

1 Biomonitoring of 21 endocrine disrupting chemicals in human hair samples using  
2 ultra-high performance liquid chromatography–tandem mass spectrometry

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12 **ABSTRACT** Rapid industrial growth has increased human exposure to a large variety of  
13 chemicals with adverse health effects. These industrial chemicals are usually present in the  
14 environment, foods, beverages, clothes and personal care products. Among these compounds,  
15 endocrine disrupting chemicals have raised concern over the last years. In the present work, the  
16 determination of 21 EDCs in human hair samples is proposed. An analytical method based on the  
17 digestion of the samples with a mixture of acetic acid/methanol (20:80, v/v) followed by a solid-  
18 liquid microextraction and analysis by ultra-high performance liquid chromatography–tandem  
19 mass spectrometry (UHPLC–MS/MS) was developed and validated. The most influential  
20 parameters affecting the extraction method were optimized. The method was validated using  
21 matrix-matched calibration and recovery assays. Limits of detection ranged from 0.2 to 4 ng g<sup>-1</sup>,  
22 limits of quantification from 0.5 to 12 ng g<sup>-1</sup>, and inter- and intra-day variability was under 15%  
23 in all cases. Recovery rates for spiked samples ranged from 92.1 to 113.8%. The method was  
24 applied for the determination of the selected compounds in human hair. Samples were collected  
25 weekly from six randomly selected volunteers (three men and three women) over a three-month  
26 period. All the analyzed samples tested positive for at least one of the analyzed compounds.

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28 **KEYWORDS:** Endocrine disrupting chemicals; Biomonitoring; Human hair analysis; UHPLC–  
29 MS/MS

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## 32 1. Introduction

33

34 Humans are constantly exposed to new chemicals with adverse health effects. These  
35 potentially hazardous compounds, are ubiquitous in the environment, foods, beverages, and  
36 personal care products. Among these emerging contaminants, endocrine disrupting chemicals  
37 (EDCs) have raised concern over the last years. EDCs are synthetic and natural substances that  
38 can interfere with the hormonal systems in wildlife and humans. The biological effects of EDCs  
39 are related to their ability to mimic/antagonize endogenous hormones, or alter the synthesis and  
40 metabolism of endogenous hormones and receptors (Sonneschein and Soto, 1998). Bisphenol A  
41 (BPA) and its chlorinated derivatives, parabens (PBs), benzophenone-UV filters (BPs) and  
42 perfluoroalkyl compounds (PFCs) are included in this group of compounds (Fei et al., 2009;  
43 Kavlock et al., 2013; Liao et al., 2009; Liu et al., 2007; Paris et al., 2002; Rivas et al., 1997; U.S.  
44 Environmental Protection Agency, 2003).

45 BPA is mainly used for the manufacturing of epoxy resin and polycarbonate plastics, with an  
46 annual production of over 2-3 million tons. BPA has attracted considerable attention of  
47 governments and the scientific and medical community (Bisphenol A website). The European  
48 Food Safety Authority (EFSA) and the US-Environmental Protection Agency (US-EPA) have  
49 established the maximum acceptable level of exposure to BPA of 50  $\mu\text{g kg}^{-1}$  body weight/day  
50 (Nicolucci et al., 2013). There are many sources of human exposure to BPA, with diet considered  
51 the main source (Calafat et al., 2008). Interestingly, literature has reported the formation of  
52 chlorinated derivatives of BPA during the chlorination process of drinking water before  
53 consumption (Yamamoto and Yasuhara, 2002). PBs are the alkyl esters of *p*-hydroxybenzoic acid.  
54 They are used as antimicrobial preservatives, mainly against mold and yeast. PBs are used as  
55 additives in the manufacture of cosmetic products, pharmaceuticals, food and beverages. In the  
56 last years, several studies have demonstrated the ability of PBs to disrupt physiological functions

57 in both in vitro (van Meeuwen et al., 2008), and in vivo models (Boberg et al., 2010; Soni et al.,  
58 2005). Inhalation, dermal contact and ingestion are the main sources of exposure to PBs (El  
59 Houssein et al., 2010). The existing European Union (EU) limit for total PB concentration in  
60 cosmetics is 0.8 % (w/w) and 0.4 % (w/w) as acid for mixtures or individually, respectively  
61 (European Union Regulation No. 1223/2009). BPs are one of the most common UV filters used in  
62 sunscreens because they absorb UVA (320 to 400 nm) and UVB (290 to 320 nm) radiation. BPs  
63 have been reported to enter the human body through the food chain (Cuderman and Heath, 2007)  
64 and skin absorption (Jiang et al., 1999). Finally, PFCs are commonly used in the manufacture of  
65 finished products such as paints, adhesives, waxes, polishes, electronics, fire-fighting foams and  
66 caulks, as well as grease-proof coatings for food packaging (Corsini et al., 2014; Giesy and  
67 Kannan, 2001; Henkster et al., 2003; Rivière et al., 2014). Although food is the major exposure  
68 route for PFCs, drinking water, indoor air and house dust can also be potential sources of PFC  
69 exposure (Domingo, 2011).

70 Hair analysis has been traditionally used for drug testing, especially in cases of drug abuse  
71 (Olivezira et al., 2007). In addition, hair has been used as a bio-indicator of exposure to pesticides  
72 in children and exposure to organochlorine pollutants in adults. Although human milk, placental  
73 tissue, serum, blood or urine are conventionally used to determine exposure to EDCs, hair analysis  
74 could be used as a complementary tool, especially in cases of chronic exposure or when traditional  
75 matrices are not available. The main advantage of hair analysis is its large window of detection,  
76 which allows us to establish a chronological profile of exposure based on hair length (Cooper et  
77 al., 2012). Additional advantages of hair analysis include non-invasive collection and easy  
78 monitoring, difficult tampering with samples, storage and transportation at room temperature, and  
79 most contaminants are chemically stable (Caplan and Golberger, 2001). Therefore, the analysis  
80 of hair samples can might be of great interest for the assessment of exposure to different EDCs  
81 (Król et al., 2013), but not many analytical methods have been validated for the determination of

82 the EDCs mentioned in hair samples. Moreover, the methods described in the literature focused  
83 on the individual determination of BPA (Tzatzarakis et al., 2015), PBs (Sakol et al., 2015), or both  
84 chemicals (Martín et al., 2015).

