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Title: OCCURRENCE OF COMMON PLASTIC ADDITIVES AND CONTAMINANTS IN MUSSEL SAMPLES: VALIDATION OF ANALYTICAL METHOD BASED ON MATRIX SOLID-PHASE DISPERSION

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Corresponding Author: Dr. Rosa M^a Garcinuño, Dr

Corresponding Author's Institution: Facultad de Ciencias, UNED

First Author: Raquel Cañadas

Order of Authors: Raquel Cañadas; Esther Garrido; Rosa M^a Garcinuño, Dr; Gema Paniagua; Pilar Fernández

Abstract: A new matrix solid-phase dispersion (MSPD) extraction methodology, combined with high-performance liquid chromatography equipped with a diode-array detector, was developed and validated for the simultaneous determination of 10 compounds in mussels from Galician Rias (Spain). These pollutants are compounds commonly used for plastic production as additives, as well as common plastic contaminants. The compounds selected were bisphenol-A, bisphenol-F, bisphenol-S, nonylphenol-9, nonylphenol, diethyl phthalate, dibutyl phthalate, di-2-ethylhexyl phthalate, dichlorodiphenyltrichloroethane, dichlorodiphenyldichloroethane, and dichlorodiphenyldichloroethylene. The parameters affecting the MSPD extraction efficiency such as the type of sorbent, mass sample-sorbent ratio, and extraction solvent were optimised. The proposed method provided satisfactory quantitative recoveries (80–100%), with relative standard deviations lower than 7%. In all cases, the matrix-matched calibration curves were linear in the concentration range of 0.32–120.00 µg/kg, with quantification limits of 0.25–16.20 µg/kg. The novel developed MSPD-high-performance liquid chromatography methodology provided good sensitivity, accuracy, and repeatability for quality control analysis in mussels.

Highlights

- A novel approach for endocrine-disrupting chemicals determination in mussels.
- Development of a matrix solid phase dispersion methodology for mussel samples.
- Simultaneous separation and quantification of 11 EDCs by HPLC-DAD.
- Limits of quantification ranged between 0.25-16.20 µg/Kg.

1 **OCCURRENCE OF COMMON PLASTIC ADDITIVES AND CONTAMINANTS IN**
2 **MUSSEL SAMPLES: VALIDATION OF ANALYTICAL METHOD BASED ON**
3 **MATRIX SOLID-PHASE DISPERSION**

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5 R. Cañadas, E. Garrido Gamarro¹, R. M. Garcinuño Martínez*, G. Paniagua González*,
6 P. Fernández Hernando

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8
9 *Departamento de Ciencias Analíticas, Facultad de Ciencias, Universidad Nacional de*
10 *Educación a Distancia (UNED); Paseo Senda del Rey nº 9, 28040, Madrid, Spain*

11
12 ¹*Food Safety and Quality Officer, Fisheries and Aquaculture, Department, Food and*
13 *Agriculture Organization of the United Nations (FAO); Viale delle Terme di Caracalla,*
14 *00153 Roma RM, Italy*

15
16
17
18
19
20
21
22 *Corresponding authors: Dra. Rosa M^a Garcinuño Martínez
23 Address: C/ Senda del Rey nº 9, 28040 Madrid, Spain.
24 Telephone number: + 34 91 398 73 66
25 E-mail address: rmgarcinuno@ccia.uned.es

26
27 Dra. Gema Paniagua González
28 Address: C/ Senda del Rey nº 9, 28040 Madrid, Spain.
29 Telephone number: + 34 91 398 72 71
30 E-mail address: gpaniagua@ccia.uned.es
31

35 **Abstract**

36 A new matrix solid-phase dispersion (MSPD) extraction methodology, combined with high-
37 performance liquid chromatography equipped with a diode-array detector, was developed and
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65 **1. Introduction**

66 Plastic production has increased significantly since the 1950s, reaching 359 million tonnes in
67 2018. Due to mismanagement of plastic waste, estimates suggest that between 4.8 million
68 tonnes to 12.7 million tonnes of plastics entered the ocean in 2010 (Jambeck et al., 2015). An
69 added problem is the fact that plastics can be fractionated into millimetre-sized particles called
70 microplastics (MP), usually with a size of $<5 \mu\text{m}$ (Ivleva, Wiesheu & Niessner, 2016), where MP
71 are recognised as emerging contaminants due to fact that the smaller microplastics as well as
72 nanoplastics (from $0.001 \mu\text{m}$ to $0.1 \mu\text{m}$) could potentially penetrate the capillaries of
73 organs (Yoo, Doshi, & Mitragotri, 2011). Dumping of plastics in the ocean has led to the
74 ingestion of microplastics by aquatic animals, and to the presence of microplastics in common
75 fisheries and aquaculture commodities. The composition of microplastics and their capability
76 to sorb (either absorb or adsorb) and desorb chemicals such as
77 dichlorodiphenyldichloroethane (DDD), dichlorodiphenyldichloroethylene (DDE), and
78 dichlorodiphenyltrichloroethane (DDT), which are known to be persistent, bioaccumulative,
79 and toxic substances (PBTs), from the surrounding environment could pose a food safety
80 threat.

81 Microplastics tend to accumulate in the gastrointestinal tract of aquatic animals, where
82 plastics smaller than $1.5 \mu\text{m}$ can cross the gastrointestinal barrier. Thus, certain food
83 commodities consumed whole are of special interest in terms of dietary exposure to
84 microplastics and associated contaminants (Garrido et al., 2020). The consumption of
85 invertebrates that feed by filtration, such as bivalve molluscs (mussels, clams, oysters, etc.),
86 seems the most likely route of exposure (Li et al., 2019). Wild mussels are complex samples,
87 rich in water, proteins, vitamins, and minerals, where their composition is affected by their
88 environment. Wild mussels have been commonly used as bioindicators for monitoring
89 pollutants in coastal waters due to their extensive distribution, easy sampling, abundance,
90 their ventilation of large volumes of water for nutrition, respiration, and excretion, low
91 mobility, and ecological and economic importance (Chiesa et al., 2018).

92 Several countries are building regulatory frameworks to reduce the production and use of
93 plastics, as well as the impact of plastic composition in food safety. For instance, the European
94 Union (EU) has adopted different directives and regulations in recent years, such as the new
95 EU Directive 2018/852 (European Parliament, 2018), prohibiting single-use packaging and
96 waste. The EU legislation also set limits for bisphenol-A (BPA) in food-contact plastic materials,

97 establishing a specific migration limit of 0.05 mg of BPA per kilogram of food (European
98 Commission, 2018). For *bis*(2-ethylhexyl) phthalate (DEHP) and dibutyl phthalate (DBP), the EU
99 legislation specifically established limits in products intended for food contact of 1.5 mg/kg for
100 DEHP and 0.3 mg/kg for DBP in its Commission Regulation (EU) N° 10/2011 (European
101 Commission, 2011). Since February 2015, the production of phthalate esters has been
102 forbidden, unless authorisation has been granted for a specific use, although these compounds
103 may still be imported in consumer products (European Commission, 2015).

