

**MULTI-CLASS DETERMINATION OF UNDESIRABLES IN AQUACULTURE
SAMPLES BY GAS CHROMATOGRAPHY/TANDEM MASS
SPECTROMETRY WITH ATMOSPHERIC PRESSURE CHEMICAL
IONIZATION: A NOVEL APPROACH FOR POLYCYCLIC AROMATIC
HYDROCARBONS**

T. Portolés^{1*}, B. Garlito¹, Jaime Nácher-Mestre², M.H.G. Berntssen³, J. Pérez-Sánchez².

¹ Research Institute for Pesticides and Water, University Jaume I, 12071 Castellón, Spain.

² Institute of Aquaculture of Torre la Sal (IATS, CSIC), 12595 Ribera de Cabanes, Castellón, Spain.

³ National Institute of Nutrition and Seafood Research, PO Box 2029 Nordnes, N-5817 Bergen, Norway.

* Phone: 34-964-387339. Email: tportole@uji.es.

ABSTRACT

In this work, a method for the analysis of 24 PAHs in 19 different matrices, including fish tissues, feeds and feed ingredients, has been developed using gas chromatography coupled to triple quadrupole tandem mass spectrometry with atmospheric pressure chemical ionization source (GC-APCI-MS/MS). The method is based on a modification of the unbuffered QuEChERS method, using freezing as an additional clean-up step and applying a 20-fold dilution factor to the final extract. The procedure was also tested for 15 pesticides and 7 polychlorinated biphenyl (PCB) congeners in order to widen the scope of the method.

The excellent sensitivity and selectivity provided by GC-APCI-MS/MS allowed the dilution of the sample extracts and quantification using calibration with standards in solvent for all the 19 matrices tested. The developed method was evaluated at 2, 5 and 50 ng·g⁻¹ spiking levels. LOQs were 2 ng·g⁻¹ for most compounds, and LODs ranged from 0.5 to 2 ng·g⁻¹.

Analysis of real-world samples revealed the presence of naphthalene, fluorene, phenanthrene, fluoranthene and pyrene at concentration levels ranging from 4.8 to 187 ng·g⁻¹. No PCBs, DDTs and pesticides were found in fillets from salmon and sea bream.

KEYWORDS

Polycyclic aromatic hydrocarbons; atmospheric pressure chemical ionization; gas chromatography; triple quadrupole; aquaculture; fish; feed.

INTRODUCTION

Farming of carnivorous marine fish species such as Atlantic salmon (*Salmo salar*) and gilthead sea bream (*Sparus aurata*) has traditionally relied on the use of marine feed ingredients such as fish oil and meal in their formulated feeds [1–3]. Limited access to fish meal and fish oil for the rapidly growing aquaculture industry has led to a need to develop more sustainable aquafeeds that rely less on marine ingredients from feral fish stocks [4,5]. Vegetable oils and proteins are the main alternative to marine feed ingredients in aquafeeds [2,5].

The development of new sustainable plant based feeds for marine fish farming introduces new challenges concerning contaminants that were previously not relevant when using traditional marine feed ingredients. Unrefined plant oils obtained from oilseeds such as soybeans, rapeseeds, olive seeds, and sunflower seeds are known to contain elevated levels of polyaromatic hydrocarbons (PAHs) [6–11]. Although fish oil also contains PAHs from environmental pollution [12], the use of plant oils causes increased PAH levels in plant-based salmon feeds compared to marine-based feeds [13]. In contrast, in a study on fish oil replacement in sea bream (*Sparus aurata*) feeds, plant oils did not have elevated PAH levels compared to fish oil, and it was concluded that feed PAH levels did not correlate with the amount of fish or plant oil used [14]. The genotoxic and carcinogenic “heavy” PAHs (>4-6 rings), such as benzo[a]pyrene (B[a]P), benzo[a]anthracene (B[a]A), chrysene (Chr), benzo[b]fluoranthene (B[b]F), have received special attention with regards to food safety [15]. Studies related to plant oil PAH contamination, however, mainly focus on light (2-4 rings) PAHs such as fluoranthene, naphthalene, anthracene, phenanthrene, as they are most dominantly present in unrefined plant oils. These light PAHs are also on the US EPA

(Environmental Protection Agency) list for environmental relevant PAHs but are mostly not carcinogenic and genotoxic [10,16].

The European union has set an upper limit for 4 PAHs (B[a]P +B[a]A+Chr+B[b]F) in food products to protect the consumer safety [15]. No upper limits for PAH in feed ingredients or animal feeds currently exist. The PAH levels of raw ingredients for feeds for farmed terrestrial animals such as cow, chicken, and pig have been reported in only few studies [11] and little is known on PAH levels in feeds for farmed fin fish species [17,18], especially for feed ingredients used in novel plant based fish feed [13,14,19].

Traditionally, the determination of PAHs in solid fatty matrices have involved time-consuming extractions prior to GC-MS with electron ionization (EI) analysis [20]. However, the sensitivity restrictions of the GC-EI-MS(/MS) systems [21,22] in addition to the inherent matrix effects make pre-concentration and time consuming clean-up steps, as gel permeation chromatography [23] or solid phase extraction [19,24], necessary. Recently, some methodologies have applied the QuEChERS procedure in aquaculture samples [25,26], reducing notably the time and cost of sample treatment. However, despite all efforts to minimize the presence of interferences, the still remaining matrix effects make necessary to use matrix-matched calibration for quantification purposes [22]. In the last years, the application of atmospheric pressure chemical ionization (APCI) source in combination with last generation triple quadrupole instruments has made possible to notably increase sensitivity and selectivity of the determination of organic contaminants in food safety, environment and doping control fields [27–30] in selected reaction monitoring (SRM) based methods. The improved performance of GC-APCI-MS/MS is more than welcome in the aquaculture field, where the analysis of a wide diversity of “difficult” matrices (oils, plant ingredients, marine ingredients, terrestrial animal ingredients, fish, etc.) with this

technique can be simplified by reducing the matrix content in the final extract by simple dilution which minimizes the matrix effect and avoids the need of using matrix-matched calibrations.

In this work, we explore the potential of the new GC-APCI-MS/MS technique for quantification and confirmation of PAHs, together with some selected pesticides and PCBs, in very different sample matrices from aquaculture field. Sensitivity of this new technique has been exploited in order to avoid the time-consuming and expensive purification steps commonly used in these complex matrices and with the aim of obtaining a cheap and efficient sample treatment. The developed methodology has been applied to the quantification of compounds that were previously found in wide-scope screening of aquaculture samples by combined use of GC&LC-QTOF MS. Up to our knowledge this is the first application of GC-APCI-MS/MS with triple quadrupole to the analysis of PAHs in samples from aquaculture field.