85 The aim of the present work was to develop a multi-class method for the determination of 21  
86 EDCs in human hair samples. A sensitive multi-residue method based on the digestion of hair  
87 samples followed by a solid-liquid microextraction prior to the UHPLC–MS/MS analysis was  
88 developed. BPA and its chlorinated derivatives (mono- bi- tri- and tetrachlorobisphenol A), four  
89 PBs (methyl-, ethyl-, n-propyl- and butylparaben), six BP-UV filters (benzophenone-1,  
90 benzophenone-2, benzophenone-3, benzophenone-6, benzophenone-8 and 4-  
91 hidroxybenzophenone) and six PFCs (perfluorobutanoic to perfluorooctanoic acid and  
92 perfluorooctane sulfonate) were selected as target analytes. After validation, the method was  
93 applied to determine the free pollutant content in samples from six randomly selected volunteers  
94 (three men and three women) from Granada, Spain.

95

96

## 97 **2. Experimental**

98

### 99 *2.1. Chemicals and reagents*

100

101 The reagents used for the experiments were of high analytical grade and purity. A Milli-Q  
102 system from Millipore (Bedford, MA, USA) was used for water purification (18.2 MΩ cm).  
103 Perfluorobutanoic acid (PFBuA) (98%), perfluoropentanoic acid (PFPeA) (97%),  
104 perfluorohexanoic acid (PFHxA) ( $\geq 97\%$ ), perfluoroheptanoic acid (PFHpA) (99%),  
105 perfluorooctanoic acid (PFOA) (96%) and perfluorooctanesulfonic acid (PFOS) ( $\geq 98\%$ ),  
106 perfluoro-n-[1,2,3,4- $^{13}\text{C}_4$ ]octanoic acid (MPFOA), methylparaben (MPB), ethylparaben (EPB), n-

107 propylparaben (PPB), and n-butylparaben (BPB) were supplied by Alfa Aesar (Massachusetts,  
108 MA, USA). Bisphenol A (BPA), tetrachlorobisphenol A (Cl<sub>4</sub>-BPA), deuterium labelled bisphenol  
109 A (BPA-d<sub>16</sub>), benzophenone-1 (BP-1), benzophenone-2 (BP-2), benzophenone-3 (BP-3),  
110 benzophenone-6 (BP-6), benzophenone-8 (BP-8), 4-hydroxybenzophenone (4-OH-BP),  
111 deuterium labelled benzophenone (BP-d<sub>10</sub>), deuterium labelled ethylparaben (EPB-d<sub>5</sub>), LC-MS  
112 grade methanol, water and acetonitrile, acetic acid and ammonia (25%, w/v) were supplied by  
113 Sigma-Aldrich (Madrid, Spain). Mono-, di- and trichlorobisphenol A (Cl-BPA, Cl<sub>2</sub>-BPA, Cl<sub>3</sub>-  
114 BPA) were synthesized in our laboratory (purity > 99 %) by chlorination of BPA (Vilchez et al.,  
115 2003). Stock solutions of 1000 mg L<sup>-1</sup> of each compound, prepared in methanol, were stored in a  
116 freezer at -20 °C. Working solutions were prepared fresh weekly by diluting the stock standard  
117 solutions in methanol or in the initial mobile phase. They were maintained at 4 °C in the  
118 refrigerator. All of the solutions were stored in the dark, in order to prevent photodegradation.

119

## 120 *2.2. Instrumentation and software*

121

122 A chromatographic system Acquity UPLC™ H-Class (Waters, Manchester, UK), provided  
123 with a binary solvent manager was used for chromatographic separation. Analyte detection was  
124 carried out using a triple quadrupole mass spectrometer (Waters) Xevo TQS with an orthogonal  
125 Z-spray™ electrospray ionization (ESI) source. The stationary phase was an Acquity UPLC® BEH  
126 C<sub>18</sub> column (50 mm × 2.1 mm i.d., 1.7 μm particle size). Other laboratory equipment such as a  
127 vortex-mixer (IKA, Staufen, Germany), an ultrasound-HD bath (Selecta, Barcelona, Spain), a  
128 Spectrafuge™ 24D centrifuge from Labnet International, Inc. (New Jersey, USA) and a sample  
129 concentrator (Stuart, Staffordshire, UK) were also used. Statistical analysis of data was performed  
130 with Statgraphics Plus version 5.1 (Statpoint Technologies Inc., Virginia, USA).

131

132 2.3. *Sample collection and storage*

133

134 Hair samples were collected from six healthy volunteers (three men and three women) aged  
135 18 or older from the city of Granada, Spain. Hair samples were collected weekly for a three-month  
136 period. All volunteers were informed about the scope and nature of the study. Hair specimens were  
137 cut from the posterior vertex region of the head, as this region is associated with the smallest  
138 variation in growth rate, and as close as possible to the scalp. Sample length was of roughly 3–5  
139 cm. Samples were anonymized and stored in aluminum foil, at room (ambient) temperature until  
140 further processing and analysis. One of the concerns in hair analysis is the need to differentiate  
141 between external contamination (from air or dust) and internally incorporated chemicals. The ideal  
142 decontamination procedure should remove external contamination (deposited on the hair cuticle)  
143 without affecting internally incorporated compounds (present in the hair bulk).

144 To remove endogenous substances as well as adsorbed chemicals from the surface of the hair  
145 (Schramm, 2008), samples were washed twice with MilliQ water (ultra-sonicated for 5 min),  
146 washed with SDS (for 5 min), and finally rinsed with MilliQ water (for 5 min). In forensic  
147 protocols, SDS has shown to be significantly less effective for removing external contamination  
148 of hair with lipophilic substances compared with more aggressive substances such as MeOH or  
149 DCM. For this work, however, water and SDS were used since, as reported by the scarce literature  
150 available, the use of organic solvents could extract internally deposited contaminants (Kucharska  
151 et al., 2015), especially polar organic pollutants like PBs or BPA. SDS and water are used by most  
152 authors to determine other organic pollutants like polybrominated diphenyl ethers (PBDEs) (Tadeo  
153 et al., 2009; Zheng et al., 2011; Krol et al., 2014), polychlorinated biphenyls (PCBs) (Zhao et al.,  
154 2008; Wielgomas et al., 2012; Liang et al., 2014) or pesticides (Altshul et al., 2004; Cuong et al.,  
155 2012) in hair samples. After washing, samples were cut in 2–3 mm pieces, dried at room

156 temperature and wrapped in aluminum foil until analysis. Thus, all pollutant concentrations  
157 determined are related to the weight of the dry hair.