104 The effects of exposure to the specific compounds mentioned above have been studied, where
105 these compounds are known to be endocrine disruptor chemicals (EDCs) or PBTs (bisphenol-A,
106 *bis*(2-ethylhexyl) phthalate, nonylphenols (NPs), polychlorinated biphenyls (PCBs), and
107 organochlorine pesticides), but others such as bisphenol-F (BPF), bisphenol-S (BPS), and diethyl
108 phthalate (DEP) are still to be evaluated (European Commission, 2008; EFSA 2017). To
109 understand the magnitude of the problem and the possible implications of these chemicals in
110 the environment and in food safety, it is necessary to develop standardised analytical
111 methodologies to promote consistency in the results obtained. The most used analytical
112 methodologies for the determination of these compounds are gas chromatography (GC)
113 (Filipkowska & Lubecki, 2016; Sánchez-Avila et al., 2011; Wang et al., 2010) and liquid
114 chromatography (LC) (Yang et al., 2014; Salgueiro-González et al., 2016; Luo et al., 2018). High-
115 performance liquid chromatography (HPLC) coupled with different detectors (diode-array
116 detector (DAD) and/or mass detector (MS)) provides advantages over the GC methods, such as
117 simple sample preparation without the need for derivatisation techniques. The liquid
118 chromatography detection method has previously been used for the analysis of EDCs in food
119 samples such as oil (Xian et al., 2017), eggs (Song et al., 2019), hotpot seasoning (Dong, Zeng &
120 Bai, 2018), fish (Tran et al., 2019), and mussels (Ocharoen et al., 2018). The most frequently
121 applied extraction techniques are solid-phase extraction (Azzouz et al., 2019; Filipkowska &
122 Lubecki, 2016; Yang et al., 2014), Soxhlet extraction (Movahedinia, Salamat & Kheradmand,
123 2018), sonication (Blackburn, Kirby & Waldock, 1999), and pressurised liquid extraction
124 (Salgueiro-González et al., 2016). However, these extraction methodologies generally require a
125 clean-up step to reduce the presence of interferents during the subsequent chromatographic
126 determination. Matrix solid-phase dispersion (MSPD) is an effective alternative technique to
127 traditional methods for sample preparation. It offers several advantages such as simplicity and
128 flexibility. This methodology integrates disruption, homogenisation, extraction, and clean-up of
129 the sample into one step (Barker et al., 2007; Qi, 2010; García-Mayor et al., 2012). Sample
130 preparation methods based on MSPD have been efficiently applied to the determination of a

131 wide variety of micropollutants (PCBs, DEPH, ethoxylates, pesticides, etc.) in complex
132 biological matrices such as mussels (Carro et al., 2017; Rodríguez-González et al., 2015;
133 Rombaldi, 2015; Wianowska & Gil, 2019). However, to date, MSPD has not been applied for
134 the simultaneous extraction of endocrine disruptors, including different groups of compounds,
135 (bisphenols, phthalates, etc.) from food samples.

136 The aim of this work is to develop an effective, simple, and reliable multi-residue analytical
137 methodology based on a matrix solid-phase dispersion extraction technique (MSPD) coupled
138 to high-performance liquid chromatography with a diode-array detector (HPLC-DAD) for the
139 quantitative extraction and simultaneous determination of several EDCs, including common
140 plastic additives (bisphenols (BPA, BPF, and BPS), phthalate esters (DBP, DEHP, and DEP),
141 alkylphenols (NP and NP-9), and other organic pollutants such as organochlorine pesticides
142 (DDD, DDE, and DDT) in wild raw mussel samples. The chemicals studied were selected
143 because they are commonly present in microplastics and some of them are known to be highly
144 potent EDCs that are environmentally persistent (McCombe, 2020) and have already been
145 detected in wild raw mussels (Salgueiro-González et al., 2016; Suárez et al., 2013), indicating
146 their potential for bioaccumulation.

147 **2. Material and methods**

148 2.1. Chemicals

149 All reagents were of analytical grade unless specified otherwise. Water (18 M Ω /cm) was
150 purified with a Milli-Q water system (Millipore Ibérica, Madrid, Spain). Bisphenol-A (BPA)
151 (purity \geq 99%), bisphenol-F (BPF) (purity \geq 98%), bisphenol-S (BPS) (purity 98%), diethyl
152 phthalate (DEP) (purity \geq 99%), dibutyl phthalate (DBP) (purity 99%), *bis*(2-ethylhexyl)
153 phthalate (DEHP) (purity \geq 99.5%), nonylphenol (NP) (purity \geq 99%), 9-*n*-nonylphenol (NP-9)
154 (purity \geq 99%), dichlorodiphenyldichloroethane (DDD), dichlorodiphenyldichloroethylene
155 (DDE), and dichlorodiphenyltrichloroethane (DDT) were supplied by Sigma-Aldrich (Madrid,
156 Spain). Stock standard solutions (1000 mg/L) of each analyte were prepared in acetonitrile
157 (ACN), except for the nonylphenols, which were prepared in methanol (MeOH). The standard
158 solutions were stored in dark bottles at 4 °C until use and remained stable for at least three
159 months. Organic solvents (acetonitrile, methanol, and hexane) and sodium hydroxide were
160 purchased from Scharlab (Madrid, Spain). The solid-phase materials used for MSPD were
161 Florisil from Sigma-Aldrich (Madrid, Spain), anhydrous sodium sulfate (Na₂SO₄) from Panreac
162 (Barcelona, Spain), and washed sea sand (0.25–0.30 mm) from Symta (Madrid, Spain). The
163 silanised glass wool used herein was supplied by Panreac (Barcelona, Spain). All solutions

164 collected from the MSPD procedure were filtered through 0.22 µm Teflon membrane disc
165 filters from Merck (Madrid, Spain).

166 2.2. Mussel sample collection and preparation

167 Mussels (*Mytilus galloprovincialis*), which are common species from Arousa stuary in Galicia
168 (Northwest, Spain), from aquaculture production were used as the raw material. A total of 30
169 live specimens, representing this type of species, were purchased from a local market. The
170 samples were immediately transported to the laboratory, protected from possible
171 contamination, and refrigerated for sample preparation. The samples were dissected with a
172 clean scalpel blade to separate the mussel from the shell. The whole raw mussels were then
173 homogenised and stored at 4 °C prior to the analyses. Measures were taken to avoid
174 contamination by the equipment used for sample collection, transportation, and preparation.

