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Title: Optimization of hollow-fiber liquid phase microextraction for polychlorinated biphenyls in human breast milk

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Abstract: A reliable and sensitive analytical approach has been optimized for the extraction of seven polychlorinated biphenyls (PCBs) from human breast milk. Hollow fiber liquid phase microextraction (HF-LPME) was applied for the first time for the extraction and pre-concentration of the analytes. Analytes were separated by gas chromatography with electron capture detector (GC- μ ECD) for the sensitive detection and mass spectrometry for the unequivocal identification. A rotatable central composite design (RCCD) was performed for the multivariate optimization of the method. The best results were obtained at 40°C during 30 min and 600 rpm of stirring speed using a hollow fiber length of 5 cm and toluene as an extractant phase and salt addition was not required. The detection limits were in the range 7-14 ng L⁻¹ for PCBs. The coefficients of determination of the calibration curves indicated good linearity ($R^2 > 0.96$) and the enrichment factors ranged from 74-143. This type of study is of great importance due to the deleterious effect that the presence of contaminants can produce in infants health related to the immature character of the defense system. Moreover, exclusive breastfeeding is recommended by neonatologists up to six months of life and as complementary food during the first two years.

Dear Editor,

Please find enclosed the amended version of the paper “Optimization of hollow-fiber liquid phase microextraction for polychlorinated biphenyls in human breast milk” by M.C. Villegas-Álvarez, B. Callejón-Leblic, G. Rodríguez-Moro, A. Arias-Borrego and T. García-Barrera, to be considered for publication in Journal of Chromatography A. The manuscript has been considerably improved taking into account all the points raised by the reviewers. A detailed point-by-point response to each comment is also attached.

The present paper describes for the first time the application of hollow fiber liquid phase microextraction (HF-LPME) for polychlorinated biphenyls (PCBs) in human breast milk. A new method has been optimized using multivariate approaches. The new method is very sensitive and reliable in comparison with other proposed in the literature.

Sincerely yours,

Tamara García Barrera

Highlights

- A HF-LPME method was developed for the determination of PCBs in human breast milk.
- The HF-LPME method is very simple since a sample preparation step is not required.
- A multivariate design (RCCD) was applied to optimize the extraction conditions.
- Extraction time, temperature and length of the fiber were main experimental factors.
- The proposed approach provides limits of detection in the range of 7-14 ngL⁻¹

1 **Optimization of hollow-fiber liquid phase**
2 **microextraction for polychlorinated biphenyls in**
3 **human breast milk**

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16

17 **ABSTRACT**

18 **A reliable and sensitive** analytical approach has been optimized for the **extraction** of seven
19 polychlorinated biphenyls (PCBs) **from** human breast milk. **Hollow fiber liquid phase**
20 **microextraction (HF-LPME)** was applied for the first time for the extraction and pre-
21 concentration of the analytes. Analytes were separated by gas chromatography **with** electron
22 capture detector (GC- μ ECD) for the sensitive detection and mass spectrometry for the
23 unequivocal identification. A rotatable central composite design (RCCD) was **performed** for the
24 multivariate optimization of the method. The best results were obtained at 40°C during 30 min
25 and 600 rpm of stirring **speed using** a hollow fiber length of 5 cm and toluene as an **extractant**
26 **phase and salt addition was not required. The detection limits were in the range 7-14 ng L⁻¹ for**
27 **PCBs. The coefficients of determination of the calibration curves indicated good linearity ($R^2 >$**
28 **0.96)** and the enrichment factors **ranged from 74-143**. This type of study is of great importance
29 due to the deleterious effect that the presence of contaminants can produce in infants health
30 related to the immature character of the defense system. Moreover, **exclusive breastfeeding is**
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32 first two years.

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34 **Keywords:** Hollow Fiber Liquid Phase Microextraction, breast milk, Polychlorinated
35 Biphenyls, multivariate optimization, gas chromatography.

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38 1. Introduction

39 Polychlorinated biphenyls (PCBs) are environmental pollutants with deleterious effects for
40 humans, characterized by a high persistence and bioaccumulation capacity [1–3]. They are a
41 class of persistent organic pollutants (POPs), considered a major public concern [4]. The
42 existing 209 congeners contain from one to eight atoms of chlorine in different positions, which
43 belong to poly-, mono- and non-ortho substituted PCBs. They are semivolatile, hydrophobic and
44 have the basic unit of biphenyl, being structurally similar to thyroid hormones, which in turn led
45 to thyroid disruption [5].

46 They were widely produced between 1930 and 1985 from industry, where they were used as
47 plasticizers, insulating fluids in electric instruments and as sealants additives [6]. As a rule, the
48 main source of PCBs are industries [7], leakage from landfills, incomplete incineration,
49 inappropriate disposal, volatilization [7,8] and port activities [9,10]. They are considered as
50 probable human carcinogens [2] by the Environmental Protection Agency (EPA) and as Group I
51 human carcinogens by the International Agency for Research on Cancer (IARC) classified them
52 [11]. During the prenatal period, exposure to PCBs comes from placental transfer and human
53 breast milk (HBM) [12] causing health problems such as low birth weight [13], reduced head
54 circumference [14], neurological effects [15] and thyroid disruption [5]. For this reason, several
55 studies have suggested the use of breast milk as an indicator of the level of POPs in humans
56 [16,17]. The content of lipids in breast milk is very high ranging from 3.5-4.5 per 100 g (mature
57 milk) and due to the lipophilic character of polychlorinated biphenyls, they are accumulated and
58 transferred to infants during breastfeeding [18].

59 For this reason, analytical methods for the sensitive and selective extraction, pre-concentration
60 and determination of PCBs in this biological fluid are claimed. To this end, different analytical
61 methods have been described for the extraction of PCBs from breast milk to be later analyzed
62 by gas chromatography, like liquid-liquid extraction (LLE) [19], solid-phase microextraction
63 (SPME) [20], accelerated solvent extraction (ASE) [21] and QuEChERS [16,18,22].
64 However, although nowadays hollow-fiber liquid phase microextraction (HP-LPME) is a very
65 promising analytical technique, there is not any HP-LPME method described in the literature for
66 the extraction of PCBs from breast milk. The most important advantages of HP-LPME are
67 among others, the simplicity of operation and required equipment [23], low organic solvent
68 consumption, robustness, low cost, disposable character (avoiding memory effects). The
69 technique is based on the use of a supported liquid membrane containing the extractant phase
70 and separated to the sample phase, which led to high surface area for the extraction and high
71 enrichment factors [24].

72 After the extraction from breast milk, PCBs are usually determined by gas chromatography
73 (GC) with mass spectrometry (MS) and/or electron capture detection (ECD) [16,18,22,25]. The
74 sensitivity of ECD for halogenated compounds is usually better than MS detector, although this
75 last detector allows the unequivocal identification of the compounds by mass spectra databases
76 and tandem mass spectrometry [26].