158

## 159 *2.4. Basic procedure*

160

### 161 *2.4.1. Preparation of spiked samples*

162

163 No certified reference materials are commercially available for the determination of these  
164 compounds in hair. Therefore, spiked samples were prepared for optimization and validation of  
165 the method. The samples were spiked at eight concentration levels (from 0.5 to 500.0 ng g<sup>-1</sup>) for  
166 calibration standards, validation and quality control, and at 100.0 ng g<sup>-1</sup> for method optimization.  
167 Aliquots of 0.05 g were weighted into 2 mL Eppendorf tubes and spiked with 100 μL of a methanol  
168 solution (50.0 ng mL<sup>-1</sup>) of the four surrogates: MPFOA, BPA-d<sub>16</sub>, EPB-d<sub>5</sub> and BP-d<sub>10</sub> (100.0 ng g<sup>-1</sup>  
169 <sup>l</sup> final concentration in hair). After shaking for 1 min, the solvent was allowed to evaporate at room  
170 temperature and the spiked samples were ready for the optimization and/or validation experiments.

171

### 172 *2.4.2. Sample treatment*

173

174 An aliquot of the hair sample (0.05 g) containing the surrogates was weighed in a 2 mL  
175 Eppendorf tube, and incubated with 0.5 mL of a mixture acetic acid/methanol (20:80, v/v) at 38  
176 °C for 12 hours. After cooling to room temperature, 1 mL of acetonitrile was added for the  
177 extraction of analytes. The mixture was shaken for 15 min and centrifuged for 5 min at 16,300 ×  
178 g. The organic phase (acetonitrile) containing the analytes was separated into a glass vial and  
179 evaporated to dryness under a nitrogen stream at room temperature. The residue was dissolved

180 with 250  $\mu\text{L}$  of the initial mobile phase. After stirring for 30 s in a vortex and centrifugation for 5  
181 min at  $16,300 \times g$ , the sample was injected into the UHPLC system.

182

### 183 *2.4.3. Ultra-high performance liquid chromatography–tandem mass spectrometry conditions*

184

185 Chromatographic separation of compounds was performed using an Acquity UPLC<sup>®</sup> BEH C<sub>18</sub>  
186 column (50 mm  $\times$  2.1 mm i.d., 1.7  $\mu\text{m}$  particle size). A gradient mobile phase consisting of 0.1%  
187 (v/v) ammonium in water (solvent A) and methanol (solvent B) was used. The optimized gradient  
188 was: 0.0–2.0 min, 30% B; 2.0–5.0 min, 30–90% B; 5.0–5.1 min, 90–100% B; 5.1–7.0 min, 100%  
189 B and back to 30% in 0.1 min (total run 10 min). Flow rate was 0.25 mL  $\text{min}^{-1}$ , injection volume  
190 10  $\mu\text{L}$ , and column temperature 40  $^{\circ}\text{C}$ .

191 Selected reaction monitoring mode (SRM) was used for operation of the triple quadrupole  
192 mass spectrometer. A unit mass resolution was set at Q1 and Q3 quadrupoles. ESI was performed  
193 in negative ion mode. The ion source temperature was 150  $^{\circ}\text{C}$ . Capillary voltage was 0.60 kV;  
194 source temperature 150  $^{\circ}\text{C}$ ; desolvation temperature, 500  $^{\circ}\text{C}$ ; cone gas flow, 150 L  $\text{h}^{-1}$ ; desolvation  
195 gas flow, 500 L  $\text{h}^{-1}$ ; collision gas flow, 0.15 mL  $\text{min}^{-1}$ ; and nebulizer gas flow, 7.0 bars. Nitrogen  
196 (99.995%) was used as cone and desolvation gas. The collision gas was argon (99.999%). Dwell  
197 time was fixed in 25 ms. Table 1 shows the parameters and mass transitions for each compound.

198

**Table 1**

199

### 200 *2.4.4. Quality assurance and quality control*

201

202 To assure the validity of the results obtained, simple quality assurance and quality control  
203 (QA/QC) measurements were carried out. First, to test for background contamination, procedural  
204 blanks were injected. Blanks were processed in the same way as the samples and injected into the



205 UHPLC–MS/MS system. No quantifiable amounts of target compounds were detected. To  
206 evaluate potential contamination and variability in the chromatographic system, spiked samples at  
207 two concentration levels, 0 and 250 ng g<sup>-1</sup>; and a standard in the initial mobile phase (100 ng mL<sup>-1</sup>)  
208 were injected in triplicate every 20 samples.

209

210

### 211 **3. Results and discussion**

212

#### 213 *3.1. Extraction procedure optimization*

214

215 An extraction with organic solvents was used for the isolation of the analytes from the  
216 previously digested hair samples. Key variables of the extraction step such as type of solvent and  
217 technique used were accurately optimized using aliquots of 0.1 g of sample spiked with the  
218 compounds at a concentration level of 100 ng g<sup>-1</sup>. All experiments were carried out in triplicate.

219

##### 220 *3.1.1. Selection of the extraction solvent*

221

222 Four extraction solvents (acetone, acetonitrile, ethyl acetate and methanol), which are widely  
223 used in the literature for the extraction of different families of EDCs from biological samples, were  
224 assayed (Rodríguez-Gómez et al., 2015). After incubation, spiked hair samples were mixed with  
225 1 mL of each of the studied solvents. The highest extraction efficiencies (> 80% in all cases) were  
226 obtained using acetonitrile, followed by acetone, ethyl acetate and methanol. Therefore,  
227 acetonitrile was selected for optimization.

228

##### 229 *3.1.2. Selection of the extraction procedure*

230  
231 One of the main difficulties traditionally found in hair analysis is to distinguish between  
232 external contaminants and chemicals incorporated from the human body. In order to overcome  
233 this, hair samples were thoroughly washed to remove contaminants from the surface and biogenic  
234 compounds, such as fatty acids, sphingolipids or steroids, which might affect analysis. Sample  
235 washing protocol has been described in Section 2.3. After washing, the key variables of the  
236 extraction procedure to be optimized are sample amount, solvent for extraction volume, and  
237 incubation time. A 15-run Box–Behnken experimental design including three replicates at the  
238 center point was used for fitting a second-order response surface. Three factors and three levels  
239 for each one were selected: amount of sample (0.05, 0.125 and 0.2 g), acetonitrile volume (0.5, 1.0  
240 and 1.5 mL) and incubation time (0.6 and 12 h). The data were analyzed using ANOVA, which  
241 provided determination coefficients ( $R^2$ ) > 0.90 in all cases. P-values for the *lack-of-fit* test were  
242 >0.05, which makes the model satisfactory with a 95% confidence level. Pareto charts were also  
243 obtained and statistically significant effects of the variables were screened using a Student's t-test.  
244 Variables having a confidence > 95% were considered to have a significant effect on the extraction  
245 efficiency. The desirability function was studied for the combination of the optimized  
246 experimental values for each compound. First, the responses for each analyte in the Box–Behnken  
247 design were normalized (0 to 1). The global desirability function is the geometric mean for each  
248 response. Figure 1 shows the plot of this function when sample amount *vs.* solvent volume, with  
249 fixed extraction time, is represented. The optimal values were 12 h for incubation time, 0.05 g for  
250 sample amount and 1.0 mL for solvent volume.