EXPERIMENTAL

Reagents

Benzo[*j*]fluoranthene (B[*j*]F), 5-methylchrysene (5-MC), benzo[*c*]fluorene (B[*c*]F), dibenzo[*a,e*]pyrene (D[*a,e*]P), dibenzo[*a,h*]pyrene (D[*a,h*]P), dibenzo[*a,i*]pyrene (D[*a,i*]P), dibenzo[*a,l*]pyrene (D[*a,l*]P), and cyclopenta[*c,d*]pyrene (C[*c,d*]P) individual standard solutions at 10 mg·L⁻¹ (100 mg·L⁻¹ for C[*c,d*]P), PAH MIX 9 containing 16 PAHs (10 mg·L⁻¹ in cyclohexane) and PCB mix 3 (100 ng·μL⁻¹ in isooctane) were purchased from Dr. Ehrenstorfer (Augsburg, Germany). Reference standards diphenylamine, ethoxyquin, *p,p'*-DDD, *p,p'*-DDT, tebuconazole, azoxystrobin, fluazinam and imazalil supplied by Dr. Ehrenstorfer GmbH (Augsburg, Germany); boscalid and malathion by Riedel-de Haën (Seelze, Germany); ethoxyquin dimer by

Synthetica AS (Oslo, Norway) and hexachlorobenzene (HCB), pirimiphos-methyl and chlopyrifos-methyl by Sigma-Aldrich (St Louis, MO, USA) with a purity 97-99.9% were used for standard preparation (see structures for all the compounds in **Table S1**). Stock standard solutions (around 500 mg·L⁻¹) were prepared in acetone. Two mixtures of pesticide standards (individual concentration of each pesticide around 50 mg·L⁻¹) were prepared by dilution of stock individual solutions in acetone. Working standard solutions containing all compounds were prepared by dilution of mixtures with acetone (for sample fortification in GC) and hexane (GC injection). Stock standard solutions and working solutions were stored in a freezer at -20 °C.

Stable isotopic labeled internal standards (SIL-IS) PCB 153-¹³C₁₂ and PAH surrogate cocktail containing acenaphthylene-D₈, B[a]P-D₁₂, B(g,h,i)Pe-D₁₂, fluoranthene-D₁₀, naphthalene-D₈, phenanthrene-D₁₀ and pyrene-D₁₀ were purchased from Cambridge Isotope Laboratories (Andover, MA, USA). Hexachlorobenzene-¹³C₆, tebuconazole-D₆ and p,p'-DDE-D₈ were also purchased from Dr. Ehrenstorfer.

HPLC-supragradient acetonitrile, acetone (pesticide residue analysis quality) and n-hexane (ultra-trace quality) were purchased from Scharlab (Barcelona, Spain). Anhydrous magnesium sulfate (extra pure) and anhydrous sodium acetate (reagent grade) were purchased from Scharlab. The QuEChERS commercial clean-up kits were purchased from Teknokroma (Barcelona, Spain). Each kit contains 50 mg of primary-secondary amine (PSA), 150 mg of anhydrous magnesium sulfate, and 50 mg of C₁₈, in 2 mL microcentrifuge tubes for d-SPE.

Samples

The sample material has been described before in detail [31]. A total of 76 samples (from 19 different matrices) were studied in this work. The list contains ingredients

from different origin (plant, terrestrial animals and marine), and also feeds based on these ingredients (PAPs not included), as well as fillets of Atlantic salmon and gilthead sea bream reared on these feeds. Fish individuals were fed for, respectively, 7 and 18 months, and fillet samples were taken for analysis at the end of the exposure trial and additionally for sea bream at 8 months (commercial size). The same feed compositions were provided throughout the feeding trial. The quantification was performed on feed ingredients, feeds produced from the same feed ingredients, and fish fillets of fish fed on these feeds. The feed samples were analysed at the beginning of the trial (additionally after 8 months for sea bream) and no stability assessment was made by analyzing the feed during storage (below 7°C for salmon and sea bream feeds).

Commercially available plant and marine feed ingredients were provided by Biomar (Tech Center, Brande, Denmark) feed producer and processed animal proteins (PAPs) from non-ruminants were provided by the European Fat Processors and Renderers Association (EFPPRA). The ingredients selected represent the novelties in fish feed compositions to reduce the inclusion of fish derivatives. Fish feeds for feeding trials were based on plant feed ingredients, and not PAPs, as higher levels of PAHs were found in plant feed ingredients. The feeds were produced by Biomar under commercial aquafeed production techniques based on high-temperature extrusion processes, which potentially could affect pesticide residue levels.

Sample treatment

Samples were thawed at room temperature and 1 g was accurately weighed and transferred to 15 mL falcon tubes and spiked with 0.2 mL of SIL-IS solution of 1 $\mu\text{g}\cdot\text{mL}^{-1}$. After 30 min, 2 mL of acetonitrile were added and the tube was vigorously shaken by vortex for 30 s. Then, 0.8 g of MgSO_4 were added and the tube was

immediately shaken for 30 s. Subsequently, the tube was centrifuged at 1893 rcf·g for 5 min and the upper layer of the extract was transferred to a 2 mL eppendorf tube and stored for at least two hours in a freezer to precipitate proteins and fix lipids to the tube walls (freezing cleanup). Then, a QuEChERS clean-up step was carried out prior injection in the GC-system [25]. Briefly, 1 mL of the extract was carefully transferred to the cleanup QuEChERS vial (50 mg of PSA + 150 mg of MgSO₄ + 50 mg of C18), and it was vigorously shaken for 30 s and centrifuged at 12557 rcf·g for 5 min. Then, 50 µL of the final acetonitrile extract was diluted with 300 µL of acetone and 650 µL of hexane in order to make the solution miscible (20-fold dilution factor).

Instrumentation

GC-APCI-MS/MS.

Data were acquired using a GC system (Agilent 7890A, Palo Alto, CA, USA) equipped with an autosampler (Agilent 7693) and coupled to a triple quadrupole (QqQ) mass spectrometer (Xevo TQ-S, Waters Corporation, Manchester, UK), operating in positive APCI mode. The GC separation was performed using a fused silica DB-17MS (50% phenyl-methylpolysiloxane) capillary column with a length of 30 m x 0.25 mm i.d. and a film thickness of 0.25 µm with a 1 m x 0.25 mm i.d deactivated post-column (J&W Scientific, Folsom, CA, USA). The oven temperature was programmed as follows: 70 °C (1 min); 20 °C min⁻¹ to 230 °C; 5 °C min⁻¹ to 320 °C (6 min). The injector was operated in pulsed splitless mode (50 psi), injecting 1 µL at 280 °C. Helium was used as carrier gas at a ramped flow mode programmed as follows: 2 mL·min⁻¹ (23 min); a ramp of 4 mL·min⁻¹ to 6 mL·min⁻¹ (9 min). In the SRM method, automatic dwell time (values ranging from 3 to 63 ms) was applied in order to obtain 15 points per peak. The interface and source temperatures were set to 320 °C and 150 °C respectively using N₂

as auxiliary gas at $275 \text{ L}\cdot\text{h}^{-1}$, as make-up gas at $300 \text{ mL}\cdot\text{min}^{-1}$ and as cone gas at $200 \text{ L}\cdot\text{h}^{-1}$. The APCI corona discharge pin was operated at $1.6 \mu\text{A}$. The ionization process occurred within an enclosed ion volume, which enabled control over the protonation/charge transfer processes. In order to work under proton transfer conditions, an uncapped vial containing 1.25 mL of water was placed in a designed holder into the APCI source door to enhance protonation. Targetlynx (a module of MassLynx) was used to handle and process the acquired data.