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78 In this paper, a sensitive and reliable HF-LPME method has been optimized and applied for the
79 first time to the extraction of PCBs from human breast milk. The extracts have been analyzed by
80 both GC-MS and GC- μ ECD for the determination of seven PCBs in breast milk. A multivariate
81 optimization of the variables that affect the extraction was based on the combination and
82 maximization of the response of all the PCBs. This optimization was performed using
83 chemometric tools obtaining high enrichment factors. The developed method was sensitive,
84 simple, reliable, reproducible and low-cost being easily transferable to routine analysis in
85 laboratories to control the occurrence of PCBs in breast milk.

86 2. Experimental

87 2.1. Chemical and standard solution preparation

88 The mixture of PCBs standards (PCB# 1, 5, 29, 47, 116, 136,187,194, 206) were supplied by
89 Sigma–Aldrich (Steinheim, Germany). Ultrapure water (18.2 M Ω cm) was obtained using a
90 Millipore Milli-Q® Direct 8 equipment (Watford, UK). A solution with all the PCBs was
91 prepared by dissolving in n-hexane the corresponding amount to attain the following
92 concentrations of the standards (mg L⁻¹): PCB1 (100), PCB5 (100), PCB29 (10), PCB47 (10),
93 PCB116 (10), PCB 136 (10) and PCB 187 (5). Intermediate working solutions used in direct
94 injection were daily prepared by stepwise dilution of the above described solution with n-
95 hexane. All the standard solutions were stored in darkness at 4°C until analysis. All reagents and
96 solvents used in this study were of the highest purity. N-hexane, acetone and toluene were
97 supplied by Teknokroma (Barcelona, Spain). The hollow fiber membranes (Accurel Q3/2) were
98 made of polypropylene (inner diameter 600 μ m, wall thickness 200 μ m, pore size 0.2 μ m)
99 (Wuppertal, Germany).

100

101 2.2. Samples

102 Human breast milk samples were taken after 48 h of life and stored at -80°C until their analysis.
103 Samples were collected at the Juan Ramón Jiménez Hospital, during 2017. Written informed
104 consent was obtained from all the participants and the study was approved by a local ethical
105 committee.

106 For analytical validation of the method, the sample phase was composed by 350 μ L of human
107 breast milk to a final volume of 7 mL with ultrapure water (18.2 M Ω cm).

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2.3. Instrumentation

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2.4. Extraction of PCBs by HF-LPME

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Extracts were analyzed into a gas chromatograph model 6890N coupled to a ^{63}Ni microelectron capture detector (GC- μECD) (Agilent Technologies, Hewlett Packard, Wilmington, DE, USA). Chromatographic separation was conducted on a DB-5 chromatographic column (30m x 0.25mm x 0.25 μm). The carrier gas was helium (99.999%, 1.2 mL min^{-1}). A volume of 1 μl of the extracts was injected into the GC using the splitless mode (splitless time: 30 s). The injector temperature was set at 250°C. The oven temperature was programmed at 75°C for 1 min, then increased to 150°C ramped to 30°C min^{-1} , to 180°C with a ramp to 5°C min^{-1} held for 15 min, to 190°C ramped to 5°C min^{-1} for 5 min, to 200°C with a ramp of 5°C for 3min, to 280°C with a ramp of 5°C held for 2.5 min and then raised by 50°C min^{-1} to 290°C with 4.8 min hold time. Nitrogen was used as makeup gas for the ECD detector (60 mL min^{-1} , 99.999%). The ChemStation software package (version A0903) was used for data acquisition. The extracts were simultaneously analyzed on an ion-trap GC-MS model ITQ 900 (Thermo Fisher Scientific, Thermo Fisher Scientific Spa, Rhone, Milan, Italy). The analytical column and oven temperature conditions were same than that used for GC- μECD . The injector temperature was set at 250 ° C. The carrier gas was helium (1.3 mL min^{-1}). The temperature of the transfer line between the chromatograph and the mass detector was set at 280 °C. For the separation of the peaks, the temperature program of the oven was the same above described for the GC- μECD . The ionization of analytes in the mass spectrometer was carried out by electronic impact (EI) with electronic ionization source at 70 eV in selected ion monitoring mode (SIM). PCBs were identified by retention time matching with standards, NIST database and tandem mass spectrometry.

The extraction procedures were optimized prior to the application. The U-shaped configuration of the fiber and two-phase mode were used for HF-LPME. The arrangement is shown in Figure 1. The fibers were cut into 5cm portions and then cleaned into an ultrasound bath with acetone during 10 min. Then the cleaned and air-dried fiber was inserted into one of the medical syringe needles. The extractant phase toluene (500 μL) was loaded into the lumen of the membrane by the immersion of the fiber into a vial containing that solvent, avoiding incomplete filling and air bubbles. The second end of the fiber was inserted into other syringe needle. Both syringes were introduced through of the septum into a vial containing toluene during 1 min in order to fill the pores of the membrane with the extractant phase (SLM). Afterwards, the U-shaped fiber was introduced into a 10 mL glass vial with 7 mL of sample (sample phase). The extraction was carried out in direct immersion mode during 40 min at 36°C and 600 rpm. After that, the extract

145 was collected from the lumen and poured into a 1.5 mL vial with an insert of 250 μ L by
146 separating one end of the fiber from the syringe. Finally, 10 μ L were obtained from the fiber
147 and 1 μ L was injected into GC- μ ECD/MS.

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149 **2.5. Optimization of hollow-fiber liquid phase microextraction**

150 The most influential variables in HF-LPME were optimized for the simultaneous analysis of 7
151 PCBs in human breast milk. Firstly, a preliminary screening using a univariate optimization was
152 carried out to explore the influence of the different variables as well as the range of values that
153 should be optimized. The following variables were studied in this work: extraction solvent,
154 temperature, stirring rate, ionic strength, pH, fiber length and extraction time. After that, a
155 Rotatable Central Composite Design (RCCD) of second order was used for the optimization of
156 the variables taking into account the interactions between them. To this end, the data were
157 processed with the MINITAB®Release 16 Statistical Software (State College, USA). The
158 combined response was also optimized using some home-made programs written in
159 QUICKBASIC.

160

161 **3. Results and discussion**

162 **3.1 Screening of the critical variables in the extraction process**

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164 A univariate approach was applied to screen the main factors with significant effects on the HF-
165 LPME extraction procedure. To this end, when one of the variables is modified, the other
166 features were set at the following values: temperature (25°C), extraction time (5 min), HF length
167 (4 cm), immersion time (1min), pH (not adjusted) and sample volume (350 μ L). The screened
168 factors have been matching with those selected in most of the previous works about the HF-
169 LPME extraction process of organic compounds [27,28]. The changes in the peak areas of the
170 PCBs were plotted against the variation of each factor optimized and analyzed to select the
171 ranges for the multivariate optimization.

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173 **3.1.1. Selection of the extractant phase for the HF-LPME**

174

175 Several considerations should be taken into account to select the solvent for the extractant phase
176 to be used for the HF-LPME: (i) it should be compatible with the material of the fiber
177 (polypropylene) and (ii) the solvent should be immiscible with water to form two phases for the
178 extraction, (iii) the viscosity should be adequate to allow filling the membrane pores as well as
179 the diffusion of analytes, (iv) low volatility to avoid losses during the extraction procedure [29].