251

252

### Figure 1

253

254 *3.1.3. Optimization of the re-dissolution step*

255  
256 After isolation of analytes from digested samples, the extraction solvent is removed by  
257 evaporation under a nitrogen stream, and the dry extract is then re-dissolved in the initial UHPLC  
258 mobile phase for chromatographic analysis. The effect of the volume of reconstitution solvent is  
259 an important aspect to take into account when mass spectrometry is used as the detection  
260 technique. Selectivity aspects, including ion suppression, are critical in this case. Volumes of 50,  
261 100, 250, 500 and 1000  $\mu\text{L}$  were tested, and it was observed that the use of volumes  $> 250 \mu\text{L}$   
262 provided a cleaner extraction solution with less background, while with 50 and 100  $\mu\text{L}$  the  
263 extraction solution appeared cloudy and the noise was extremely high, making quantification  
264 difficult. Since a constant response was observed for volumes  $> 250 \mu\text{L}$  in all cases, this was the  
265 selected volume.

266

### 267 3.2. Analytical performance

268

269 A calibration curve at eight concentration levels was built for each compound.  
270 Analyte/surrogate peak area ratio *versus* concentration of analyte was used for establishing the  
271 calibration function. Surrogates (MPFOA, BPA-d<sub>16</sub>, EPB-d<sub>5</sub> and BP-d<sub>10</sub>) were added at a final  
272 concentration of 100 ng g<sup>-1</sup> of hair. The presence/absence of matrix effect was estimated using two  
273 calibration curves for each EDC. The first curve was built in the initial mobile phase, and the  
274 second in samples of natural dark hair free of analytes obtained from one of the volunteers. A  
275 Student's *t*-test was used for comparison of the two calibration curves. Previously, a Snedecor's  
276 *F*-test was applied to compare the estimated variances ( $S^2_{y/x}$ ). The Student's *t*-test showed, in most  
277 of the cases, statistically significant differences between the slopes of the two calibration curves,  
278 which indicated significant matrix effects. Therefore, the validation and the application of the

279 method was carried out using matrix-matched calibration for all the studied analytes. The  
280 analytical and statistical parameters are summarized in Table 2.

281

282 **Table 2**

283

### 284 3.3. Validation of the method

285

286 Linearity, sensitivity (limits of detection, LODs, and quantification, LOQs), accuracy (in  
287 terms of trueness and precision) and selectivity were determined. The US Food and Drugs  
288 Administration (FDA) guideline for Bioanalytical Method Validation was used.

289 *Linearity.* The concentration range selected was from the minimal quantified amount (LOQ)  
290 to 125.0 ng g<sup>-1</sup> for all compounds except for BPA, its chlorinated derivatives and PFBuA which  
291 was 250.0 ng g<sup>-1</sup> (see Table 2). Linearity of the calibration curves was assessed with the  
292 determination coefficients (% R<sup>2</sup>) and the P-values (% P<sub>lof</sub>) of the *lack-of-fit* test (Analytical  
293 Method Committee, 1994). The values obtained for R<sup>2</sup> ranged from 99.4% for PFOA to 99.9% for  
294 Cl-BPA, Cl<sub>2</sub>-BPA, Cl<sub>4</sub>-BPA, MPB, EPB, BPB, BP-1, BP-2, BP-3, BP-6, BP-8, 4-OH-BP, PFBuA,  
295 PFPeA, PFHpA and PFOS. P<sub>lof</sub> values > 5% were found for all compounds. Therefore, a good  
296 linearity within the stated ranges can be assured.

297 *Sensitivity.* LODs and LOQs were calculated in order to check the sensitivity of the analytical  
298 method. These parameters were calculated from the calibration function, using the standard  
299 deviation of residual, S<sub>y/x</sub>, the slope of the curve, b, and an estimation of the standard deviation of  
300 the blank (S<sub>0</sub>), (Currie, 1999). The LOD is defined as 3·S<sub>0</sub> and the LOQ as 10·S<sub>0</sub>. The values  
301 obtained for the LODs ranged from 0.2 to 4 ng g<sup>-1</sup>, and the LOQs from 0.5 to 12 ng g<sup>-1</sup> for MPB  
302 and PFBuA respectively. The results are summarized in Table 2.

303 *Accuracy (precision and trueness).* Since no certified materials are commercially available to  
304 evaluate the precision and the trueness of the proposed method, a recovery assay with spiked hair  
305 samples (free of the analytes), at three concentrations levels for each compound, was proposed.  
306 For BPA and its chlorinated derivatives and PFBuA, 25.0, 125.0 and 250.0 ng g<sup>-1</sup> were the  
307 concentrations assayed; and 5.0, 25.0 and 125.0 ng g<sup>-1</sup> for to the rest of the EDCs analyzed. The  
308 analyses were carried out in triplicate over six days. The relative standard deviation (RSD) was  
309 used for precision determination, and % recovery was applied for trueness evaluation. Precision  
310 and trueness results are shown in Table 3. The recoveries obtained were close to 100% and RSDs  
311 were < 15% for all compounds. The values obtained for both parameters are within the acceptable  
312 limits defined by the Analytical Method Committee guidelines ( $\leq 15\%$ , except for the LOQ which  
313 should not exceed 20%) which indicate that the method is highly accurate.

314

315

**Table 3**

316

317 *Selectivity.* The chromatograms of the procedure blank and the corresponding blank of sample  
318 were analyzed. Taking into account the retention time of all analytes, no interferences from  
319 endogenous substances were detected. The spectrometric conditions used ensured high selectivity  
320 of the UHPLC-MS/MS method. Figure 2A shows the SRM chromatograms obtained from a blank  
321 hair sample spiked with the analytes (75.0 ng g<sup>-1</sup>).

322

**Figure 2**

323

#### 324 *3.4. Biomonitoring of EDCs in human hair. Method application*

325

326 After validation, the method was used for the determination of the selected EDCs in hair  
327 samples from six volunteers. In order to study the persistence of compounds over a long period of

328 time, a hair sample from each volunteer was collected weekly over a three-month period (12  
329 samples per volunteer). Three replicates were analyzed for each day. The results obtained,  
330 expressed as means of concentrations for each volunteer, are summarized in Table 4. In all the  
331 analyzed samples at least one of the compounds studied was found. Figure 2B shows the  
332 chromatograms of one of the samples (M04).