Study of matrix effects

Nine out of nineteen different matrices (fish fillets, feed and PAPs) were selected for the study of matrix effects and different dilution factors were tested (crude extract, 5-fold, 10-fold and 20-fold dilution). To this aim, matrix-matched calibrations were prepared for each sample matrix by taking between $25\text{-}500 \mu\text{L}$ of the “blank” sample extract (for crude extract, $500 \mu\text{L}$ were evaporated to dryness) and adding $50 \mu\text{L}$ ($125 \mu\text{L}$ for the $25 \text{ ng}\cdot\text{mL}^{-1}$ level) of the corresponding standard in hexane (between 10 and $100 \text{ ng}\cdot\text{mL}^{-1}$), resulting in final analyte concentrations between 1 and $25 \text{ ng}\cdot\text{mL}^{-1}$. The final volume of each calibration point was $500 \mu\text{L}$ in all cases (30% acetone, 5 % acetonitrile and 65% hexane). Matrix effect was evaluated by comparing the calibration graphs obtained with standards in solvent and in matrix (at the different dilution factors) [32]. The accepted relative error between the slopes of calibration in solvent and matrix-matched calibration was $\pm 25\%$. Once the optimum dilution factor was selected, the absence of matrix effect was also tested for the remaining 10 matrices.

Validation and recovery experiments

Eleven SIL-IS were added at the initial stage of the procedure as quality control (surrogates) in order to correct for possible losses during the overall procedure and instrumental deviations. Linearity of relative response of analytes was established by analyzing standard solutions injected in triplicate in the range of 0.01-25 ng·mL⁻¹ for all compounds except for C[c,d]P (0.005-12.5 ng·mL⁻¹). Linearity was assumed when the correlation coefficient (R^2) was higher than 0.99 with residuals lower than 30%.

A full validation of the method was carried out for two sample matrices, salmon and sea bream fillets. Accuracy (estimated by means of recovery experiments) was evaluated by analyzing six replicates spiked at 2 ng·g⁻¹ (1 ng·g⁻¹ for C[c,d]P) and 50 ng·g⁻¹ (25 ng·g⁻¹ for C[c,d]P) for the two matrices tested. Precision was expressed as repeatability in terms of relative standard deviation (RSD, %) (n = 6) calculated for each fortification level. The limit of quantification (LOQ) objective, for salmon and sea bream fillets, was defined as the lowest concentration level validated following SANTE/11945/2015 guide criteria [33] (recovery 70-120% and RSDs below 20%). For those compounds that could not be validated due to the high concentration in the “blank” sample a statistically calculated LOQ was estimated as the analyte concentration that produced a peak signal of ten times the background noise in the chromatogram at the lowest fortification level tested for each compound. The limit of detection (LOD) was estimated for all compounds considering a signal-to-noise (S/N) ratio of three.

The methodology was validated for the remaining 17 sample matrices by analysis of quality control (QC) samples, i.e. “blank” samples spiked at 5 and 50 ng·g⁻¹ injected in each batch. A total of 48 QCs prepared from 24 “blank” samples (corresponding to 17 different matrices) were analyzed. Recoveries between 60 and 140% for each individual sample were considered satisfactory following the acceptance criteria for routine recoveries according to SANTE/11945/2015 guide criteria [33].

The selectivity of the method was assured by choosing selective SRM transitions using M^{++} or $[M+H]^+$ as precursor ion, when possible. Specificity was evaluated by verification of the absence of interfering peaks at the retention time of each compound in blank samples for the acquired SRM transitions.

The q_i/Q ratio, defined as the ratio between the intensity of the confirmation ion (q_i) and the quantification ion (Q), was used to confirm peak identity in real and spiked samples. The experimental q_i/Q value for each compound was calculated as the mean value obtained from three standard solutions injected in triplicate (RSD below 15% in all cases) Confirmation of analytes detected in samples was considered positive when the q_i/Q ratio was within acceptable tolerances (30% of the experimental q_i/Q value calculated from standards) according to SANTE/11945/2015 guide criteria [33].

RESULTS AND DISCUSSION

Ionization and fragmentation behaviour of PAHs in GC-APCI

In this work, the ionization behavior of the APCI interface was tested using PAH standards in solvent. Two mechanisms of ionization were simultaneously observed: i) charge transfer in which the nitrogen plasma created by the corona discharge needle promotes the formation of M^{++} , and ii) proton transfer, where the presence of water vapor traces in the source favors the formation of the $[M+H]^+$ ion.

The PAHs studied showed a mixture of two ions, M^{++} and $[M+H]^+$, as base peak of the spectrum. In the case of acenaphthene, fluorene, phenanthrene, anthracene, fluoranthene, pyrene, B[c]F and C[c,d]P, the $[M+H]^+$ intensity was slightly higher than M^{++} (among 1.2 and 1.5 times higher). In contrast, the intensity of M^{++} was around 1.1 and 1.2 times higher than $[M+H]^+$ for naphthalene, I[1,2,3-cd]P, B[g,h,i]Pe, D[a,h]A, D[a,h]P, D[a,e]P, D[a,i]P and D[a,l]P. For the rest of PAHs, the abundance of M^{++} and

$[M+H]^+$ was similar. **Figure 1 (up)** shows the APCI spectrum of D[a,l]P where $M^{+\bullet}$ and $[M+H]^+$ can be observed as base peak of the spectrum. After that, the fragmentation of the PAHs in the collision cell was studied. $M^{+\bullet}$ and $[M+H]^+$ were selected as precursor ions for all PAHs. Fragmentation was performed at collision energies between 10-60 eV. The losses of H^\bullet and H_2 from both $M^{+\bullet}$ and $[M+H]^+$ were the most abundant and common to all PAHs studied. Losses as CH_3^\bullet or C_2H_2 among others were also commonly observed. The use of water as modifier favored the formation of the $[M+H]^+$ and the $M^{+\bullet}$ decreased in all cases (see **Figure 1, down**). However, $M^{+\bullet}$ still appeared in the spectra in a percentage among 35-80%, depending on the PAH.

Sensitivity and repeatability. Effect of different modifiers

Once the SRM transitions were optimized, the sensitivity and repeatability of the GC-MS/MS signals were evaluated for all the transitions selected (those coming from $M^{+\bullet}$ as well as from $[M+H]^+$). As explained above, both $[M+H]^+$ and $M^{+\bullet}$ were observed in the APCI spectra of all PAHs even under “dry” conditions. Thus, we investigated the effect of different modifiers in order to promote the formation of the protonated molecules and to increase the sensitivity of SRM transitions coming from $[M+H]^+$. Water, 0.5% formic acid, methanol and isopropanol were added separately in an uncapped vial, which was located within a specially designed holder placed in the source door. The sensitivity and repeatability of the GC-MS/MS signals for PAHs were evaluated under the different conditions (with and without the use of modifiers).