175 2.3. Matrix solid-phase dispersion procedure

176 The analytes were extracted from the biomatrix by performing the MSPD procedure with
177 spiked and blank mussel samples. The optimal extraction process was as follows: A 0.1 g
178 sample of homogenised raw mussels was accurately weighed and spiked with 100 µL of stock
179 standard solution containing the analytes at the desired concentration (for instance, NP and
180 NP-9 were spiked at 120 mg/L and the rest of the analytes (BPS, BPF, BPA, DEP, DBP, DDD,
181 DDT, DDE, and DEHP) at 40 mg/L in MeOH/H₂O (85:15, v/v). The mix was maintained at room
182 temperature for 20 min to allow the solution to equilibrate with the mussel matrix. The spiked
183 mussel sample was poured into a glass mortar containing 0.5 g of Florisil (dispersing sorbent,
184 <200 mesh), 0.5 g of Na₂SO₄ (anhydrous agent, purity ≥99%), and 0.2 g of washed sea sand. To
185 achieve complete disruption and dispersion of the biosample on the solid support, the mixture
186 was blended for approximately 10 min using a glass pestle, until an apparently dry and
187 homogeneous material was obtained. The homogenised mixture was then packed into a glass
188 cartridge with a plug of silanised glass wool at the bottom, which retained the entire sample. A
189 small amount of Na₂SO₄ (0.001–0.002 g) was placed on top of the mixture, just enough to
190 cover it. To ensure homogeneous packing of the column, care was taken to pour the material
191 into the tube in several portions and remove the air pockets inside the material. The sample
192 was first conditioned using 1 mL of acetonitrile. The analytes were optimally eluted from the
193 MSPD column dropwise by gravity, with 9 mL of methanol/acetonitrile (30:70, v/v), in three 5
194 min static extraction steps. The flow rate was constant at approximately 0.6 mL/min. The
195 extracts were collected and evaporated to dryness under a nitrogen stream at room
196 temperature. Finally, the residue was reconstituted in 400 µL of MeOH/H₂O (85:15, v/v) and an

197 aliquot of 20 μL was injected into the chromatographic system. The corresponding blank
198 sample was prepared by following the same procedure, excluding the spiked analytes from the
199 MSPD mixture. In addition, a solvent blank was prepared to check for background
200 contamination due to the use of laboratory plastic material.

201 2.4. HPLC-DAD conditions

202 Chromatographic separation of the analytes was performed using an Agilent Technologies
203 model 1200 series liquid chromatograph (Agilent Technologies, Germany) equipped with an
204 on-line degasser, a quaternary pump, autosampler, and a photo-diode array detector (DAD).
205 The analytes were separated on an ACE 5 C18-PFP HPLC column (150×4.6 mm, $5 \mu\text{m}$) from
206 Symta (Madrid, Spain). The column and auto sampler were maintained at room temperature.
207 The mobile phase consisting of a mixture of Milli-Q ultrapure water (solvent A) and acetonitrile
208 (solvent B) was set to a gradient programme: 0–30 min: 45–80% B; 30–31 min: 80–100% B;
209 and 100% B for 9 min, at a flow rate of 0.8 mL min^{-1} . Subsequently, the column was
210 reconditioned with 45% B under isocratic conditions for 10 min. The sample injection volume
211 was $20 \mu\text{L}$. All compounds were successfully separated within 40 min. To achieve the maximum
212 sensitivity, quantitative measurements of the peak areas were performed by selecting the
213 optimum detection wavelength for each compound. All analytes were therefore quantified at
214 210 nm. Quantification was performed using external calibration and peak area
215 measurements.

216 3. Results and discussion

217 3.1. Optimisation of conditions for HPLC-DAD analysis

218 Preliminary studies were performed using a chromatography Eclipse Plus C18 $3.5 \mu\text{m}$ ($100 \times$
219 4.6 mm) column for separation of the selected compounds. Different mobile phases
220 comprising water (eluent A) and acetonitrile or methanol (eluent B) at different ratios (20, 30,
221 50 and 70%) were tested. In this preliminary study, the signals of the analytes in the
222 chromatograms were not satisfactorily resolved.

223 To improve the feasibility for elution of all analytes, a new column, an ACE 5 C18-PFP HPLC
224 $5 \mu\text{m}$ (150×4.6 mm), having a slightly modified composition and length, was tested. Different
225 eluent gradient profiles using water (solvent A) and different ratios of methanol or acetonitrile
226 (solvent B) were tested to obtain adequate separation and resolution of the analytes. Finally, a
227 gradient programme was selected, which consisted of 45% ACN initially, followed by an
228 increase to 80% ACN in the first 30 min. The composition of the eluent was increased to 100%

229 ACN for 1 min, and kept stable at 100% ACN for 4 min. The system was allowed to equilibrate
230 for 10 min at 45% of ACN. The resulting total run time was 40 min. The flow rate was modified
231 between 0.5 and 1.0 ml/min. Because higher flow rates did not significantly improve either the
232 sensitivity or retention time, a flow rate of 0.8 ml/min was selected. The injection volume was
233 varied from 15 to 25 μ L. A final volume of 20 μ L was selected to increase the sensitivity
234 without any loss of resolution.

235 In order to achieve the optimum analytical signal for each analyte, the absorption with multi-
236 wavelength detection at 210, 230, 250, 254, and 280 nm using DAD was evaluated by
237 integrating the peaks manually. The maximum areas for all analytes was achieved at 210 nm.
238 The chromatogram corresponding to HPLC-DAD separation of a standard mixture of the
239 compounds under the optimum conditions is shown in Fig. 1. Separation of the eleven
240 compounds was achieved within 35 min.