180 Under these conditions, three organic solvents were checked to extract the target analytes (1-
181 octanol, toluene and n-hexane). Figure 2 shows the enrichment factors for all the analytes using
182 these solvents concluding that toluene is the most efficient **extractant** phase for the extraction.
183 **Longer times than 1 min to embed the pores of the HF do not enhance the enrichment factors.**

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185 **3.1.2. Effect of salt addition**

186 The ionic strength could influence the partition coefficients of the analytes when two
187 immiscible phases are in contact, which in turn affects the extraction efficiency. The ionic
188 strength could be increased by adding salt, **which could enhance** the enrichment factors of the
189 analytes in the organic solvent [27,28,30–34]. To this end, different percentages of sodium
190 chloride were assayed up to 10% (w/v), but any effect was detected. The reason maybe could be
191 the low polar character of the PCBs compounds, which enhance the extraction and does not
192 need the modification of the ionic strength [35]. This result is in agreement with our previous
193 work [27].

194

195 **3.1.3. Effect of agitation**

196 The stirring **speed** is important to enhance the mass transfer of the analytes from the **sample**
197 **phase** to the **extractant** phase, reducing also the extraction time. **Thus, the stirring rate** was
198 increased up to 1200 rpm, and the results showed a progressive increase of the enrichment
199 factors until 600 rpm (Figure 3). However, higher values of this variables decrease the
200 reproducibility of the results, mainly due to the **occurrence** of air bubbles in the solution. Based
201 on these findings, 600 rpm was **used in** further experiments.

202

203 **3.1.4. Effect of pH**

204 **The effect of pH in the extraction of PCBs by HF-LPME was optimized in the range of 4-12**
205 **and the corresponding enrichment factors are shown figure 4. Acid conditions were achieved by**
206 **adding 6M HCl and basic conditions with 1M sodium hydroxide. There was no considerable**
207 **effect in the enrichment factors of the PCBs varying the pH in the range of 6-8, in which the**
208 **sample pH is included. Thus, the sample pH was not adjusted for subsequent experiments.**

209

210 **3.3. Central composite design**

211 A RCCD was created [28] to optimize the three most important variables: **temperature (T), time**
212 **(tm) and membrane length (L)**. The number of experiments of the full factorial design can be
213 calculated by 2^k , where k is the number of variables. These eight experiments can be augmented
214 with 2k star points, which are the vertices of a cross-polytope **at** the coordinate axes of the
215 **design**. Finally, C **central** points are included as replicates. Thus, for three variables to optimize,
216 the CCD consist of 17 experiments ($2^k + 2k + C$). **The variables to be optimized has** five coded

217 levels $(-\alpha, -1, 0, 1, \alpha)$, where $\alpha = 2^{k/4} (\sim 1.68)$, the **star arm (distance) from the center to a star**
 218 **point**. The $-\alpha$ (**minimum**) and $+\alpha$ (**maximum**) **define the range of parameters to be optimized**.
 219 To avoid systematic errors the trials were randomized. Table 1 collect the trials included **in the**
 220 **RCCD** showing the real and coded levels. Table 2 shows the response (areas peak of the
 221 analytes) of the PCBs obtained by HF-LPME in the different trials for the optimization of the
 222 temperature, extraction time and membrane length. After that, a regression analysis was
 223 performed to obtain reduced models of polynomial equations to calculate the response of the
 224 analytes using the studied variables (Table 3). A significant Student's "t-test" was performed to
 225 obtain the regression coefficients $b_k, s(b_k)$. An effect is considered significant when $t_k =$
 226 $|b_k|/s(b_k)$ is greater than the tabulated $t_{critic, (v,P)}$, **with** v degrees of freedom ($v =$ number of run
 227 $-$ number of coefficients to be **calculated**), **at a P confidence level of 95%** ($p\text{-value} > 0.05$).

228 **Table 1. Experiments performed in the design of the RCCD.**

Run	Time min (X1)		Temperature °C (X2)		Length cm (X3)	
	Coded level	Real level	Coded level	Real level	Coded level	Real level
1	-1	20	-1	35	-1	4
2	1	40	-1	35	-1	4
3	-1	20	1	45	-1	4
4	1	40	1	45	-1	4
5	-1	20	-1	35	1	6
6	1	40	-1	35	1	6
7	-1	20	1	45	1	6
8	1	40	1	45	1	6
9	0	30	0	40	0	5
10	0	30	0	40	0	5
11	-1.68	13.2	0	40	0	5
12	1.68	46.8	0	40	0	5
13	0	30	-1.68	31.6	0	5
14	0	30	1.68	48.4	0	5
15	0	30	0	40	-1.68	3.32
16	0	30	0	40	1.68	6.68
17	0	30	0	40	0	5

229

230 **Table 2. Peak areas of the analytes after the analysis by HF-LPME from the RCCD trials.**

Run	PCB 1	PCB 5	PCB 29	PCB 47	PBC 116	PCB 136	PCB 187
1	1445437	151166	341421	2982558	992053	576920	1079141
2	1224615	315737	459125	2072237	2018372	1072200	1138348
3	2532024	978107	290671	144663	445166	1094476	684602
4	1514127	106678	381923	151113	739466	986393	670114
5	2090428	13532203	1635603	803381	471934	2719882	241383

6	1787659	18015125	1041421	873296	823271	2091914	385883
7	1981037	17599705	395431	754873	774625	1484487	533688
8	215935	10079523	249864	784821	1092053	1363427	421684
9	32821417	8971327	1635202	1906952	14896527	1270088	565888
10	47586625	9125733	1444278	1706678	13739257	1592765	542409
11	8670406	338390	810645	229711	236551	1098930	394146
12	10514127	13151374	1637315	1893551	21003809	1177669	498725
13	4670909	11119787	1980343	1935806	15188631	1962856	569321
14	5709354	1298494	455265	617351	749486	669072	431516
15	996262	145265	1044196	287919	1859647	645741	887167
16	8670406	15131803	753395	436889	1941386	1285688	565888
17	4670909	9599705	1867114	1383234	1305880	1221292	581988

231

232 As we can see in Table 3, the temperature (T) and the HF length (L) have the most significant
 233 effects in the response of all the target analytes (the variables **with** squared terms indicate a high
 234 effect in the response of the analyte). The extraction time also has a significant effect in the
 235 response of PCB116, as can be seen in its equation model. In **almost all the trials**, the interaction
 236 between the temperature (T) and HF length (L) **enhance** the peak areas of the **PCBs**.