As can be seen in **Figure 2 (up)**, under charge transfer conditions (without adding a modifier), SRM transitions coming from $M^{+\bullet}$ were more sensitive than those coming from $[M+H]^+$ for acenaphthene, fluorene, B(c)F, C[c,d]P, chrysene, 5-MC, B(b)F and B(j)F (among 2 and 4 times higher). In contrast, the sensitivity of SRM transitions

coming from $[M+H]^+$ was higher than $M^{+\bullet}$ (among 3 and 14 times) for naphthalene, acenaphthylene, phenanthrene, anthracene, fluoranthene, pyrene and B[g,h,i]Pe. For the rest of PAHs, they were mostly similar. When adding water as modifier, the sensitivity of transitions coming from the $M^{+\bullet}$ decreased between 1.5 to 3 times for all compounds, except for D[a,h]A which was rather similar; and consequently, the intensity for $[M+H]^+$ transitions improved 1.5 to 2 times for high molecular weight PAHs (HMW-PAHs, with four or more aromatic rings).

When adding HCOOH 0.5%, the sensitivity of $[M+H]^+$ transitions remained almost the same as with only water (**Figure 2 (down)**). Similar results were obtained using MeOH as modifier, except for some compounds, such as naphthalene, fluorene, phenanthrene, fluoranthene, chrysene and B(b)F, among others, whose responses decreased considerably. Regarding isopropanol, its use was not beneficial because sensitivity extremely decreased for most of the compounds. Thus, the modifiers MeOH and isopropanol were discarded for further optimization.

Once these modifiers were discarded, the repeatability of the response ($n=10$ at $10 \text{ ng}\cdot\text{mL}^{-1}$) was also evaluated under the remaining working conditions (without water, with water and with HCOOH 0.5%) (**Figure S1**). Our data showed a slightly poorer repeatability for $M^{+\bullet}$ and $[M+H]^+$ transitions working under dry conditions (relative standard deviations, RSD, between 20-30% for most compounds) being a bit better (RSD 10-17%) for fluoranthene, pyrene and low molecular weight PAHs (LMW-PAHs, 2-3 aromatic rings). This parameter was improved under wet conditions using water as modifier for $M^{+\bullet}$ (RSD 7-20%) and specially for $[M+H]^+$ transitions (RSD 2-8%), but dramatically get worst when adding HCOOH 0.5%, especially for $M^{+\bullet}$ transitions (RSD 14-39%).

Overall, the use of water as modifier was selected for further experiments, selecting the transitions from $[M+H]^+$ for most of PAHs. In some particular cases, the most sensitive transitions using $M^{+\bullet}$ as precursor ion were used for confirmation purposes (except for b[j]F which was used for quantification) (**Table 1**).

In a later stage, the optimum conditions selected for PAHs were tested for the PCBs and pesticides in order to widen the applicability of the methodology to other relevant pollutants included in the method. In previous works, it was reported that sensitivity for halogenated hydrocarbons without any other heteroatoms, e.g. PCBs and many organochlorine pesticides (such as hexachlorobutadiene, pentachlorobenzene, hexachlorobenzene, HCHs, DDTs, trans-chlordane, mirex) was negatively affected by the introduction of water in the source [34,35]. Despite this fact, the sensitivity and repeatability of PCBs, p,p'-DDT, p,p'-DDD, p,p'-DDE and HCB was still enough to reach the maximum residue level regulated. Regarding the other pesticides included in the method, ionization efficiency was favored under proton-transfer conditions, i.e. selecting $[M+H]^+$ as precursor ion, except for ethoxyquin and ethoxyquin dimer that were scarcely affected.

Selectivity of the SRM transitions for PAHs

Selectivity of the selected SRM transitions, especially in cases where coelution occurred, was also studied during the optimization process. In such cases, the possibility of selecting different precursor ions for the same compound ($M^{+\bullet}$ or $[M+H]^+$) was of great advance in terms of specificity. This was the case of the determination of cyclopenta(c,d)pyrene ($C_{18}H_{10}$, MW=226), which nearly coelutes with chrysene ($C_{18}H_{12}$, MW=228) (see **Figure S2**). Chrysene shows 226 as an in-source fragment (coming from $M^{+\bullet}$ 228). Thus, when monitoring C(c,d)P with the transition coming

from its M^{+} 226, an interference from chrysene (coming from an in-source fragment) is observed in the chromatogram, which makes the peak area integration troublesome, especially when the concentration of chrysene is higher than cyclopenta(c,d)pyrene. However, quantification and confirmation transitions for C(c,d)P coming from $[M+H]^{+}$ 227 did not suffer from this interference and showed higher specificity, which is commonly observed on EI-based methods [36]. A similar case occurred for the determination of indeno(1,2,3-cd)pyrene ($C_{22}H_{12}$, MW=276) due to the nearly coelution of dibenzo(a,h)anthracene ($C_{22}H_{14}$, MW=278). D(a,h)A shows an in-source fragment at m/z 276, so the transition coming from M^{+} 276 for I(1,2,3,cd)P was interfered by D(a,h)A, while those coming from $[M+H]^{+}$ 277 were much cleaner.

Study of matrix effects

Matrix effects observed under GC-APCI-MS come from the GC liner (normally signal enhancement) and from the APCI source (normally signal suppression) [27]. As a summary of the matrix effect study for 9 out of 19 matrices (where different dilution factors were tested: crude extract, 5-fold, 10-fold and 20-fold dilution), signal enhancement/suppression was observed for: i) 36% of the combinations matrix/PAHs, 58% for matrix/pesticides and 52% for matrix/PCBs (crude extract); ii) 35% for matrix/PAHs, 46% for matrix/pesticides and 17% for matrix/PCBs (5-fold dilution factor); iii) 26% of matrix/PAHs, 33% for matrix/pesticides and 14% for matrix/PCBs (10-fold dilution factor); and iv) 11% for matrix/pesticides (20-fold dilution factor). As an illustrative example, **Figure 3** shows how the matrix effect for PAHs in sea bream fillet decreases when the different dilution factors studied are applied. Thus, 20-fold dilution of the final extract was finally selected for further experiments.

As a summary, for a 20-fold dilution, signal enhancement was observed only for ethoxyquin and ethoxyquin-dimer, whereas signal suppression was observed for boscalid and azoxystrobin (mainly in plant and marine ingredients). No significant matrix effect was observed for PAHs, PCBs and the rest of pesticides (**Figure S3**). According to our data, 20-fold dilution of the final sample extract eliminated matrix effect for most analyte/matrix combinations, and this allowed us to perform the quantification in the analysis of the 19 aquaculture matrices using calibration curve in solvent. The remaining matrix effects for boscalid and azoxystrobin could be corrected with the use of SIL-IS. Matrix effect for ethoxyquin and ethoxyquin-dimer could not be properly evaluated in feed, krill meal and fishmeal matrices because the analyte concentration in these samples was typically higher than the spiking levels assayed.

Analytical parameters

Validation of the analytical procedure was carried out in salmon and sea bream fillets considering the parameters of linearity, accuracy, precision, LODs and LOQs. Relative responses to the selected SIL-IS were used (see **Table 1**).

The study of the linearity revealed that correlation coefficients (R^2) were higher than 0.99, with residuals lower than 30% for most compounds in the range 0.01-25 ng·mL⁻¹. Some exceptions were: 0.0125-12.5 ng·mL⁻¹ (C[c,d]P); 0.025-25 ng·mL⁻¹ (naphthalene, acenaphthylene, acenaphthene, anthracene, B[c]F, B[a]A, chrysene, B[b]F, B[k]F, B[j]F, D[a,h]A, PCB 153 and 180, diphenylamine, tebuconazole, ethoxyquin dimer, boscalid and azoxystrobin); 0.1-25 ng·mL⁻¹ (D[a,l]P and D[a,e]P, HCB and p,p'-DDE); 0.5-25 ng·mL⁻¹ (D[a,i]P, D[a,h]P, ethoxyquin, fluazinam, imazalil, p,p'-DDD and p,p'-DDT).