241 3.2 Optimisation of MSPD extraction procedure

242 Selection of the most favourable extraction conditions is crucial for achieving adequate
243 efficiency of the MSPD procedure. Therefore, the dispersing sorbent, the sample mass/sorbent
244 ratio, and the type and volume of the eluent were optimised to obtain clean final extracts and
245 to achieve the highest analyte recoveries and the lowest matrix interferences from mussel
246 samples. Three types of dispersing sorbents (Florisil, Alumina, and C-18) at three loadings (0.5,
247 1.0, and 1.5 g) were tested with an initial mussel sample mass of 0.5 g and a fixed amount of
248 0.5 g of Na_2SO_4 and 0.2 g of washed sea sand to improve the sample dispersion. The amount of
249 mussel sample (0.5, 0.3, and 0.1 g) was then optimised using the optimum 0.5 g of Florisil as a
250 clean-up/dispersing sorbent, 0.5 g of Na_2SO_4 , and 0.2 g of washed sea sand. Finally, 0.1 g of
251 mussel sample, 0.5 g of Florisil as a dispersing sorbent, 0.5 g of Na_2SO_4 , and 0.2 g of washed
252 sea sand were chosen as the optimal MSPD mixture. In these mentioned studies, the mussel
253 sample was spiked with 40 ppm of BPS, BPF, BPA, DEP, DBP, DDD, DDT, and DDE, and 120 ppm
254 of NP-9, and dispersed with the corresponding amount of Florisil, Na_2SO_4 , and washed sea
255 sand. The homogenised mixtures in each assay were then transferred to a glass cartridge to
256 prepare the MSPD columns by following the procedure described in Section 2.3. Thereafter,
257 the column was conditioned with 1 mL of methanol and the analytes were extracted with 9 mL
258 of methanol as the starting solvent. Subsequently, the type and volume of the eluent was
259 optimised. The selection of an adequate eluent to desorb the analytes from the MSPD column
260 has an important role in this extraction procedure. A set of assays to test the feasibility of
261 several solvents with different polarity (ACN, MeOH) and mixtures (MeOH/ H_2O (80/20) and

262 MeOH/ACN (80/20, 50/50, 40/60, 30/70, 20/80)) was carried out. Different volumes (3, 6, 9,
263 12, and 15 mL) of eluent that afforded the elution of all analytes were tested in static cycles of
264 5 min using 3 mL of eluent per cycle. Prior to the elution process, in all cases, the column was
265 conditioned with 1 mL of the test eluent. The obtained extracts were treated as indicated in
266 Section 2.3 before injection into the HPLC system, and analysed by following the procedure
267 outlined in Section 2.4. From the tests, it was concluded that 9 mL was the most efficient
268 eluent volume for recovering the analytes. Figure 2 shows the results for these experiments
269 using 9 mL of eluent in consecutive 5 min static extraction cycles (3 mL per cycle). When
270 methanol was used as the eluent, the recovery for all analytes was in the range of 70–96%,
271 except for DEP, for which extraction was not possible. When the methanol/water mixture
272 (80:20, v/v) was used, the recovery was 70–100% for most analytes. However, DEP and NP-9
273 were not extracted at all. The use of acetonitrile as the eluent allowed extraction of all the
274 analytes, with recoveries of approximately 100% for some of them, such as DEP, DBP, and NP,
275 and recoveries ranging between 54 and 82% for BPS, BPF, BPA, DDT, DDD, DDE, and DEHP. In
276 this assay, it was also possible to detect NP-9 with a recovery of approximately 30%. When
277 methanol/acetonitrile mixtures were used, it was not possible to obtain quantitative recovery
278 of all the analytes in any case, and some of them were not isolated (DEP with 80:20
279 methanol/acetonitrile; NP-9 with 50:50, 40:60, 30:70, 20:80 methanol/acetonitrile).
280 Considering the results (Fig. 2), and to develop a methodology capable of simultaneously
281 determining the greatest number of analytes with quantitative recovery, the mixture of 30:70
282 methanol/acetonitrile was selected as the eluent for method validation. In this case, the
283 recovery for all detected analytes was in the range of 80–100%. Based on the results obtained
284 by HPLC-DAD analysis, it can be concluded that 0.1 g of mussel sample, 0.5 g of Florisil, 0.5 g of
285 Na₂SO₄, 0.2 g of washed sea sand, and 9 mL of 30:70 methanol/acetonitrile eluent are the
286 most favourable extraction conditions. These conditions afforded the highest intensity
287 chromatographic analyte peaks, indicating a greater percentage recovery of the analytes.

288 In order to further improve the recovery of the analytes, the possibility of using an additional
289 washing step was evaluated using 0.01 M sodium hydroxide or hexane. For these assays, 3 mL
290 of 0.01 M sodium hydroxide solution or hexane was added to the cartridge to eliminate
291 possible matrix interferences, and elution with methanol/acetonitrile 30:70 was then
292 accomplished. These additional washing steps did not improve the previously obtained results,
293 and in the case of sodium hydroxide, even made it impossible to analyse the samples due to
294 hydrolysis of the phthalic acid esters.

295 Figure 3 presents a comparison of the chromatograms for an enriched mussel sample (where
296 30 mg/L of nonylphenols and 10 mg/L of bisphenols, phthalates, and pesticides were added)
297 and an unenriched mussel sample (blank sample) subjected to the developed method. The
298 chromatogram of a standard solution at the concentration level of the spiked samples is
299 included in Fig. 3. Signals corresponding to DBP and DEHP were observed in the chromatogram
300 of the blank. Hence, the quantification of these analytes was carried out by subtracting the
301 blank signal from the sample signal. The peaks of both analytes were confirmed by mass
302 spectrometry.

303 3.3. Method validation

304 The developed MSPD method was validated in terms of linearity, precision, recovery, and the
305 limits of detection and quantification, using spiked mussel samples. Calibration plots were
306 constructed by preparing spiked mussel samples in triplicate, where the samples contained
307 increasing concentrations of each compound, in the corresponding range (Table 1). The results
308 showed good linearity for all analytes, with correlation coefficients (R^2) of 0.9873–0.9991.

309 The precision was calculated in terms of the intra-day repeatability ($n = 3$) and inter-day
310 reproducibility (three successive days) from triplicate assays at three spiking levels (between
311 2.00 and 120.00 $\mu\text{g}/\text{kg}$). The lowest spiking level for each analyte was always higher than the
312 corresponding LOQ, which allowed correct quantification of its recovery. The intra-day
313 repeatability, expressed as the relative standard deviation (RSD), ranged from 0.8% to 6.9%.
314 The inter-day repeatability was lower than 6.6% for all analytes. These results demonstrate
315 that the analytical method is sufficiently accurate. As shown in Table 1, good recoveries
316 (80–100%) were obtained for all investigated compounds, except for nonylphenol-9. The non-
317 quantification of NP-9 is caused by the ill-defined chromatographic peak. The peak of NP-9 was
318 broken down into smaller associated peaks, forming a wide peak without an observable clear
319 maximum (Fig. 1).

320 The limits of detection (LODs) and quantification (LOQs) were calculated at signal-to-noise
321 ratios of 3 and 10, respectively, by applying the following equations: $\text{LOD} = 3.3 \times \sigma/S$ and $\text{LOQ} =$
322 $10 \times \sigma/S$, where σ is the standard deviation of 3 samples spiked at the experimental estimated
323 LOQ, and S is the slope of the corresponding calibration curve. The obtained results were
324 calculated according to the FDA Guidance for Industry (FDA, 2015). The LODs of the detected
325 analytes were between 0.06 and 6.00 $\mu\text{g}/\text{kg}$, and the LOQs were between 0.25 and 16.22
326 $\mu\text{g}/\text{kg}$ (see Table 1).