237 **Table 3. Mathematical models of responses and variables.**

Model equation
PCB1=285974-900493T ² -913135L ²
PCB5=917345+607208L
PCB29=168089-346052T-372945L ²
PCB47=165103-521151T+577754T*L+412916L ²
PCB116=103614+27027tm-186973T-65655L
PCB136=134154-271583T+366794L-299454T*L
PCB187=557634-185368L+148860T*L+77180L ²

Equation model: $y=B_0+\sum B_{0i}X_i+\sum B_{ii}X_i^2+\sum B_{ij}X_iX_j$

Response (peak area)

Variables: *T* (temperature), *tm* (extraction time), *L* (HF length)

238

239 In order to have the compromise of the best response of each analyte according to the variables
 240 used, **the responses of all the analytes were combined for a better optimization**. In this way, the
 241 analytical operations are simplified and avoided possible rounding mistakes and without
 242 neglecting any term. The response variable, namely “Combined Response” is **the** weighed sum
 243 of **all the** responses using the equation (Eq.1):

$$COMB = \sum_i \left\{ kw_i \left| \frac{Y_i - G_i}{R_i} \right| \right\}$$

244 Where w_i is the weight of the variable, which can be chosen by the user (generally the unity), Y_i
245 are the different responses for the optimization and R_i their range. When the aim is the
246 maximization of the combined response, then $G_i = \text{minimum}$ in the range $(Y_i - 1/2R_i)$ and $K=1$.
247 The randomized optimization of the combined surface (tm, L, T) is carried out using a home-
248 made algorithm and the simultaneous optimum is the maximum of the COMB surface. Several
249 previous studies have used this optimization method with very good results [27,36]. Figure 5
250 shows the 3D combined response and contour plots against the time, length and temperature as
251 well as the optimum point. The optimum coded values were fixed at the following coordinates:
252 temperature (T)=0.00, extraction time (tm)=0.00, and HF length (L)=0.00, which correspond to
253 the decoded values of 40°C for the extraction temperature, 30 min for the extraction time and
254 5cm of length.

255

256 3.4. Method validation

257 For method validation, the figures of merit of the optimized HF-LPME GC- μ ECD method were
258 calculated for the analysis of PCBs (Table 4) such as: coefficients of determination (R^2), linear
259 range, limits of detection (LOD) and quantification (LOQ), relative standard deviation (%RSD)
260 and enrichment factors (EF). The experiments were performed using a spiked sample phase
261 (350 μ L of human breast milk diluted with ultrapure water to a final volume of 7 mL). PCBs
262 were under the detection limits in the samples used for method validation. The linearity was in
263 the range 0.02 to 140 $\mu\text{g L}^{-1}$ for PCB29 and PCB47, 0.05 to 70 $\mu\text{g L}^{-1}$ for PCB187, and in the
264 range of 0.03 to 140 $\mu\text{g L}^{-1}$ for other PCBs determined, and the determination coefficients
265 ranged from 0.96 to 0.98. Seven different concentration levels of the PCBs under study were
266 prepared and analyzed to determine the LODs and LOQs. The limits of detection were
267 calculated as the concentration corresponding to a peak area 3-fold higher than the background
268 noise, and the limits of quantification as the concentration corresponding to a peak area 10-fold
269 higher the background noise of the chromatogram ($\text{LOQ} = 10S/N$). The LOD values ranged from
270 8 to 14 ng L^{-1} with the proposed method. These results are very low when compared with other
271 methods previously published in literature Table 5. Under optimized HF-LPME conditions, the
272 enrichment factors (EF), calculated as the ratio between the PCBs concentration in the
273 extractant phase (Ca) and their initial concentration in the sample phase (Cd) [28,37], were
274 satisfactory in the range of 74 to 143-fold that is in good agreement with other published works
275 related with the use of HP-LPME for other food samples [27]. The results have confirmed that
276 the features of the method are very good and the methodology can be successfully applied for
277 the analysis of samples.

278

279 **Table 4. Figures of merit of the HF-LPME/GC- μ ECD approach for the analysis of PCBs**
 280 **in human breast milk.**

	Precursor ion m/z*	Recovery (%)	Repeatability RSD (%) ^a	Reproducibility RSD (%) ^b	EF	Range (ng L ⁻¹)	R ²	LOD (ng L ⁻¹)	LOQ (ng L ⁻¹)
PCB 1	188.08	73	5	7	124	30-1400	0.96	10	33
PCB 5	222.07	70	7	10	168	30-1400	0.98	8	27
PCB 29	256.05	82	6	9	128	20-1400	0.98	7	23
PCB 47	291.97	117	6	8	128	20-1400	0.97	8	27
PCB 116	325.98	97	9	11	144	30-1400	0.98	9	29
PCB 136	359.96	80	7	10	118	30-1400	0.97	10	33
PCB 187	395.88	72	4	7	75	50-700	0.97	14	47

281 ^aIntra-day precision, calculated by extracting n=5; ^bInter-day precision, calculated by extracting n=5;

282 *The compounds are unequivocally identified by GC-MS.

283

284 The optimized method has been applied to human breast milk for the analysis of a group of
 285 seven PCBs. In spite of human breast milk is a sample matrix very complex the proposed
 286 method is very simple because it does not include a sample preparation step, which is mainly
 287 due to the excellent clean-up capability of the HF-LPME technique. In fact, this procedure
 288 involves only an initial dilution of the sample in ultrapure water, followed by two-phase HF-
 289 LPME. To this end, the extraction recoveries (ER) were calculated using the sample phase
 290 fortified at three different concentrations (100 μ g L⁻¹ of PCB1 and PCB5, 5 μ g L⁻¹ of PCB187
 291 and 10 μ g L⁻¹ of other PCBs). Recoveries obtained ranged from 70 to 109 % at those spiked
 292 level, which are in good agreement with other methods proposed in the literature (Table 4 and
 293 5) [18,27,38]. For method precision evaluation, the reproducibility (inter-day) and repeatability
 294 (intra-day) were calculated using the same spiked sample phase than that for the extraction
 295 recoveries. Five replicates were analyzed in the same day for the repeatability and the results
 296 were very good ranging from 4 to 9 % for PCBs, when they are compared with other methods
 297 [18,27,38]. The reproducibility was calculated with five replicated analysis over 3 different
 298 days. The results were very good in the range of 7 to 11 % [18,27,38]. Figure 6 show the
 299 chromatograms obtained from a sample of human breast milk spiked with 10 μ g L⁻¹ of PCBs
 300 before and after the extraction using HF-LPME followed GC- μ ECD. As can be seen, the
 301 enrichment factors are very high, as well as the selectivity obtained after the accurate
 302 optimization of the HF-LPME combined with the high selectivity and sensitivity of the ECD
 303 detector for halogenated compounds.

304

305 **3.5. Comparison of the developed method with current methods**

306 Taking into account the extraction efficiency (enrichment factors) and sensitivity (LODs), the
 307 proposed approach was in good agreement with other methods for the extraction and
 308 preconcentration of the analytes from human breast milk. On the basis of the analytical features

309 collected in Table 4, the **optimized** method revealed that the LODs were very close to **other**
 310 **analytical methods** previously proposed in the literature, **and the repeatability and (%RSD) the**
 311 **reproducibility (%RSD) were lower than those of other reported (Table 4 and 5)**. On the other
 312 hand, it is **remarkable** that the extraction time of the HF-LPME method **is** shorter than those of
 313 other methods. The comparison of the results revealed that the **developed** method is rapid,
 314 **accurate and** simple for the sensitive determination of PCBs in human breast milk samples.