333

334

#### Table 4

335

336 As shown in Table 4, 13 of the 21 analyzed compounds were detected in almost one of the  
337 samples but were not quantified in all of them. BPA was detected in only three of the samples.  
338 Chlorinated derivatives of BPA were not detected in any of the analyzed samples. For PBs, at least  
339 one of the compounds was detected and quantified in five of the samples. Regarding BPs, only  
340 four of the six compounds were detected and/or quantified in at least one sample. BP-1 and BP-3  
341 showed the highest concentration values. Finally, among the six PFCs analyzed, PFOA was the  
342 most commonly found compound, followed by PFOS. Both compounds were quantified in 83%  
343 of the samples. PFPHxA and PFPHpA were also detected in five of the samples but quantified in  
344 one, with concentrations close to the LOQs for both compounds.

345 Because of the limited information available on the selected pollutants in human hair, the  
346 comparison of the results previously published in hair and other biological matrices is of great  
347 interest. Perez et al. (2012) studied the presence of PFCs in human hair and urine and found a  
348 similar pattern of PFCs to ours, but they found much higher concentration in urine than in hair  
349 samples. PFOS and PFOA were the compounds more frequently quantified. PFOA was present  
350 in the 0.1–6 ng g<sup>-1</sup> range among the positive samples, whereas PFOS was between 3.7 and 7.0 ng  
351 g<sup>-1</sup>, while concentrations up to 49 ng L<sup>-1</sup> were reported in urine samples. In a large study by  
352 Vassiliadou et al. (2010), concentrations ranging from 7.5 to 15 ng mL<sup>-1</sup> for PFOS and from 2 to

353 4 ng mL<sup>-1</sup> for PFOA were found in blood samples from three different groups of adults living in  
354 Greece. Some of the selected analytes have also been previously analyzed by the authors of the  
355 present work in human milk (Rodríguez-Gómez et al., 2014). BPA, PBs and BPs were detected  
356 and quantified in almost one of the analyzed samples. However, none of the chlorinated derivatives  
357 of BPA were detected. BPA was quantified at concentrations ranging from 3.2 to 10.8 ng mL<sup>-1</sup>.  
358 Among the four PBs analyzed, EPB and PPB were the most abundant quantified in most milk  
359 samples, with concentrations ranging from 0.8 to 43.5 ng mL<sup>-1</sup> for PPB and from 0.8 to 15.0 ng  
360 mL<sup>-1</sup> for EPB. Martín et al. (2015) reported concentration levels of up to 1980 ng g<sup>-1</sup> for EPB and  
361 PPB in human hair from German individuals. Hines et al. (2015) reported concentrations in  
362 multiple matrices (milk, urine and serum) of PBs and BPA. These data suggest that urine is the  
363 best matrix for measurements, with concentration levels similar to those found in this study, 993  
364 and 279 ng mL<sup>-1</sup> for MPB and PPB respectively. In a recent study to estimate the burden of BPA  
365 in urban and rural populations, Tzatzarakis et al. (2015) reported concentration levels in hair  
366 between 17.7–192.8 ng g<sup>-1</sup> in urban population and between 13.1–72.8 ng g<sup>-1</sup> in rural population.

367

368

#### 369 **4. Conclusions**

370

371 A new analytical method for the biomonitoring of 21 EDCs (BPA and its chlorinated  
372 derivatives, 4 PBs, 6 BP-UV filters and 6 PFCs) in human hair samples has been successfully  
373 validated with good analytical features using a simple extraction procedure with acetonitrile after  
374 incubation of samples, and UHPLC-MS/MS analysis. The method was successfully validated,  
375 obtaining low LODs (between 0.2 ng g<sup>-1</sup> for MPB; and 4 ng g<sup>-1</sup> for PFBuA), high recoveries (>90%  
376 and <115%) and precision (RSD <15%). The proposed procedure of sample preparation was  
377 considered optimal because of its high extraction yield and easy operation.

378 The method has been applied to hair samples obtained from six volunteers over a three-month  
379 period. To our knowledge, this is the first validated method that allows the biomonitoring of such  
380 a large number of EDCs. At least one of the compounds analyzed was found in 100% of the  
381 samples, although quantification was not always possible.

382 The main advantage of the present study is the possibility of performing a multiresidue and  
383 multiclass analytical determination in only two steps while providing the sensitivity and selectivity  
384 necessary for the detection of compounds in human hair samples. Since hair is a representative  
385 sample for providing evidences of EDC exposure, the sampling protocol is crucial. Hair should be  
386 cut from the posterior vertex region of the head, as close as possible to the scalp (root) to minimize  
387 external exposure, with a length of roughly 3–5 cm. The main limitation of hair analysis is  
388 distinguishing between internal and external exposure. For this reason, hair samples were carefully  
389 washed. After washing, samples were cut in 2-3 mm pieces. Hair natural pigmentation and hair  
390 treatments are also possible limitations of hair analysis.

391 Finally, one of the limitations of the present work is the small amount of samples, and a larger  
392 cohort is required for the determination of global exposure to the target analytes. Nonetheless, this  
393 method is a powerful analytical tool that can be used in further determinations of human exposure  
394 to EDCs.