327 4. Conclusions

328 The novel developed method proposed in this research illustrates the potential of combining
329 MSPD extraction with HPLC-DAD for the analysis of traces of endocrine disruptors in mussel
330 samples. This MSPD-HPLC-DAD method provides good sensitivity, accuracy, and repeatability
331 for quality control analysis in real seafood samples. For most of the analytes, satisfactory
332 recoveries of 80–100% were achieved, with an RSD of less than 7% and LODs and LOQs of
333 0.06–6.00 µg/kg and 0.25–16.22 µg/kg, respectively. Advantageously, the proposed
334 methodology requires a small amount of mussel sample (0.1 g) for extraction and does not
335 require an additional clean-up stage, as it offers the possibility to carry out sample extraction
336 and clean-up steps at the same time. These advantages make the method a promising
337 alternative to other time-consuming and multi-step analytical procedures. Furthermore, to the
338 best of our knowledge, there is no documentation of methods implementing MSPD and HPLC-
339 DAD for the sensitive simultaneous analysis of this set of compounds in mussel samples.

340 The determination of plastic additives and plastic contaminants, derived from the
341 contamination of production waters, in fishery and aquaculture products requires further
342 research. The developed method constitutes a significant advance in terms of simplicity,
343 efficiency, and sensitivity, allowing realisation of the first potential simultaneous
344 determination of ten chemical compounds from microplastics that are commonly present in
345 mussel samples.

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464

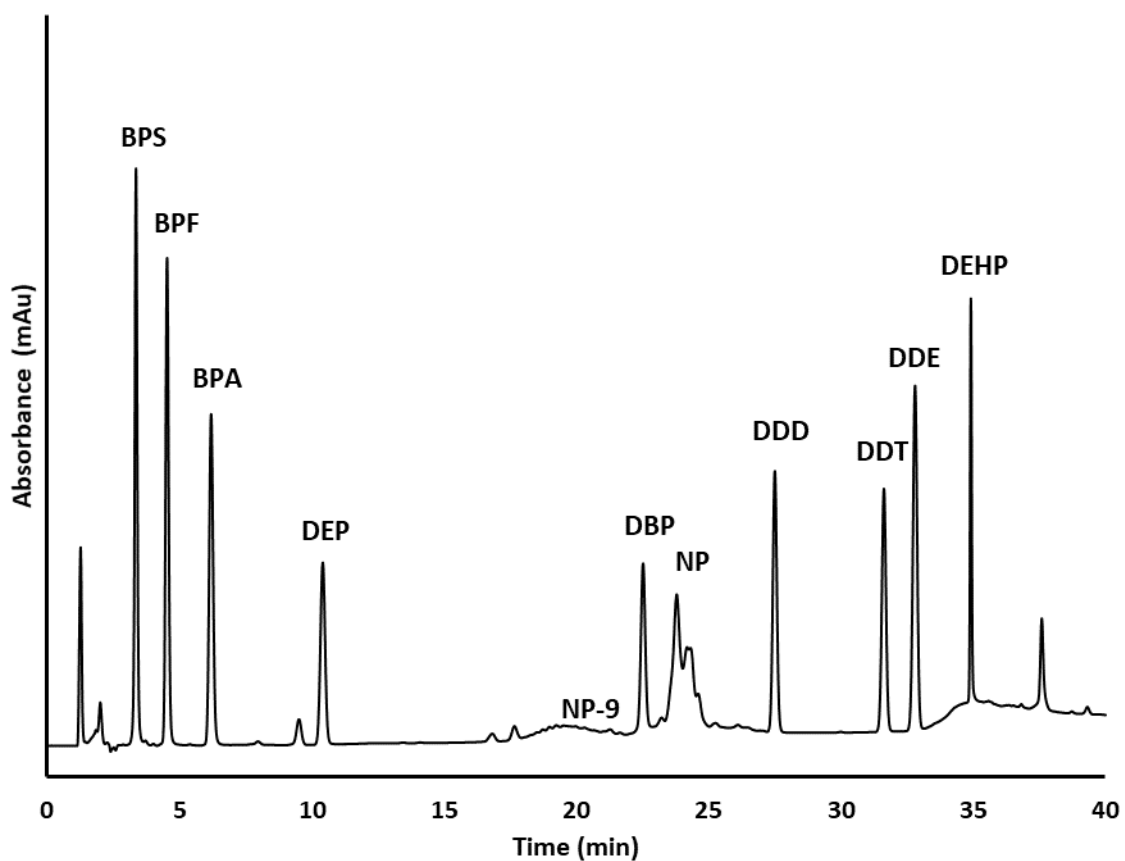


Fig. 1. Chromatogram obtained at 210 nm for standard mixture of EDCs (10 mg/L of BPS, BPF, BPA, DEP, DBP, DDD, DDT, DDE, and DEHP, and 50 mg/L of NP and NP-9) using HPLC-UV-DAD with MeOH/H₂O (85/15, v/v). The optimum experimental conditions are described in the text.