Trueness and precision data for salmon and sea bream fillets are shown in **Table 2** which shows satisfactory recoveries and precision for the wide majority of compounds at the two concentrations studied, 2 and 50 ng·g⁻¹. An LOQ objective of 2 ng·g⁻¹ was empirically demonstrated for most compounds in both matrices, using samples spiked at this level and subjected to the overall analytical procedure. A statistical LOQ of 5 ng·g⁻¹ (S/N = 10) was estimated for naphthalene, fluorene, fluoranthene and pyrene, as the high concentrations present in the “blank” samples did not allow the calculation of recoveries and precision below that concentration. Dibenzopyrenes have poor response, and therefore the LOQ was established in 50 ng·g⁻¹. The same occurred for HCB, fluazinam, imazalil, p,p'-DDE, p,p'-DDD and p,p'-DDT which could not be detected at the low level of 2 ng·g⁻¹ being 50 ng·g⁻¹ still the normally applied MRL for these compounds (as commented above, these compounds were negatively affected by the introduction of water as modifier in the source). Similarly to some light PAHs, ETQ and ETQ-D were found at quite high concentrations in the samples impeding validation at 2 ng·g⁻¹ level. Thus, LOQs were statistically calculated for S/N=10. LODs were found to be between 0.5-2 ng·g⁻¹ for 80 % of the PAHs, all the PCBs and 50% of the pesticides studied. The rest of compounds showed LODs between 5 and 50 ng·g⁻¹.

In relation to the remaining 17 matrices, a total of 48 QC samples, spiked at 5 ng·g⁻¹ and 50 ng·g⁻¹, were analyzed. **Figure 4** shows the box-plot diagrams representing the recovery values for PAHs in all matrices at the two concentrations assayed. Some recoveries could not be calculated (e.g. naphthalene, fluorene, phenanthrene, pyrene, D[a,e]P, D[a,i]P and D[a,h]P) at 5 ng·g⁻¹ spiking level due to the high analyte concentration present in “blank” sample. It can be emphasized that all QC recoveries were in the range 60 - 140%, with median values between 88 and 115 %, at 5 ng·g⁻¹, and between 92 and 116% at 50 ng·g⁻¹. Among the studied matrices, PAPs showed

higher complexity giving individual recoveries out of 70-120% range without a clear tendency for specific compounds. Data from above illustrate that the methodology applied was robust and satisfactory for the determination of very low PAH concentrations in different sample matrices, some of them of high complexity. The excellent sensitivity of GC-APCI-MS/MS allowed the extra dilution (x20) of the sample extracts, minimizing matrix effects but still reaching low LOQs, comparable and, in some cases, even lower than those reported in the literature [21,22] with the great advantage of using solvent calibration for quantification of the compounds in all matrices tested.

Analysis of samples

In order to demonstrate the applicability of the methodology developed in this paper, it was applied to the analysis of a notable number of samples from aquaculture production: 12 diets, 12 fish fillets, 19 terrestrial animal ingredients, 6 marine ingredients, 19 plant ingredients and 8 plant oils. A reagent blank and two spiked samples (5 and 50 ng·g⁻¹) for each matrix analyzed were included in each batch. The results, in **Table S2**, are expressed in a wet weight basis. The q_i/Q ratios obtained for all positive samples were within the range of the tolerance accepted (30%) around the experimental q_i/Q value obtained from reference standards.

LMW-PAHs together with HMW-PAHs as fluoranthene and pyrene were found in most of the samples. Regarding diets, fish fillets, plant ingredients and PAPs, concentrations ranged from <5 ng·g⁻¹ for acenaphthene to 126.6, 77.1, 88.6 and 177.5 ng·g⁻¹ for naphthalene, respectively. In marine ingredients, they ranged from <5 ng·g⁻¹ for acenaphthene and acenaphthylene to 129.0 ng·g⁻¹ for phenanthrene. For plant oil ingredients, concentrations ranged from <5 ng·g⁻¹ for acenaphthylene to 186.9 ng·g⁻¹ for

phenanthrene. In the present study only two fish oils were analysed, and clearly a large variation exist in the environmental background levels of PAHs in fish oils [13,14]. **Figure S3** shows the identification of 5-MC in one fish meal. This PAH was below the LOQ in all samples analyzed. As can be seen, three transitions were acquired and q_i/Q ratios were in agreement with SANTE/11945/2015 identification criteria followed in the current research [33].

Regarding pesticides, the non-organochlorine pesticides that are currently used on crop such as chlorpiriphos-methyl and pirimiphos-methyl were only detected in the novel plant ingredients and not in marine feed ingredients such as fish oil and meal [25]. Malathion, imazalil and tebuconazole were only found in some ingredients while boscalid and azoxystrobin were indeed found in feed. It is worth to mention that studied PCBs, DDTs and pesticides were not found in fillets from salmon and sea bream. The anti-oxidant ethoxyquin and its main metabolite ethoxyquin dimer are deposited to marine feed ingredients such as fish meal to prevent lipid oxidation and spontaneous combustion during overseas transport and storage (Lundebye et al. 2010). The ethoxyquin levels were highest in fish meal and krill (32462 and 40076 $\text{ng}\cdot\text{g}^{-1}$, respectively), but surprisingly high levels were also found in fish oil (37137 $\text{ng}\cdot\text{g}^{-1}$) where normally other antioxidant such as BHT and BHA are used [37]. All other feed ingredients, including terrestrial and plant, also contained ethoxyquin, albeit at 100-150 fold lower levels than the marine ingredients (**Table S2**). PCB congeners were not detected in any sample.

CONCLUSIONS

The use of APCI has been evaluated for GC-MS/MS analysis of PAHs. The high sensitivity of this technique allowed the simultaneous quantification of 19 different

complex matrices (from aquaculture field) using solvent calibration. The key aspect for this aim was the elimination of matrix effect, without the need of time-consuming purification steps, only by a 20-fold dilution factor of the final QuEChERs extract, being this aspect a great contribution of the present work. Despite this, LOQs of the developed method were $2 \text{ ng} \cdot \text{g}^{-1}$ for most analytes, in the same order or better than those reported in previously published methods for similar matrices [21,22] showing higher efficiency [19]. Also, the use of both M^{++} or $[\text{M}+\text{H}]^+$ gave an additional value to the selectivity in the determination/identification capabilities.

FIGURE CAPTIONS

Figure 1. APCI mass spectra of D[a,l]P under charge transfer conditions (up) and under proton transfer conditions with water as modifier (down).

Figure 2. Absolute sensitivity of SRM transitions coming from M^{++} and $[\text{M}+\text{H}]^+$ for PAHs under charge transfer and proton transfer (with water) ionization conditions (up); absolute sensitivity of SRM transitions coming from $[\text{M}+\text{H}]^+$ under proton transfer conditions using different modifiers (bottom).