315

316 **Table 5. Comparison of parameters validation of the proposed method with others**
 317 **previously published for the extraction of PCBs in human breast milk.**

Method	SPME [22]			QuEChERS [16,18]			
	LOD (ngL^{-1})	LOQ (ngL^{-1})	Recovery (%)	LOD (ngL^{-1})	Intra-day RSD (%)	Inter-day RSD (%)	Recovery (%)
PCB 1	-	-	-	-	-	-	-
PCB 5	-	-	-	-	-	-	-
PCB 29	1650	5000	83	200	4	11	96
PCB 47	2240	6790	99	200	14	10	102
PCB 116	1840	4330	101	100	9	9	107
PCB 136	1140	3470	101	100	7	10	104
PCB 187	1000	3040	96	50	8	11	101

318

319

320 4. Conclusions

321 **In this work**, a simple, effective and selective HF-LPME method **has been accurately optimized**
 322 **for the analysis** of PCBs from human breast milk, involving low consumption of organic
 323 solvents, high pre-concentration factors of the analytes and relatively low cost. **This is the first**
 324 **time that HF-LPME is applied to breast milk samples for the extraction of PCBs, which are**
 325 **environmental pollutants with important deleterious effects in human health**. A rotatable second-
 326 order central composite multivariate design was **applied to determine the operational variables**
 327 **which allows obtaining maximum peak areas** of seven PCBs in human breast milk. **The**
 328 **proposed method allows obtaining** limits of detection in the range of 7-14 ngL^{-1} with average
 329 recoveries of 85% for the target analytes and enrichment factors very satisfactory **when**
 330 **compared with previous** analytical methods for the analysis of PCBs in breast milk. The
 331 proposed method can be easily automatizable, converted into a green analytical technique and
 332 **transferrable to** routine applications due to its **low cost, sensitivity**, simplicity, robustness and
 333 utility.

334

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342

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- 481
- 482

483 **Legends of Figures**

484 Figure 1. U-shaped HF-LPME configuration for two-phase extraction of PCBs from
485 human breast milk.

486 Figure 2. Mean values (n=3) of the enrichment factors for the analytes using different solvents.

487 Figure 3. Mean values (n=3) of the enrichment factors for the analytes at different stirring
488 speeds.

489 **Figure 4. Mean values (n=3) of the enrichment factors for the analytes at different pH.**

490 **Figure 5.** 3D combined response and contour plots of the surface shape against the
491 temperature, time and length axis as well as the optimum point.

492 **Figure 6.** Chromatograms obtained from a human breast milk sample spiked with $10 \mu\text{g L}^{-1}$ of
493 PCBs before (blue line) and after (red line) the extraction using HF-LPME followed GC- μ ECD.

494

Figure 1
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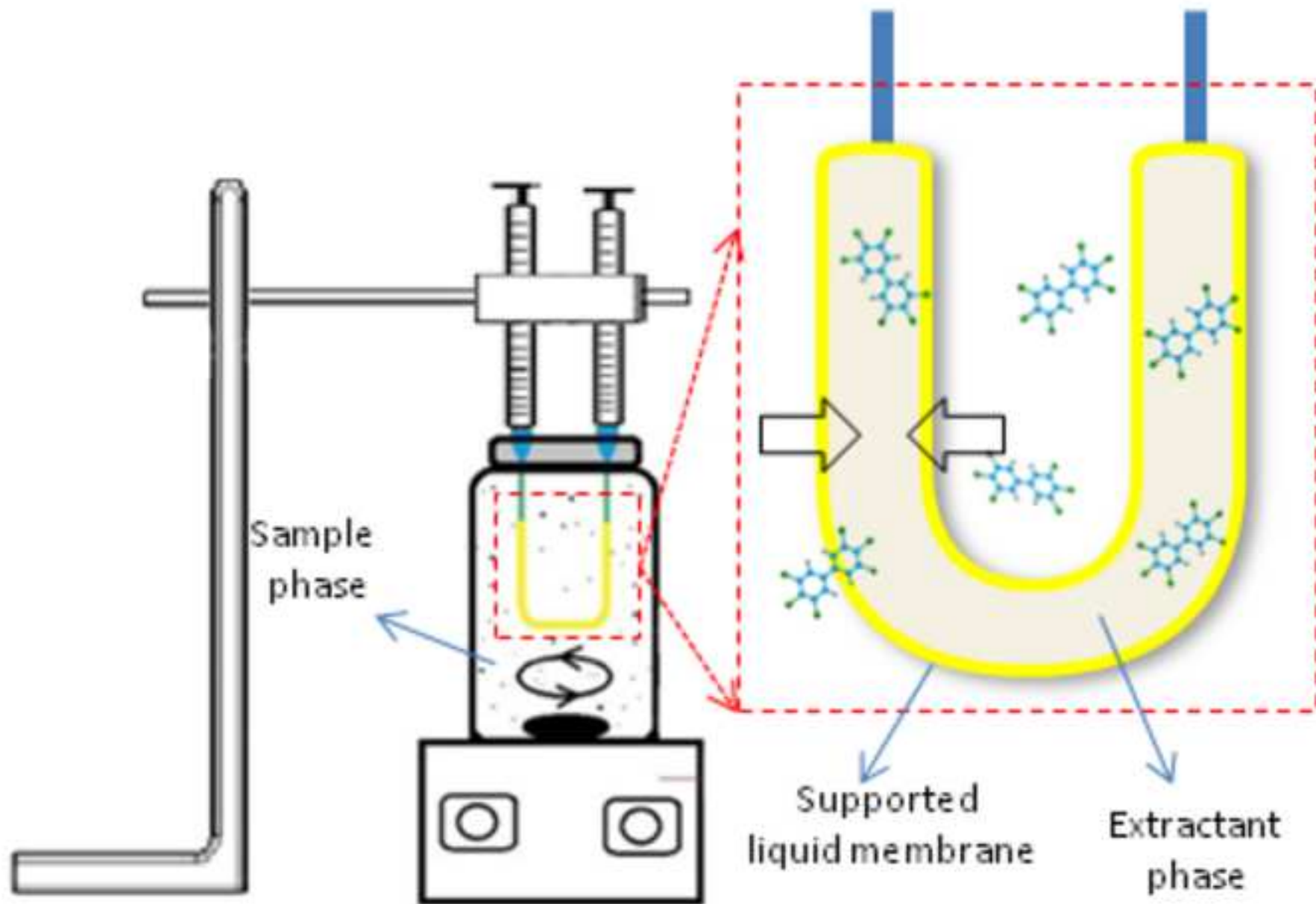