395

396

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398

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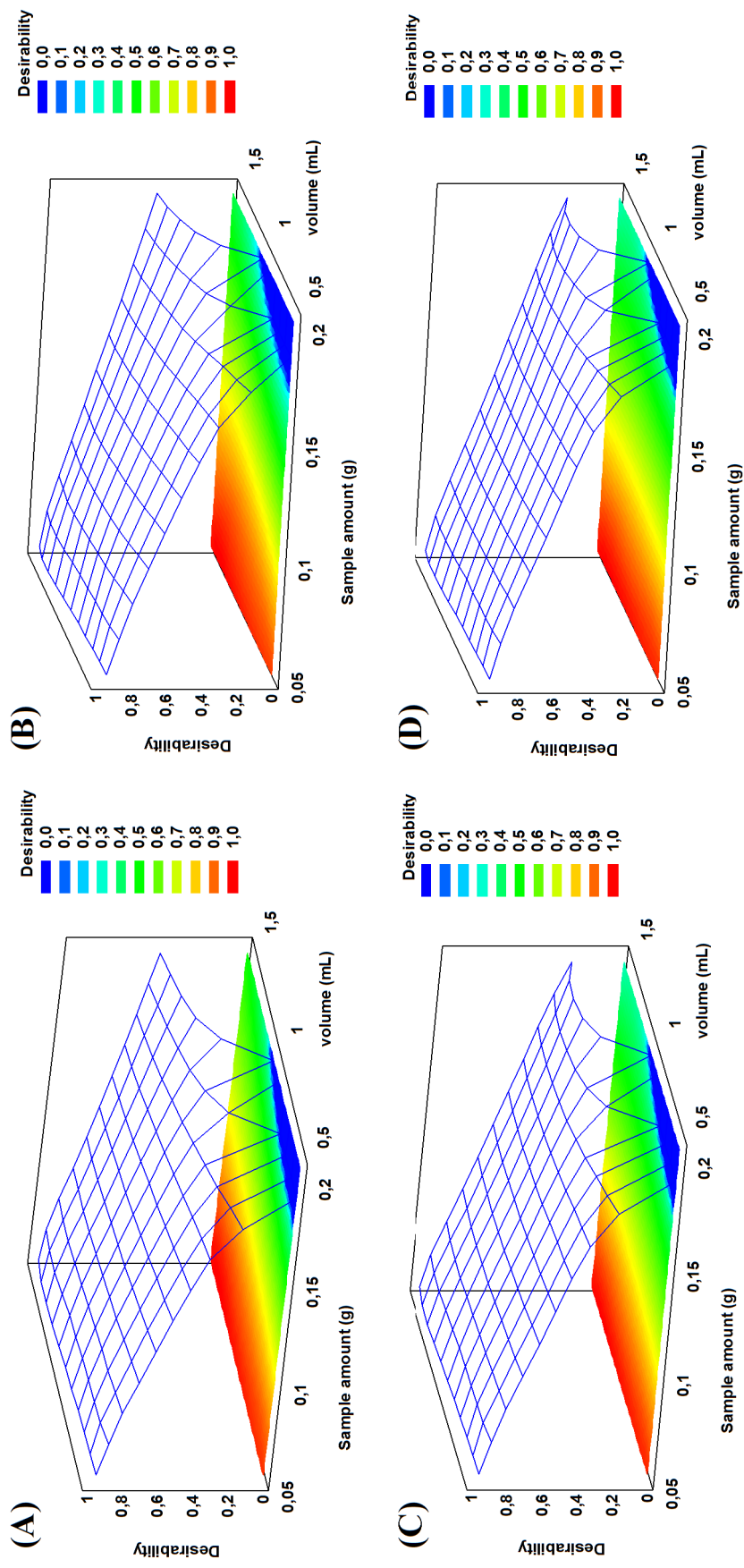
586 **Figure captions**

587

588 **Fig. 1.** Optimization of the sample amount, solvent volume and incubation time. Response  
589 surfaces obtained for Box–Behnken design: (A) BPA and chlorinated derivatives; (B)  
590 PBs; (C) BPs and (D) PFCs. Time of digestion was fixed in 12 hours.

591

592 **Fig. 2.** SMR chromatograms of: (A) a spiked sample standard mixture ( $75.0 \text{ ng g}^{-1}$  of each  
593 analyte), (B) a natural hair sample (Sample M04).