Figure 3. Matrix effect study for PAHs in sea bream fillet. Relative error between the slopes of calibration in solvent and matrix-matched calibration at different dilution factors.

Figure 4. Box plots that shows the recoveries of the 48 QCs (corresponding to 17 different matrices) spiked at 5 (up) and 50 $\text{ng} \cdot \text{g}^{-1}$ (down).

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Table 1. Experimental conditions of the optimized GC-APCI-MS/MS method using water as modifier. Quantifier (Q) and qualifier (q) transitions.

Rt (min)	Compound	Internal Standard	Precursor ion	Product ion	CE (eV)	q/Q ratio	Rt (min)	Compound	Internal Standard	Precursor ion	Product ion	CE (eV)	q/Q ratio	
5.53	Naphthalene-D ₈		136	134.1	30	Q	13.98	B[c]F	Pyrene-D ₁₀	217	215.9	20	0.129	
5.55	Naphthalene	Naphthalene-D ₈	129	128.1	20	Q					215	30	0.370	
				126.8	20	0.273					202	30	Q	
				103	20	0.011					200.1	40	0.048	
				79	20	0.002				216	215	30	0.393	
			128	102.1	20	0.063	14.27	PCB 138	¹³ C ₁₂ -PCB 153	360	325	20	0.683	
7.65	Acenaphthylene-D ₈		160	158.3	30	Q					290	40	Q	
7.66	Acenaphthylene	Acenaphthylene-D ₈	153	152	20	Q					218	60	0.559	
				150.1	30	0.010					253	50	0.150	
				125.9	30	0.018	14.31	4,4'-DDT	4-4'-DDE D ₈	235	165	20	Q	
				126.1	30	0.018					115	50	0.026	
7.82	Acenaphthene	Acenaphthylene-D ₈	155	154	20	0.539					99	40	0.023	
				153	20	Q	14.64	Tebuconazole-D ₆		314	296	10	Q	
				129.2	20	0.367	14.70	Tebuconazole	Tebuconazole-D ₆	308	290	10	Q	
				128	30	0.317					165	20	0.430	
				153	20	0.852					151	20	0.246	
8.42	Fluorene	Acenaphthylene-D ₈	167	166	20	0.349					125	40	0.865	
				165.1	20	Q	15.67	PCB 180	¹³ C ₁₂ -PCB 153	392	322	40	0.557	
				152	20	0.956					394	324	30	Q
				115	30	0.064					396	361	20	0.454
				166	165.1	20	0.607				254	60	0.448	
8.72	Diphenylamine	Tebuconazole-D ₆	170	93	30	Q	16.80	B[a]A	Pyrene-D ₁₀	229	228	40	Q	
				168	30	0.035					226.9	30	0.262	
				153	20	0.086					226	40	0.629	
				152	20	0.070					202.2	30	0.148	
				169	168	20	0.506				226	40	0.568	
8.87	HCB	HCB- ¹³ C ₆	282	247	30	Q	17.07	C[c,d]P	Pyrene-D ₁₀	227	226	30	Q	
				212.1	30	0.718					225	40	0.076	
				177	40	0.557					200.8	40	0.002	
				142	50	0.435					226	224.9	40	0.039
8.86	HCB- ¹³ C ₆		292	222	40	Q	17.11	Chrysene	Pyrene-D ₁₀	229	227.9	40	Q	
9.01	Ethoxyquin	Tebuconazole-D ₆	218	202	20	0.576					227	30	0.440	
				174	20	Q					226	40	0.842	
				148	20	0.671					202.1	30	0.219	
				160	30	0.510					226	40	0.579	
				217	202	10	0.273	18.75	5-MC	Pyrene-D ₁₀	243	242	30	0.043
9.81	Phenanthrene-D ₁₀		188	186.4	20	Q					228	30	Q	
9.85	Phenanthrene	Phenanthrene-D ₁₀	179	178.1	20	Q					226.1	40	0.261	
				176.7	30	0.025					202.1	40	0.066	
				176.1	40	0.044					242	241.1	20	0.227
				152.1	30	0.083	19.96	Ethoxyquin dimer	Tebuconazole-D ₆	433	418	10	0.076	
				152	30	0.083					216	20	Q	
9.88	Anthracene	Phenanthrene-D ₁₀	179	178	20	Q					188	30	0.744	
				176.7	30	0.025					432	417	20	0.307
				176.1	50	0.026					173	50	0.354	
				152.1	30	0.069	21.06	B[b]F	B[a]P-D ₁₂	253	251.9	30	Q	
				152	30	0.112					250.8	40	0.064	
9.98	PCB 28	4-4'-DDE D ₈	256	221	20	0.060					250.1	50	0.349	
				186	30	Q					226	50	0.051	
				151	40	0.177					252	250	40	0.188
10.07	Fluazinam	Tebuconazole-D ₆	465	372.9	20	Q	21.16	B[k]F	B[a]P-D ₁₂	253	251.9	30	Q	
				337.7	40	0.087					250.8	40	0.046	
				303.9	40	0.077					250.1	50	0.174	
				268.9	50	0.045					226	50	0.011	
10.25	Chlorpyrifos-methyl	Tebuconazole-D ₆	322	290	10	0.161					252	250	40	0.165
				212	30	0.065	22.28	B[j]F	B[a]P-D ₁₂	253	251.9	30	0.899	
				177	40	0.021					248.6	60	0.020	
				125	20	Q					226	50	0.269	
10.39	PCB 52	4-4'-DDE D ₈	290	220	30	Q					224.1	60	0.120	
				292	257	20	0.605				252	250	40	Q
				220	30	0.689	21.97	Boscalid	Tebuconazole-D ₆	343	307	20	0.336	
				150	50	0.334					140	20	Q	
10.41	Pirimiphos-methyl	Tebuconazole-D ₆	306	164	20	Q					112	40	0.413	
				108	40	0.505					342	140	20	0.039
				125	30	0.269	22.75	B[a]P-D ₁₂		264.3	262.3	30	Q	
				95	20	0.145	22.66	B[a]P	B[a]P-D ₁₂	253	252.1	30	Q	
10.71	Malathion	Tebuconazole-D ₆	331	211	10	0.026					250.7	40	0.023	
				143	30	0.017					250.2	50	0.201	
				125	30	Q					226	50	0.037	
				117	20	0.047					252	250	40	0.146
				99	10	0.732	24.97	Azoxystrobin	Tebuconazole-D ₆	404	372	10		
11.69	PCB 101	4-4'-DDE D ₈	324	254	40	Q					344	20	0.243	
				326	291	20	0.938				329	30	0.414	
				256	40	0.936					172	50	0.212	
				184	50	0.081					156	50	0.211	
12.17	Fluoranthene-D ₁₀		212.2	210.3	30	Q	25.19	I[1,2,3-cd]P	B[g,h,i]Pe -D ₁₂	277	276.1	30	Q	
12.22	Fluoranthene	Fluoranthene-D ₁₀	203	202.1	30	Q					275.4	50	0.123	
				200	40	0.069					274.3	50	0.192	
				175.9	50	0.007	25.22	D[a,h]PA	B[a]P-D ₁₂	279	278.1	30	Q	
				152	50	0.014					276.8	40	0.161	
				202	200.8	30	0.053				275.9	40	0.333	
12.30	4,4'-DDE-D ₈		324	254	40	Q					263.3	30	0.267	