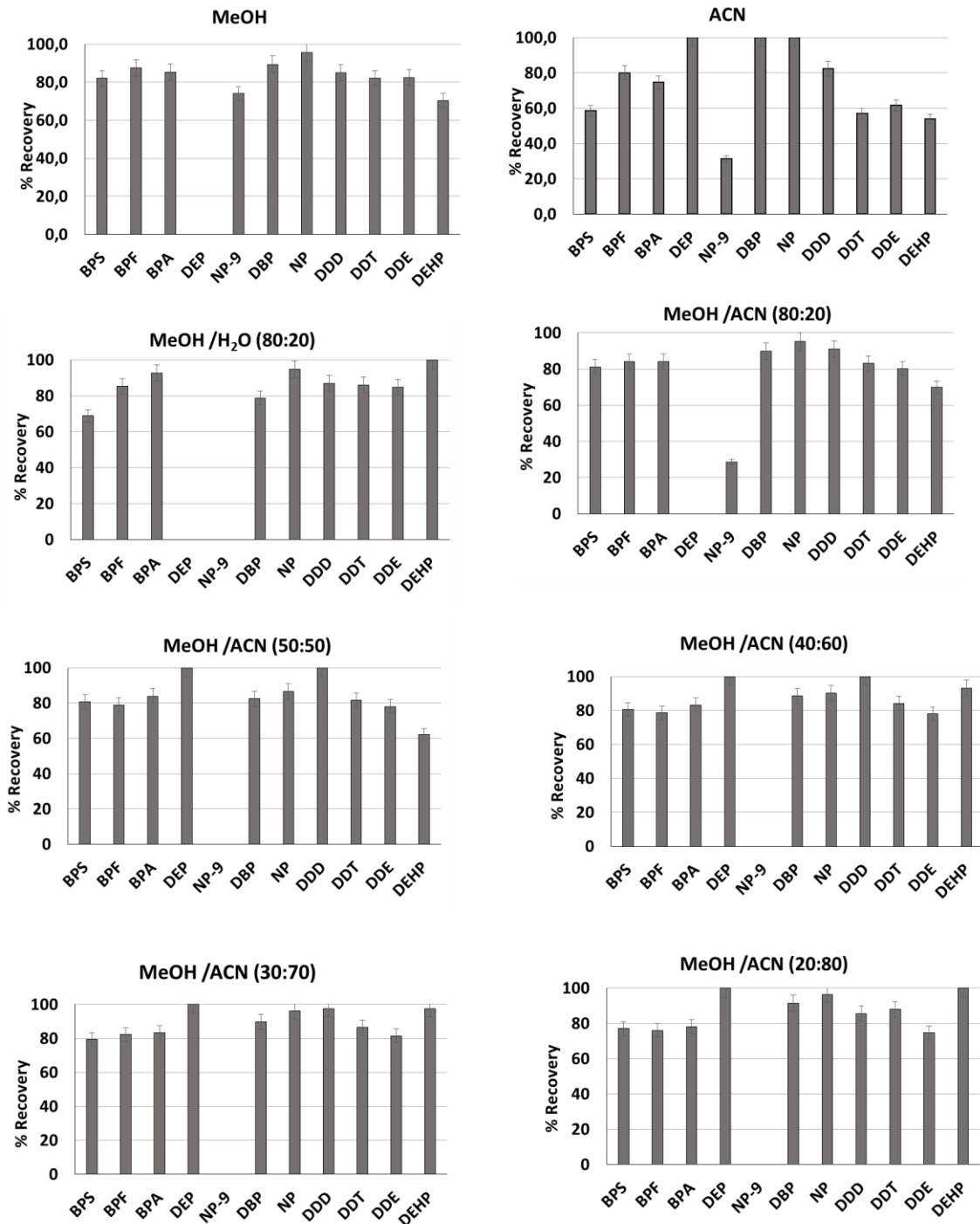


Fig. 2. Elution percentages of target analytes on MSPD columns using 9 mL of different eluents in consecutive 5 min static extraction cycles (3 mL per cycle).

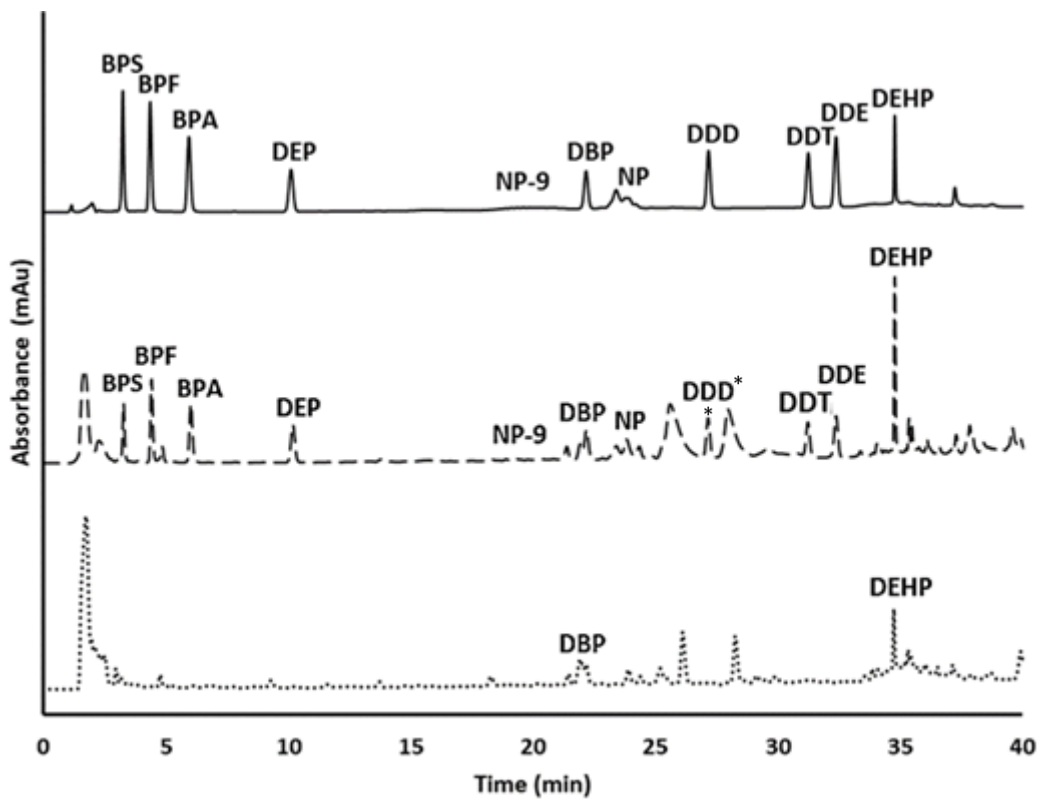


Fig. 3. Chromatograms of a standard solution of 10 mg/L BPS, BPF, BPA, DEP, DBP, DDD, DDT, DDE, and DEHP and 30 mg/L of NP-9 and NP (—), spiked mussel sample with same concentration of compounds (---), and blank mussel sample (...) at 210 nm.