**Figure 1**

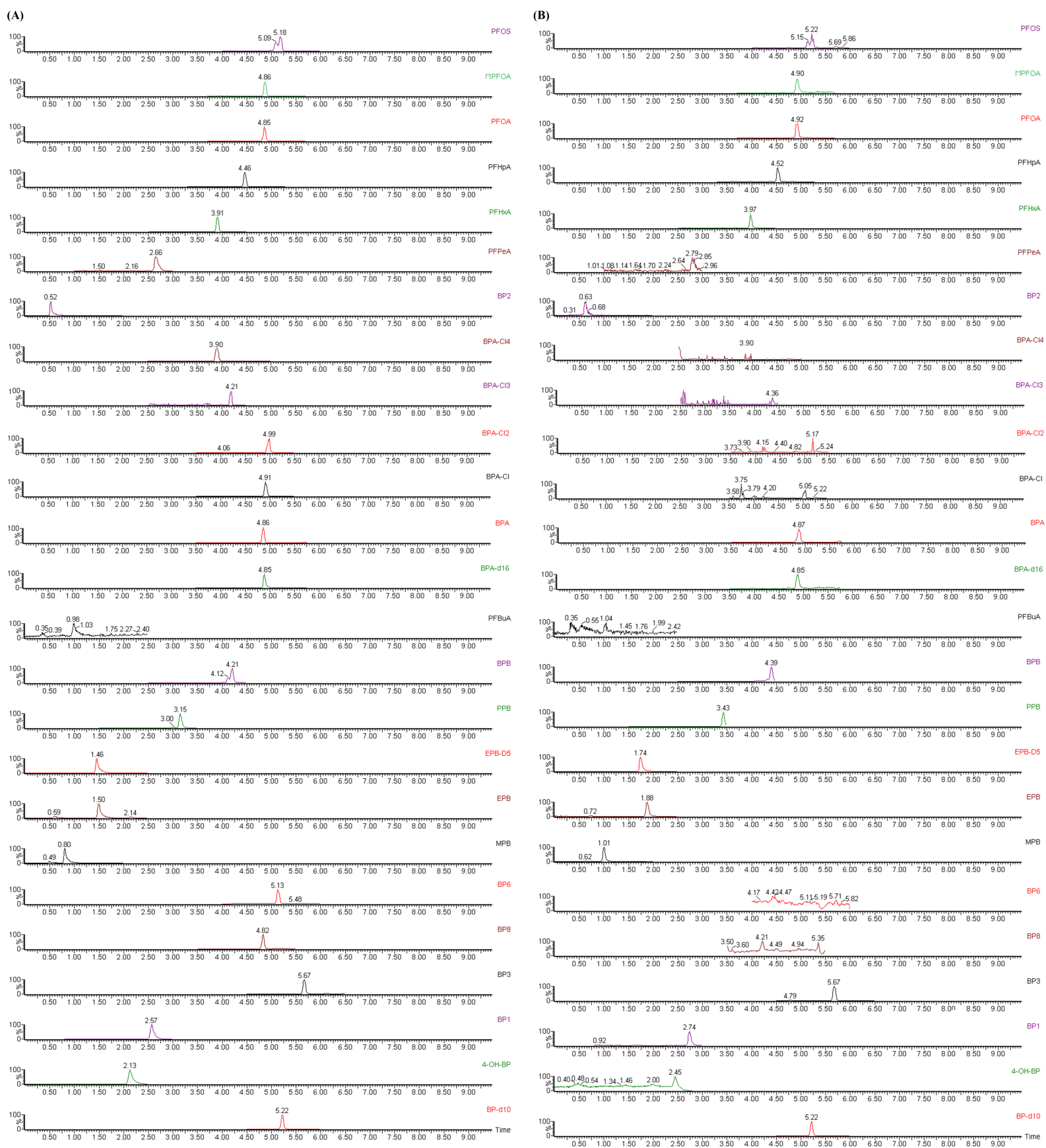


Figure 2

**Table 1**

Transitions and optimized potentials for UHPLC–MS/MS analysis

	Transitions	CV	CE		Transitions	CV	CE
BPA	227.2 → 211.9 <sup>a</sup>	-50	-22	BP-8	245.0 → 120.9 <sup>a</sup>	14	20
	227.2 → 132.9 <sup>b</sup>	-50	-26		245.0 → 150.9 <sup>b</sup>	14	20
Cl-BPA	261.1 → 182.0 <sup>a</sup>	-56	-30	4-OH-BP	199.0 → 120.8 <sup>a</sup>	36	20
	261.1 → 210.0 <sup>b</sup>	-56	-22		199.0 → 104.8 <sup>b</sup>	36	18
Cl <sub>2</sub> -BPA	295.0 → 215.9 <sup>a</sup>	-74	-30	PFBuA	213.0 → 169.0 <sup>a</sup>	36	20
	295.0 → 243.9 <sup>b</sup>	-74	-24		213.0 → 5.6 <sup>b</sup>	36	20
Cl <sub>3</sub> -BPA	329.0 → 249.8 <sup>a</sup>	-52	-32	PFPeA	263.0 → 219.0 <sup>a</sup>	36	20
	329.0 → 277.9 <sup>b</sup>	-52	-24		263.0 → 89.7 <sup>b</sup>	36	20
Cl <sub>4</sub> -BPA	365.0 → 313.9 <sup>a</sup>	-50	-28	PFHxA	313.0 → 269.0 <sup>a</sup>	36	20
	365.0 → 285.9 <sup>b</sup>	-50	-32		313.0 → 119.0 <sup>b</sup>	36	20
MPB	151.1 → 91.8 <sup>a</sup>	-38	-22	PFHpA	363.0 → 319.0 <sup>a</sup>	36	20
	151.1 → 135.8 <sup>b</sup>	-38	-14		363.0 → 333.0 <sup>b</sup>	36	20
EPB	165.1 → 91.9 <sup>a</sup>	-38	-24	PFOA	413.0 → 369.0 <sup>a</sup>	36	20
	165.1 → 136.6 <sup>b</sup>	-38	-16		413.0 → 194.0 <sup>b</sup>	36	20
PPB	179.1 → 91.8 <sup>a</sup>	-42	-24	PFOS	499.0 → 80.0 <sup>a</sup>	36	20
	179.1 → 136.1 <sup>b</sup>	-42	-16		499.0 → 52.0 <sup>b</sup>	36	20
BPB	193.1 → 91.4 <sup>a</sup>	-42	-24	BPA-d <sub>16</sub>	241.2 → 223.0 <sup>a</sup>	-46	-22
	193.1 → 136.1 <sup>b</sup>	-42	-16		241.2 → 141.9 <sup>b</sup>	-46	-32
BP-1	214.9 → 136.8 <sup>a</sup>	2	18	EPB-d <sub>5</sub>	170.1 → 92.1 <sup>a</sup>	-38	-24
	214.9 → 105.1 <sup>b</sup>	2	32		170.1 → 136.0 <sup>b</sup>	-38	-16
BP-2	245.1 → 134.8 <sup>a</sup>	-40	-16	BP-d <sub>10</sub>	193.1 → 109.8 <sup>a</sup>	18	16
	245.1 → 108.9 <sup>b</sup>	-40	-22		193.1 → 81.8 <sup>b</sup>	18	30
BP-3	229.0 → 150.8 <sup>a</sup>	4	20	MPFOA	417.0 → 371.0 <sup>a</sup>	36	20
	229.0 → 104.9 <sup>b</sup>	4	18		417.0 → 168.0 <sup>b</sup>	36	20
BP-6	275.0 → 150.9 <sup>a</sup>	14	18				
	275.0 → 94.9 <sup>b</sup>	14	34				

<sup>a</sup> SRM transition used for quantification; <sup>b</sup> SRM transition used for confirmation

CV: Cone voltage (V); CE: Collision energy (eV)

**Table 2**  
Analytical and statistical parameters

	b (g ng <sup>-1</sup> )	S <sub>b</sub> (g ng <sup>-1</sup> )	% R <sup>2</sup>	LOD (ng g <sup>-1</sup> )	LOQ (ng g <sup>-1</sup> )	LDR (ng g <sup>-1</sup> )
BPA	8.40 · 10 <sup>-4</sup>	2.72 · 10 <sup>-6</sup>	99.8	2	7	7 - 250
Cl-BPA	1.00 · 10 <sup>-3</sup>	8.72 · 10 <sup>-5</sup>	99.9	1	4	5 - 250
Cl <sub>2</sub> -BPA	3.30 · 10 <sup>-4</sup>	7.09 · 10 <sup>-6</sup>	99.9	3	9	9 - 250
Cl <sub>3</sub> -BPA	6.47 · 10 <sup>-4</sup>	3.43 · 10 <sup>-6</sup>	99.8	2	7	7 - 250
Cl <sub>4</sub> -BPA	9.50 · 10 <sup>-4</sup>	4.02 · 10 <sup>-6</sup>	99.9	3	9	9 - 250
MPB	5.24 · 10 <sup>-3</sup>	2.09 · 10 <sup>-5</sup>	99.9	0.2	0.5	0.5 - 125
EPB	3.50 · 10 <sup>-3</sup>	1.70 · 10 <sup>-5</sup>	99.9	0.6	2	2 - 125
PPB	7.20 · 10 <sup>-3</sup>	1.80 · 10 <sup>-4</sup>	99.8	0.3	1	1 - 125
BPB	6.00 · 10 <sup>-3</sup>	8.92 · 10 <sup>-5</sup>	99.9	0.4	1	1 - 125
BP-1	1.40 · 10 <sup>-1</sup>	3.45 · 10 <sup>-3</sup>	99.9	0.5	2	2 - 125
BP-2	8.00 · 10 <sup>-3</sup>	3.48 · 10 <sup>-5</sup>	99.9	0.3	1	1 - 125
BP-3	8.60 · 10 <sup>-2</sup>	3.50 · 10 <sup>-4</sup>	99.9	0.5	2	2 - 125
BP-6	3.96 · 10 <sup>-2</sup>	7.40 · 10 <sup>-4</sup>	99.9	0.7	2	2 - 125
BP-8	7.40 · 10 <sup>-2</sup>	1.00 · 10 <sup>-3</sup>	99.9	0.9	3	3 - 125
4-OH-BP	5.30 · 10 <sup>-1</sup>	1.45 · 10 <sup>-3</sup>	99.9	0.8	3	3 - 125
PFBuA	1.90 · 10 <sup>-3</sup>	2.57 · 10 <sup>-5</sup>	99.9	4	12	12 - 250
PFPeA	5.73 · 10 <sup>-3</sup>	3.45 · 10 <sup>-5</sup>	99.9	1	4	4 - 125
PFHxA	1.75 · 10 <sup>-2</sup>	1.20 · 10 <sup>-3</sup>	99.5	2	5	5 - 125
PFHpA	1.75 · 10 <sup>-2</sup>	2.80 · 10 <sup>-4</sup>	99.9	2	5	5 - 125
PFOA	3.00 · 10 <sup>-2</sup>	2.30 · 10 <sup>-3</sup>	99.4	1	4	4 - 125
PFOS	3.64 · 10 <sup>-2</sup>	5.00 · 10 <sup>-4</sup>	99.9	0.6	2	2 - 125