12.34	4,4'-DDE	4-4'-DDE D ₈	316	246.1	30	Q	25.87	B[g,h,i]Pe -D ₁₂		288.2	286.3	30	Q
				210	40	0.083	25.93	B[g,h,i]Pe	B[g,h,i]Pe -D ₁₂	277	276.2	40	Q
				281	20	0.162					275.1	50	0.153
				176	50	0.483					274	50	0.154
12.51	Imazalil	Tebuconazole-D ₆	297	254.8	10	0.031				276	272.7	60	0.005
				201.1	10	0.050					275.1	50	0.053
				176.3	20	0.334	28.86	D[a,l]P	B[g,h,i]Pe -D ₁₂	303	302.1	30	Q
				159	20	Q					300.8	30	0.067
				109	20	0.259					300	50	0.455
12.93	Pyrene-D ₁₀		212.2	210.3	30	Q					276	40	0.057
12.97	Pyrene	Pyrene-D ₁₀	203	202.2	30	Q				302	300	50	0.288
				200.8	40	0.067	29.92	D[a,e]P	B[g,h,i]Pe -D ₁₂	303	302.1	30	Q
				200	50	0.041					300.1	50	0.341
				176.1	50	0.003					298.6	60	0.008
				202	30	0.039					297.9	60	0.022
13.03	PCB 118	4-4'-DDE D ₈	326	291	20	0.107				302	300	50	0.183
				256	40	Q	30.59	D[a,i]P	B[g,h,i]Pe -D ₁₂	303	302.1	30	Q
				184	50	0.470					300.1	50	0.245
				324	40	0.968					297.9	60	0.019
13.28	PCB 153	¹³ C ₁₂ -PCB 153	360	325	20	0.478					276	50	0.033
				290	40	Q				302	300	50	0.178
				218	60	0.551	31.00	D[a,h]P	B[g,h,i]Pe -D ₁₂	303	302.1	30	Q
				358	253	0.151					300.1	50	0.274
13.27	PCB 153- ¹³ C ₁₂		376	306	40	Q					297.9	60	0.019
13.65	4,4'-DDD	Tebuconazole-D ₆	235	165	20	Q					276	50	0.033
				115	50	0.028				302	300	50	0.219
				99	40	0.029							

Table 2. Mean recoveries (%) and precision (RSD, %) for PAHs, PCBs and pesticides after application of overall analytical procedure to salmon fillet and gilthead fillet (n = 6), LOQs objective of the method.

Compounds	Salmon fillet			Sea bream fillet		
	Spiking levels (ng _{-g} -1)		LOQ objective (ng _{-g} -1)	Spiking levels (ng _{-g} -1)		LOQ objective (ng _{-g} -1)
	2	50		2	50	
Naphthalene	^a	98 (18)	5 ^b	^a	98 (17)	5 ^b
Acenaphthylene	119 (11)	101 (7)	2	106 (8)	103 (2)	2
Acenaphthene	74 (19)	72 (6)	2	93 (12)	73 (3)	2
Fluorene	^a	95 (11)	5 ^b	^a	108 (6)	5 ^b
Phenanthrene	^a	116 (9)	2 ^b	^a	113 (9)	2 ^b
Anthracene	91 (8)	104 (5)	2	101 (8)	107 (6)	2
Fluoranthene	^a	103 (7)	5 ^b	^a	96 (11)	5 ^b
Pyrene	^a	106 (7)	5 ^b	^a	99 (14)	5 ^b
B(c)F	98 (11)	94 (8)	2	90 (3)	90 (2)	2
B(a)A	114 (5)	101 (5)	2	113 (9)	73 (2)	2
C(c,d)P*	108 (7)	115 (4)	1	112 (19)	112 (7)	1
Chrysene	107 (12)	101 (4)	2	116 (8)	76 (6)	2
5-MC	117 (5)	102 (5)	2	108 (7)	75 (3)	2
B(b)F	80 (13)	104 (7)	2	106 (20)	96 (4)	2
B(k)F	71 (8)	103 (6)	2	78 (7)	86 (4)	2
B(j)F	104 (17)	76 (4)	2	115 (18)	102 (5)	2
B(a)P	104 (3)	88 (7)	2	113 (14)	86 (5)	2
I(1,2,3,cd)P	83 (12)	104 (5)	2	112 (9)	102 (3)	2
D(a,h)A	113 (11)	100 (5)	2	109 (11)	81 (6)	2
B(g,h,i)Pe	84 (8)	101 (4)	2	98 (20)	101 (3)	2
D(a,l)P	-	97 (4)	50	-	77 (6)	50
D(a,e)P	-	95 (8)	50	-	72 (9)	50
D(a,i)P	-	86 (13)	50	-	95 (13)	50
D(a,h)P	-	81 (10)	50	-	73 (6)	50
Diphenylamine	119 (20)	100 (17)	2	-	98 (17)	50
HCB	-	112 (11)	50	-	116 (17)	50
Ethoxyquin	^a	^a	15 ^b	^a	^a	75 ^b
Fluazinam	-	76 (4)	50	-	107 (15)	50
Chlorpiriphos-methyl	77 (10)	100 (6)	2	113 (13)	80 (7)	2
Pirimiphos-methyl	92 (12)	82 (4)	2	115 (13)	110 (9)	2
Malathion	95 (20)	94 (13)	2	90 (11)	104 (10)	2
4,4'-DDE	-	95 (19)	50	-	76 (19)	50
Imazalil	-	87 (8)	50	-	112 (14)	50
4,4'-DDD	-	112 (9)	50	-	114 (18)	50
4,4'-DDT	-	109 (9)	50	-	115 (11)	50
Tebuconazole	103 (19)	109 (11)	2	99 (16)	113 (12)	2
Ethoxyquin dimer	^a	^a	5 ^b	^a	^a	15 ^b
Boscalid	115 (7)	103 (17)	2	103 (4)	106 (11)	2
Azoxystrobin	95 (16)	102 (14)	2	119 (10)	109 (9)	2
PCB-28	106 (13)	117 (7)	2	110 (11)	120 (8)	2
PCB-52	107 (19)	100 (12)	2	98 (20)	97 (12)	2
PCB-101	106 (17)	120 (3)	2	109 (15)	118 (3)	2
PCB-118	110 (21)	105 (3)	2	98 (19)	121 (3)	2
PCB-153	95 (20)	120 (14)	2	87 (15)	115 (16)	2
PCB-138	115 (6)	118 (13)	2	117 (8)	116 (13)	2
PCB-180	105 (11)	111 (13)	2	81 (18)	108 (3)	2

^a Recoveries could not be calculated due to the high concentration in the blank samples

^b LOQ was estimated as the analyte concentration that produced a peak signal of ten times the background noise in the chromatogram at the lowest fortification level studied for each compound

