites were consistently decreased upon exposure to the tested chemicals (Figure  
344 5). There is ample information that there is a cross-talk among serotonergic,  
345 glutamatergic and glycinergic signaling pathways and that SSRI and SNRIs may interact  
346 with those pathways (Aprison et al., 1996; Dale et al., 2015; Musazzi et al., 2013; Ortiz  
347 et al., 2016; Pehrson and Sanchez, 2014; Rodrigues et al., 2015). Despite that *Daphnia*  
348 has an active histaminergic system (McCoole et al., 2011), any of the metabolites  
349 involved in this system were affected by the studied compounds (Table SI3). Among  
350 other metabolites related with the neurotransmitter system, L-arginine and aspartic acid  
351 decreased upon exposure to all the studied compounds, whereas fluoxetine enhanced  
352 the level of taurine and methionine. Again, it is difficult to establish a mechanistic  
353 relationship between observed changes in these metabolites and neurotoxicological  
354 effects since these amino acids are also involved in other metabolic functions (Figure 6).  
355 However, there is information reporting that fluoxetine can enhanced the levels of taurine  
356 and that low levels of L-arginine and aspartic acid in human plasma are related with  
357 depressive disorders (Colmenares-Aguilar et al., 2017; Lu et al., 2014).

358

## 359 **CONCLUSIONS**

360 A comprehensive optimization of a multiresidue method for the analysis of 43  
361 neurotransmitters and related metabolites from different biochemical classes in *D.*  
362 *magna* using LC-Orbitrap-MS has been performed. The method has been validated in  
363 terms of reproducibility, sensitivity and mass accuracy. Moreover, a study of the  
364 fragmentation pattern was done in order to confirm the target compounds in the samples.  
365 Forty-one neurotransmitters were recovered with good values (between 75±12%-  
366 124±8%) with the exception of acetylcholine and guanidine, 55±9% and 67±3%  
367 respectively, and cytidine-5'-diphosphocholine that was not recovered. Moreover, matrix  
368 effect has been evaluated for all the neurotransmitters studied, obtaining high values of  
369 signal suppression for some compounds such as acetylcholine, histidine, 5-hydroxy-L-  
370 tryptophan, octopamine, norepinephrine, taurine, citrulline and guanosine. This  
371 analytical technique is adequate for the multi-group analysis of neurotransmitters and  
372 offers advantages with respect to previous reported methods: 43 compounds can be  
373 analyzed in a single injection; the compounds are unequivocally identified using high-  
374 resolution mass spectrometry and subsequently the excellent sensitivity of LC-Orbitrap-  
375 MS is reported.

376 Results demonstrated that the pharmaceuticals exposed to *D. magna* affected to the  
377 neurotransmitter system increasing or decreasing their concentrations respect to  
378 controls. In *D. magna* most tested pharmaceuticals altered neurotransmitters per se or  
379 their associated metabolites at environmental concentrations (0.1 ng L<sup>-1</sup>). Thus, it can be  
380 concluded that pharmaceuticals are an important cause of concern in the aquatic  
381 environment. Further studies are needed in order to know more information about these  
382 organic pollutants in terms of human risk.

383

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