Table 1. Analytical characteristics of target EDCs from spiked mussel samples

Analyte	Concentration range ($\mu\text{g}/\text{kg}$)	Regression equation	R^2	Spiking level ($\mu\text{g}/\text{kg}$)	Intra-day Recovery \pm RSD (%)	Inter-day Recovery \pm RSD (%)	LOD ($\mu\text{g}/\text{kg}$)	LOQ ($\mu\text{g}/\text{kg}$)
BPS	2.00-6.00	$y = 59.792C - 0.9654$	0.9922	40.00	82.6 ± 1.9	82.2 ± 2.0	0.29	1.12
				20.00	81.7 ± 2.5	81.1 ± 3.0		
				2.00	79.3 ± 3.2	78.5 ± 4.1		
BPF	0.80-6.00	$y = 76.526C + 8.712$	0.9953	40.00	82.7 ± 1.8	82.0 ± 0.9	0.10	0.34
				20.00	80.2 ± 2.1	83.5 ± 1.7		
				1.20	83.6 ± 3.5	82.1 ± 3.4		
BPA	0.32-2.00	$y = 110.26C - 3.810$	0.9942	40.00	87.2 ± 1.5	86.2 ± 1.8	0.07	0.25
				20.00	85.3 ± 1.9	87.0 ± 2.0		
				0.40	85.6 ± 3.7	81.5 ± 5.4		
DEP	3.20-6.00	$y = 51.2C - 11.823$	0.9918	40.00	97.8 ± 3.9	96.3 ± 6.4	0.82	2.74
				20.00	104.0 ± 5.1	100.0 ± 5.4		
				3.20	102.3 ± 6.9	101.1 ± 6.6		
DBP	6.00-32.00	$y = 42.235C - 64.875$	0.9913	40.00	89.7 ± 2.9	90.7 ± 2.7	1.65	5.52
				20.00	90.1 ± 3.5	91.4 ± 3.8		
				12.00	89.1 ± 4.1	90.5 ± 4.5		
NP	28.00-60.00	$y = 15.586C - 70.458$	0.9873	120.00	96.9 ± 1.5	97.3 ± 1.8	6.00	16.22
				60.00	97.5 ± 3.9	98.6 ± 3.7		
				28.00	96.9 ± 4.5	96.8 ± 4.7		
DDD	20.00-60.00	$y = 70.598C + 325.94$	0.9982	40.00	98.0 ± 1.2	101.0 ± 1.4	2.45	10.23
				28.00	98.0 ± 2.3	99.5 ± 2.8		
				20.00	97.9 ± 4.1	102.0 ± 3.9		
DDT	0.60-6.00	$y = 76.956C + 10.985$	0.9958	40.00	86.2 ± 0.8	86.5 ± 1.1	0.18	0.43
				20.00	86.0 ± 1.3	85.9 ± 3.5		
				2.00	86.8 ± 2.3	87.1 ± 5.1		
DDE	0.40-6.00	$y = 92.893C + 2.8236$	0.9973	40.00	83.5 ± 2.0	82.8 ± 1.2	0.06	0.25
				20.00	82.5 ± 3.5	82.6 ± 2.3		
				2.00	85.3 ± 4.2	83.9 ± 5.6		
DEHP	2.00-12.00	$y = 105.721C + 32.712$	0.9991	40.00	102.0 ± 1.5	101.4 ± 1.3	0.42	1.87
				20.00	102.3 ± 2.3	$96.7.0 \pm 4.8$		
				2.00	98.7 ± 4.8	105.3 ± 5.2		

Table 1

Analyte	Concentration range ($\mu\text{g}/\text{kg}$)	Regression equation	R^2	Spiking level ($\mu\text{g}/\text{kg}$)	Intra-day Recovery \pm RSD (%)	Inter-day Recovery \pm RSD (%)	LOD ($\mu\text{g}/\text{kg}$)	LOQ ($\mu\text{g}/\text{kg}$)
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				2.00	79.3 ± 3.2	78.5 ± 4.1		
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				20.00	102.3 ± 2.3	$96.7.0 \pm 4.8$		
				2.00	98.7 ± 4.8	105.3 ± 5.2		