*C[c,d]P spiked at 1 ng_{-g}-1 and 25 ng_{-g}-1

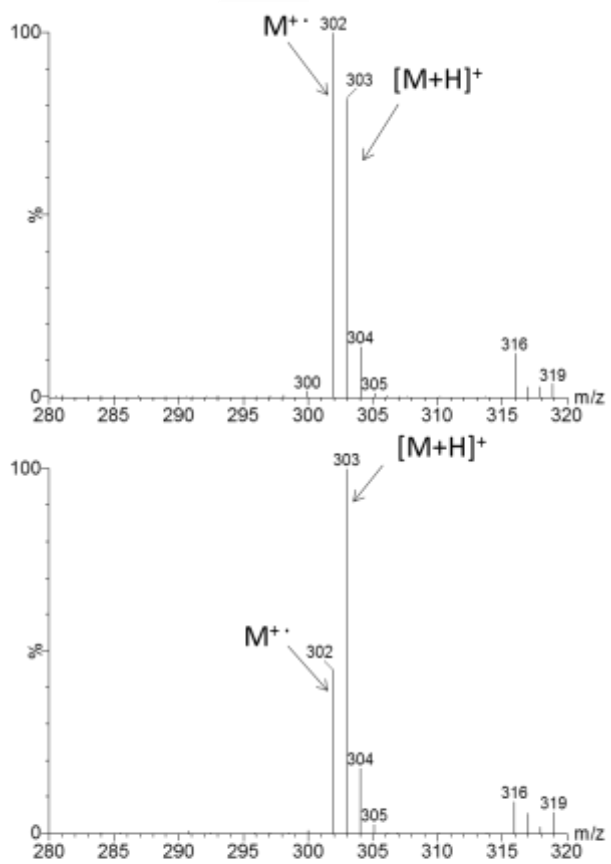


Figure 1.

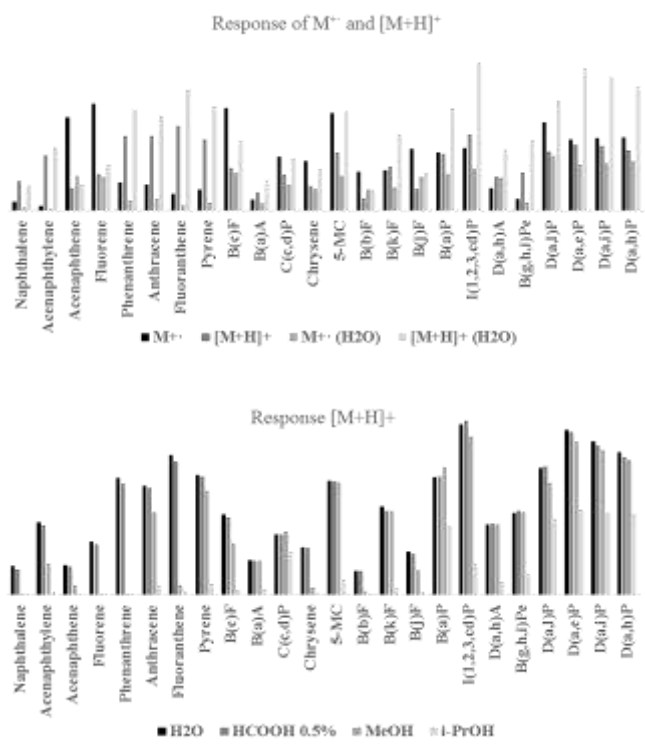


Figure 2

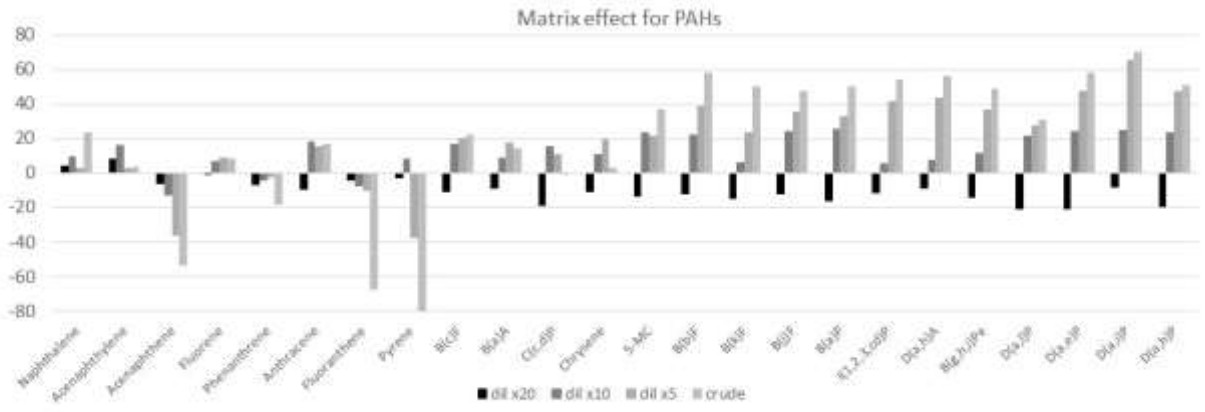
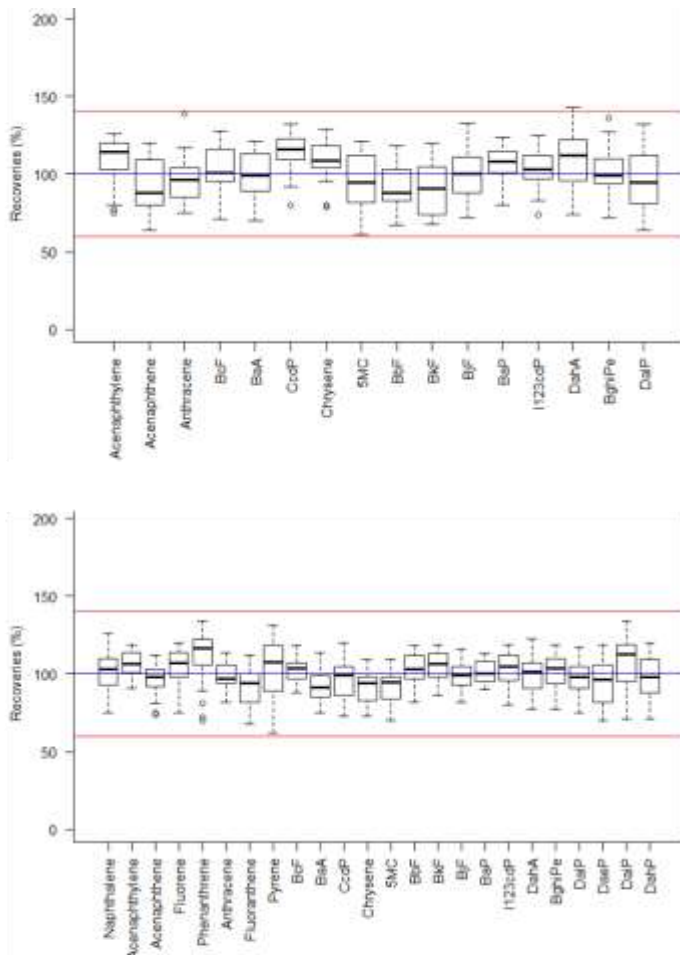


Figure 3



* Recoveries for naphthalene, fluorene, phenanthrene, pyrene, D[a,e]P, D[a,i]P and D[a,h]P could not be calculated at 5 ng/g spiking level due to the high concentration present in blank sample

Figure 4

(not colour necessary)