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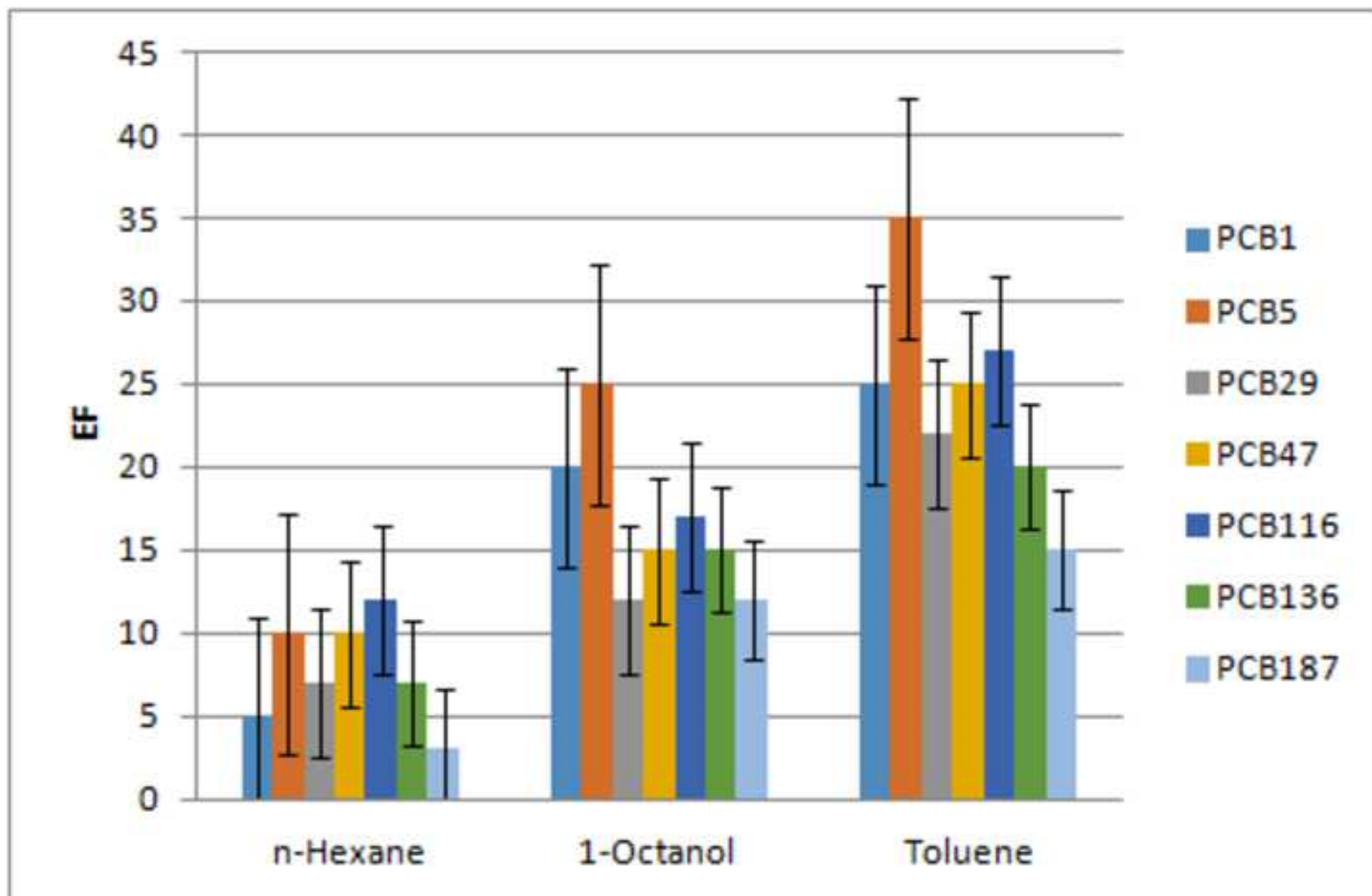


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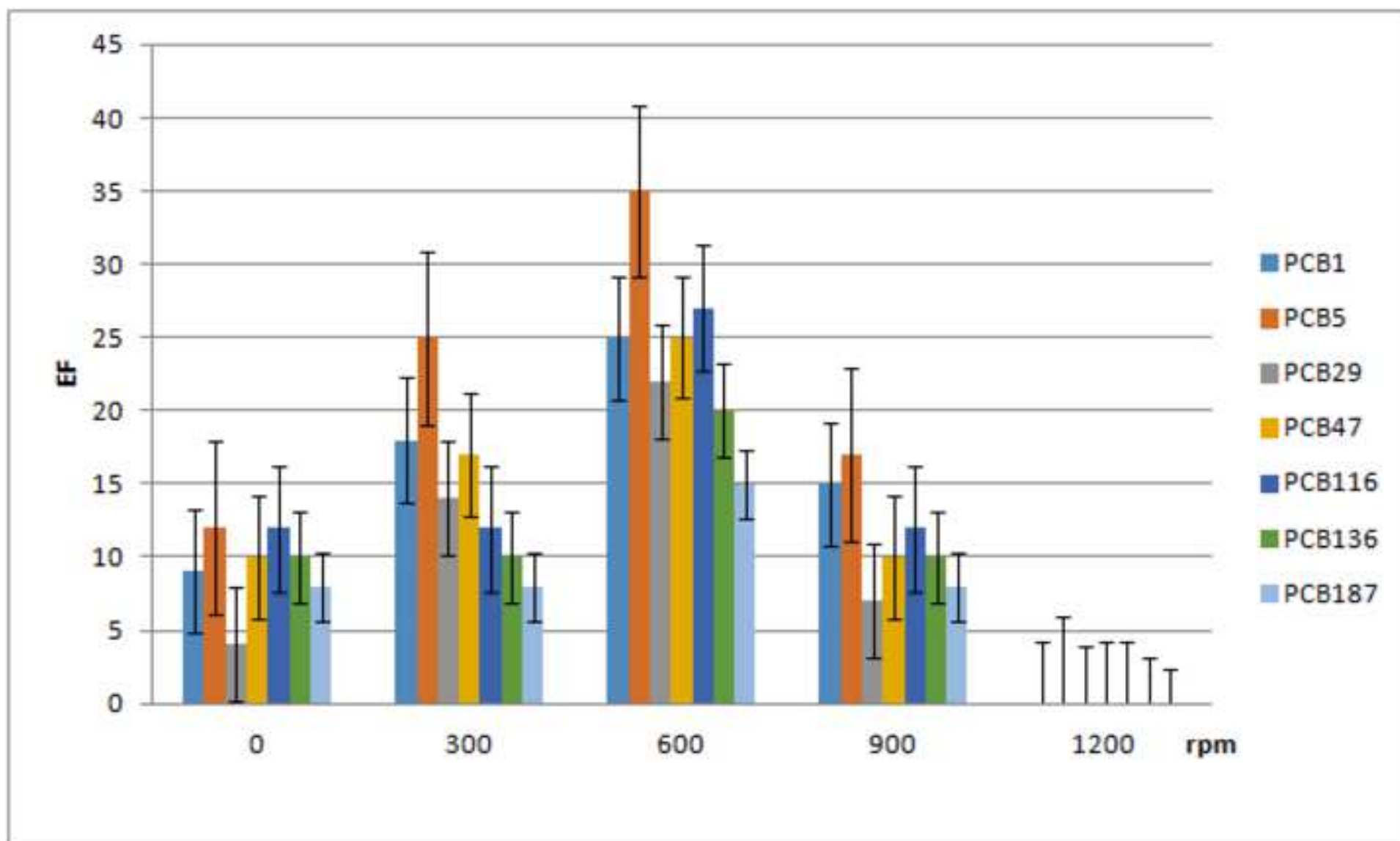