Fig. 1

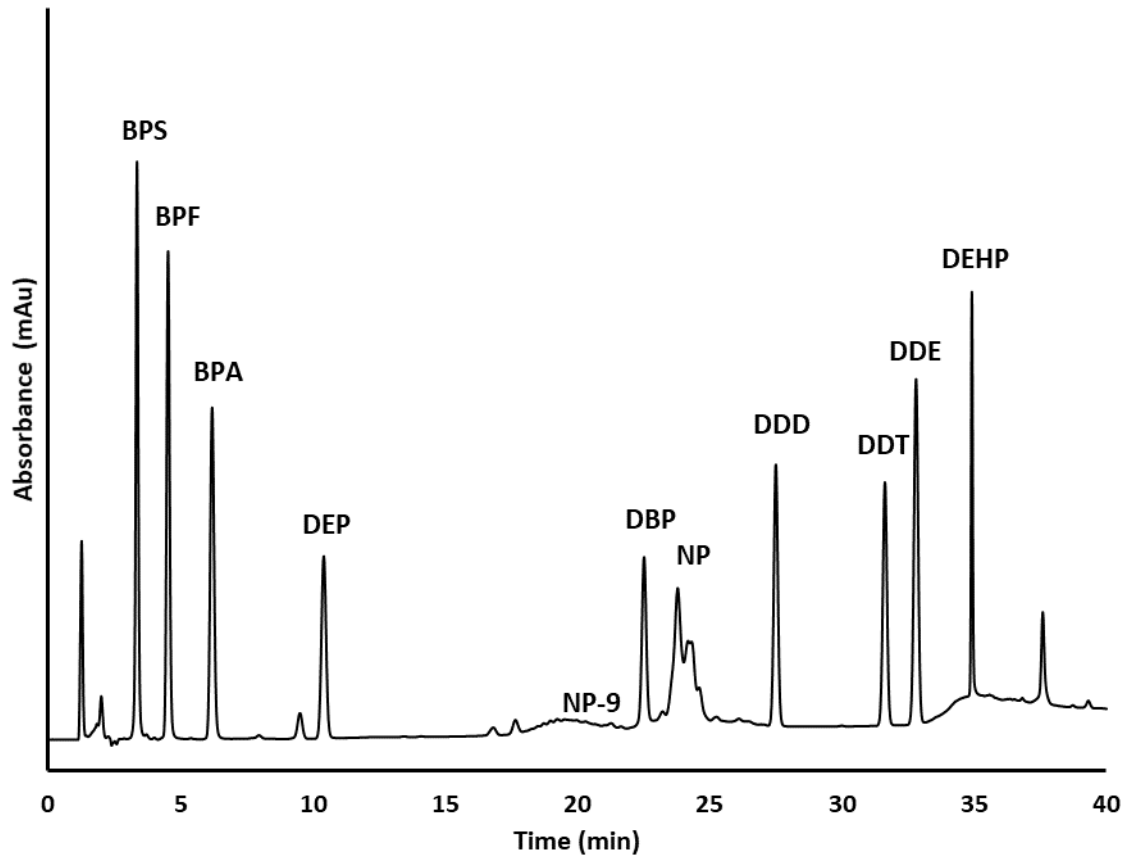


Figure 2

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Fig.2

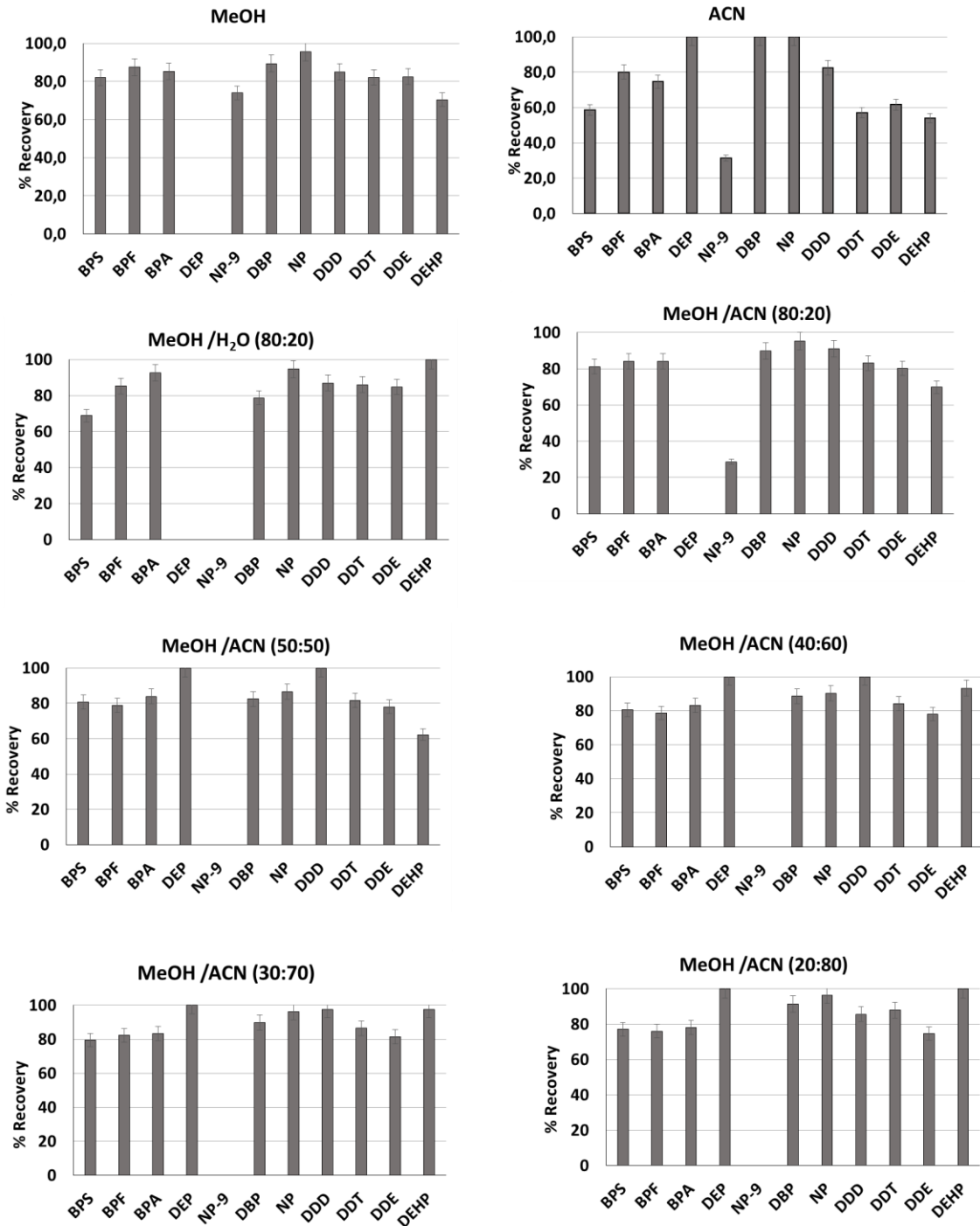
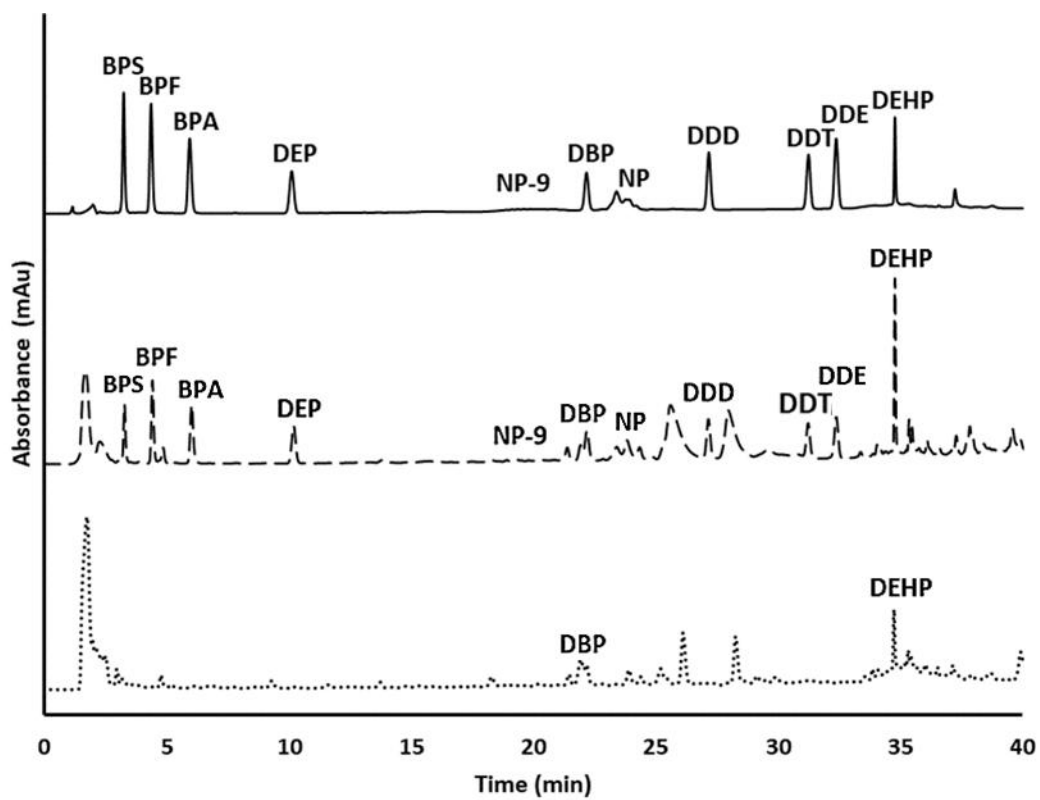


Fig. 3



CRedit author statement

R. Cañadas: Methodology, data acquisition, writing- original draft preparation

E. Garrido Gamarro: Conceptualization, methodology, reviewing and editing

R. M. Garcinuño Martínez: Methodology, validation, reviewing and editing, funding acquisition

G. Paniagua González: Resources, validation, writing- original draft preparation

P. Fernández Hernando: Supervision, reviewing and editing, funding acquisition