b, slope; S<sub>b</sub>, slope standard deviation; R<sup>2</sup>, determination coefficient; LOD, limit of detection; LOQ, limit of quantification; LDR, linear dynamic range

**Table 3**  
Recovery assay, precision and trueness of the method

	Spiked (ng mL <sup>-1</sup> )	Found <sup>a</sup> (ng mL <sup>-1</sup> )	RSD (%)	Recovery (%)		Spiked (ng mL <sup>-1</sup> )	Found <sup>a</sup> (ng mL <sup>-1</sup> )	RSD (%)	Recovery (%)
BPA	25.0	26.5	10.1	105.9	BP-3	5.0	4.6	9.3	92.1
	125.0	128.8	7.4	103.0		25.0	24.9	3.4	99.5
	250.0	247.9	3.6	99.2		125.0	126.3	3.8	101.0
Cl-BPA	25.0	24.1	8.7	96.2	BP-6	5.0	5.4	14.2	107.4
	125.0	130.9	13.4	104.7		25.0	26.8	2.5	107.2
	250.0	240.0	9.4	96.6		125.0	127.7	4.1	102.2
Cl <sub>2</sub> -BPA	25.0	24.3	8.7	97.2	BP-8	5.0	5.7	5.3	113.8
	125.0	121.3	4.3	97.1		25.0	24.8	0.8	99.1
	250.0	241.2	2.6	96.4		125.0	119.7	12.2	95.8
Cl <sub>3</sub> -BPA	25.0	27.0	8.9	108.0	4-OH-BP	5.0	5.0	5.3	99.0
	125.0	135.9	5.6	107.1		25.0	24.7	6.1	98.9
	250.0	258.0	4.3	103.2		125.0	128.0	9.7	102.0
Cl <sub>4</sub> -BPA	25.0	26.5	5.0	106.0	PFBuA	25	24.6	1.1	98.4
	125.0	124.9	8.3	99.9		125	126.5	2.9	101.2
	250.0	250.0	10.8	100.0		250	249.5	10.0	99.8
MPB	5.0	5.4	8.8	107.2	PFPeA	5.0	4.9	4.9	98.5
	25.0	25.0	10.8	100.1		25.0	25.0	7.4	100.0
	125.0	128.5	3.4	102.8		125.0	129.5	5.1	103.6
EPB	5.0	5.2	6.3	103.5	PFHxA	5.0	4.4	7.8	88.0
	25.0	23.8	7.0	95.3		25.0	24.0	8.9	96.1
	125.0	125.8	4.3	100.6		125.0	126.7	3.3	101.3
PPB	5.0	5.5	12.5	109.7	PFHpA	5.0	5.5	10.3	110.1
	25.0	26.0	6.5	104.1		25.0	23.8	5.7	95.1
	125.0	123.0	5.6	98.4		125.0	125.2	5.3	100.1
BPB	5.0	4.9	10.9	97.8	PFOA	5.0	4.9	9.1	99.1
	25.0	24.7	7.8	98.7		25.0	26.7	5.1	106.9
	125.0	125.0	9.1	100.0		125.0	141.0	5.4	112.8
BP-1	5.0	4.9	14.5	98.5	PFOS	5.0	5.5	2.0	109.4
	25.0	24.5	10.9	98.0		25.0	25.8	9.6	103.2
	125.0	127.4	11.9	101.9		125.0	128.6	2.5	102.9
BP-2	5.0	5.6	13.9	108.9					
	25.0	27.0	12.7	107.8					
	125.0	127.0	3.7	101.5					

<sup>a</sup> Mean of 18 determinations; RSD: relative standard deviation

**Table 4**

Application to hair samples

*Mean amount for each volunteer (SD) ng g <sup>-1</sup>							
	BPA	MPB	EPB	PPB	BPB	BP-1	BP-2
M01	ND	21 (1)	9.0 (0.3)	69 (4)	5.8 (0.3)	ND	2.8 (8.1)
M02	ND	ND	ND	ND	3.5 (0.3)	4.3 (0.2)	D
M03	45 (3)	33 (2)	ND	107 (4)	4.4 (0.3)	116 (4)	1.5 (5.0)
M04	12.3 (0.6)	18.7 (0.6)	D	80 (6)	9.4 (0.5)	29 (1)	5.8 (4.1)
M05	ND	ND	ND	ND	ND	ND	ND
M06	9.2 (0.6)	10.2 (0.6)	9.0 (0.6)	11.6 (0.6)	5.7 (0.3)	12.5 (0.6)	D
	BP-3	4-OH-BP	PFPHxA	PFPHpA	PFOA	PFOS	
M01	D	ND	D	D	10.6 (0.5)	4.9 (0.4)	
M02	7.0 (0.3)	ND	D	D	14.8 (0.6)	5.4 (0.2)	
M03	8.5 (0.5)	D	D	D	23.9 (0.7)	9.9 (0.5)	
M04	5.2 (0.2)	D	5.5 (0.4)	7.1 (0.4)	20.0 (0.9)	6.8 (0.3)	
M05	D	ND	ND	ND	D	D	
M06	10.5 (0.7)	ND	D	D	18.0 (0.7)	6.8 (0.4)	

\* Mean of 36 analysis (12 samples per volunteer / 3 replicates a day); ND, not detected (< LOD); D: detected (>LOD and <LOQ); SD, standard deviation