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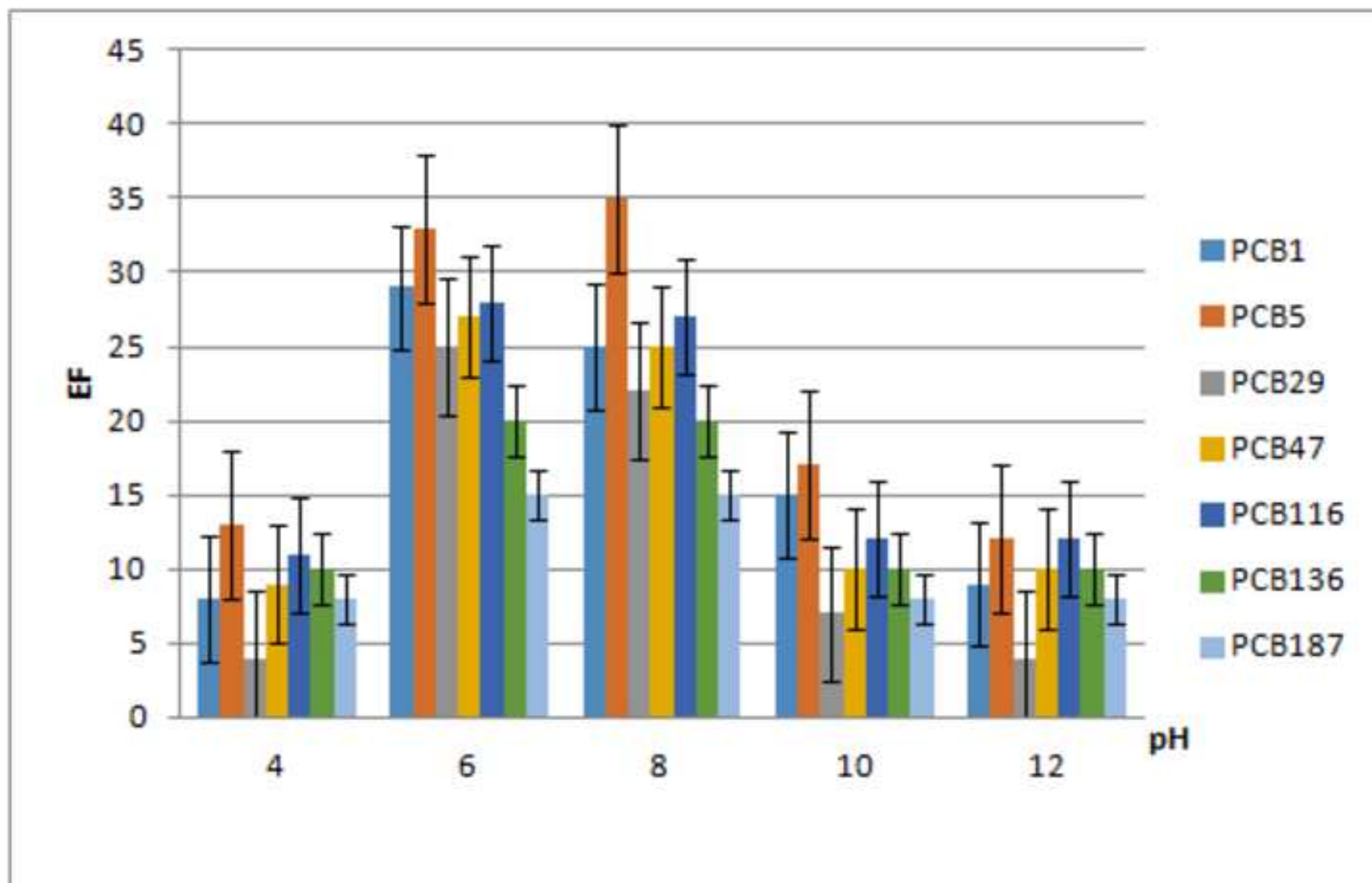


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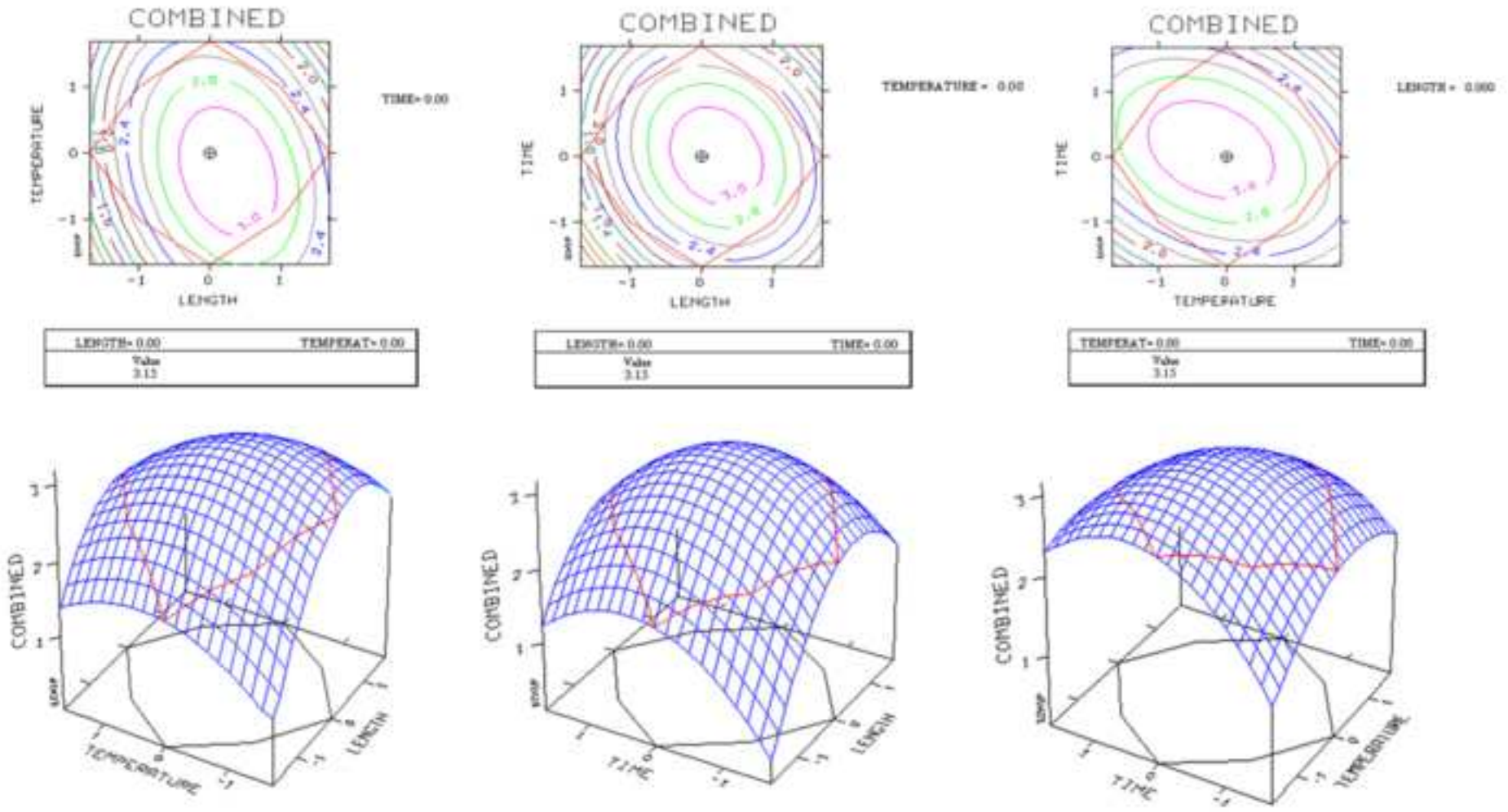
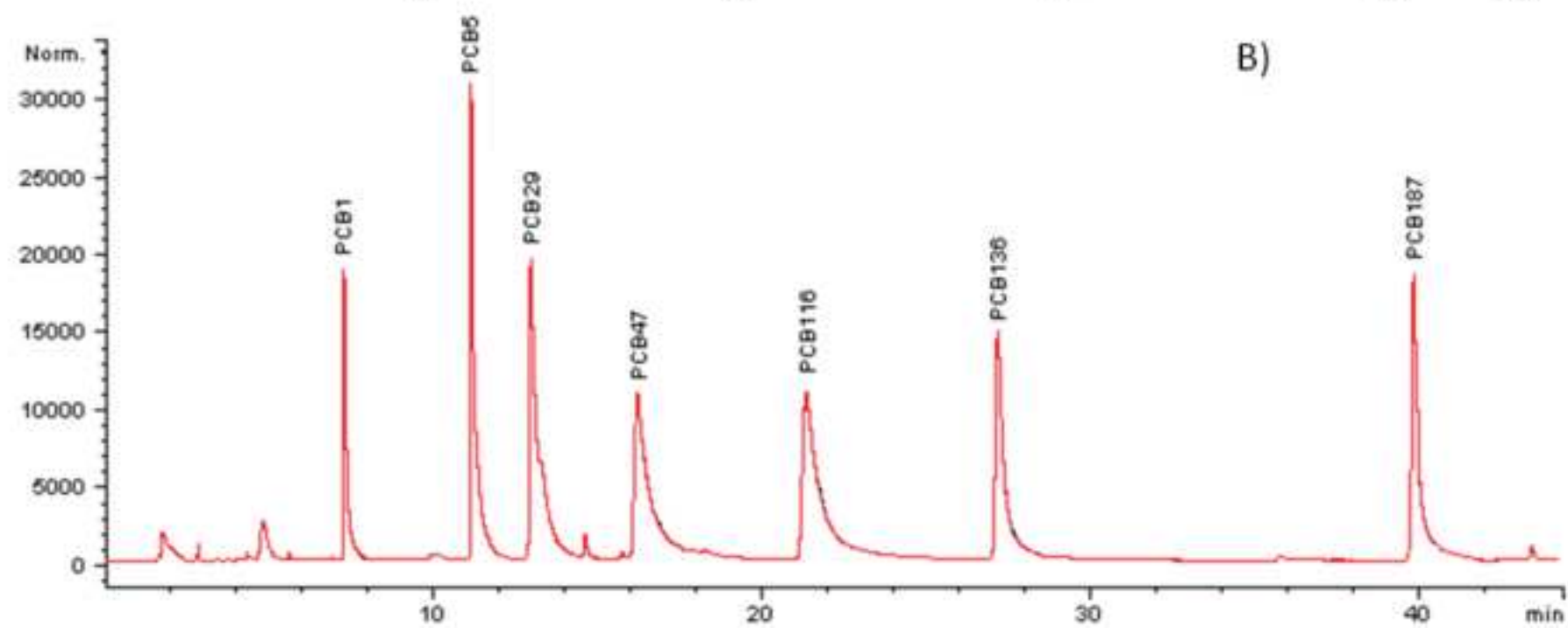
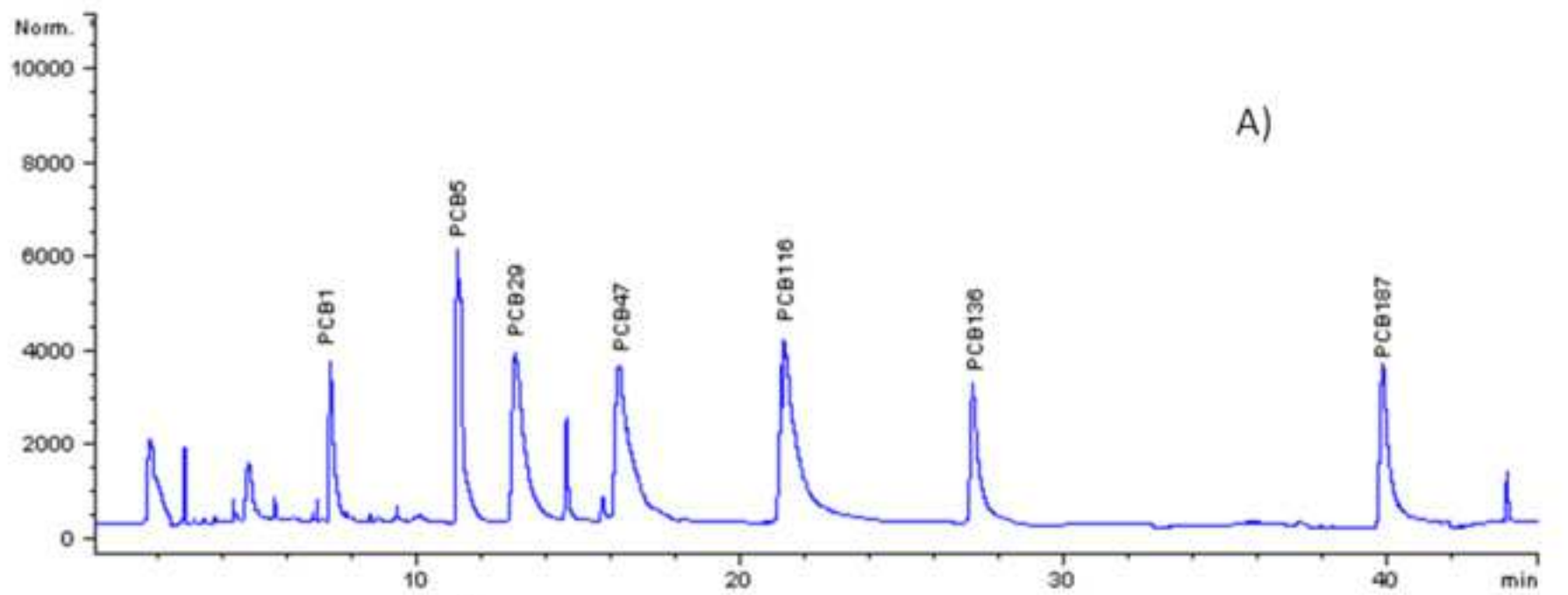


Figure 6
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