



The effect of adding a standard on the result of determination of polychlorinated biphenyls in bottom sediment samples

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ARTICLE INFO

Article history:

Received 9 February 2010

Received in revised form 22 April 2010

Accepted 10 May 2010

Available online 19 May 2010

Keywords:

Polychlorinated biphenyls

Bottom sediment

Internal standard

Extraction technique

ABSTRACT

Bottom sediments are a very important component of aquatic ecosystems. The sediment matrix is highly diverse and heterogeneous; in consequence, compounds entering the aquatic environment from different sources are considerably enriched at its surface. Bottom sediments are regarded as natural sorbents, since they accumulate many harmful substances, such as heavy metals and stable organic contaminants.

Extraction is a key stage in every analytical procedure. It is during this stage that standards are added to samples. Standards are necessary not only for estimating analyte yields but also for validating the whole procedure. The question of the addition of standard substances to sediment samples has not been widely addressed in the subject literature, and yet it is of fundamental importance as regards obtaining reliable results of determinations.

This paper describes the results of a study on the effect of standard addition techniques on the results of determination of polychlorinated biphenyls in sediment samples (certified reference material: METRANALTM2—river sediment).

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1. Introduction

Bottom sediments are a very important component of aquatic ecosystems [1]. Highly diverse and heterogeneous, the sediment matrix consists of two main parts [2–6]: an inorganic part made up of clays, silts, muds and sand, and a part of organic origin. The organic part can be further subdivided into:

- an amorphous phase (soft, plastic), consisting primarily of organic matter of animal and vegetable origin in various stages of decomposition;
- a condensed phase (hard, glassy), dominated by carbon geosorbents, e.g. various forms of carbon, including coal and kerogen [7–9].

These phases contain polar (–OH, –COOH, –NH₂, –OCH₃, =NH), non-polar and spatial (aromatic rings) fragments [10]. This diversity of composition means that compounds entering the aquatic environment from various sources undergo considerable enrichment at the sediment surface. Because they accumulate numerous harmful substances, like heavy metals and stable organic contam-

inants, bottom sediments are regarded as natural sorbents [5]. Among the stable organic contaminants that have received a lot of attention in recent years are polychlorinated biphenyls (PCBs) [11].

PCBs are entirely anthropogenic and are carried into the environment primarily with wastewaters. In sediments they are determined at considerably lower concentration levels—in only a few cases do their concentrations exceed 100 µg/kg [12,13]. The determination of both PCBs in sediments is required by international, national and local regulations.

Extraction plays an important part in isolating analytes from the sample matrix [14]. In view of the broad diversity of available techniques, recommending one that is optimal for isolating PCBs from sediments is not very efficient (17–30%) [15]: such a low yield and the diversity of the sediment matrix means that only the internal standard technique can be recommended for determining PCBs. An important stage in this procedure is therefore the addition of the internal standard to the sediment matrix. Once added to the sediment, the internal standard should be bound to it in much the same way as analytes are bound to it. The problem of adding standards to sediment samples has not been widely addressed in the subject literature, and yet it is of fundamental significance as regards achieving reliable results.

A perusal of the subject literature reveals that the yield estimated by some authors is very high, of the order of 81–119% [16] and 71–114% [17] for PCBs. However, either these authors do not

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describe precisely the technique they used to add the standard to the sediment samples, or they added the standard in an inappropriate way—to the solvent after extraction [17,18] or directly to the dry sediment [19–22]. The fault with adding standards to the final extract prior to chromatographic analysis lies in the fact that this approach takes no account of the yield of analytes from the sediment or of their loss during the successive sample preparation stages. Again, adding standards directly to the dry sediment does not ensure their proper dispersal within the sample. Standards are added only at certain points in the sediment, which does not reproduce the natural character of the analyte's bonds with the sediment [23].

That is why it is so important to choose the right techniques of adding standards to a sample—one that ensures their proper dispersal and also the reproduction of the natural character of the bonding between analyte and sediment.

This paper describes the results of a study on the effect of standard addition techniques on the results of determination of PCBs in sediment samples (certified reference material: METRANAL™2—river sediment).

2. Experimental

2.1. Reagents and standards

The solvents used during the study were dichloromethane (99.9%), methanol (99.8%) and acetone (99.9%) from Merck (Germany) and pentane (99.8%) from POCH (Poland). Individual solutions of seven selected PCB congeners (IUPAC Nos. 28, 52, 101, 118, 153, 138 and 180) [24–28] were obtained from Restek Corporation (Bellefonte, USA) as 10 µg/ml solutions. The stock solution of PCBs was prepared by mixing solutions (100 µl each) of these compounds. Certified PCB 209 (200 µg/ml in isooctane) standards were obtained from Dr Ehrenstorfer GmbH (Germany). River sediment certified reference material (METRANAL™2, Analytica Ltd.) was purchased from LGC Standards Sp. z o.o. (Poland), with certified concentrations of 15 PCB congeners. Copper powder and silica gel were from J.T. Baker.

2.2. Gas chromatographic analysis

All experiments were performed using a gas chromatograph (TRACE GC), a mass spectrometric detector (TRACE MS) and an on-column injector maintained at 280 °C. The capillary column was a ZB-5MS unit (30 m; 0.25 mm i.d.; 0.25 µm film thickness; 5% phenyl + 95% dimethylpolysiloxane). The carrier gas (helium) was maintained at a constant pressure of 70 kPa. The GC oven temperature was programmed as follows: from 40 to 120 °C at a rate of 40 °C min⁻¹; then at a rate of 5 °C min⁻¹ up to 280 °C, where it was held for 5 min. The MS was operated in electron ionization (EI) mode with the ion source temperature at 220 °C. The mass spectrometer was operated in selected ion monitoring mode; the following ions were monitored: (*m/z*) 256, 258, 290, 292, 324, 326, 358, 360, 392, 394, 496, and 494. An injection volume of 2 µl was selected for all analyses. The interface temperature was maintained at 280 °C.

2.3. Procedure for PCB determination in sediment samples

A ca. 1-g sample of sediment was extracted with 5 ml dichloromethane in shaker for 24 h. The extract obtained was decanted and then evaporated to a volume of 1 ml under a gentle nitrogen stream. Then extract was transferred to SPE columns filled with SiO₂ and activated copper (added to bind sulfur containing compounds). Prior to use freshly activated copper (in 5 ml HCl–water 1:1, v:v) was placed at the column front After column

loading the analytes were eluted with dichloromethane (1 ml/min) and 8-ml fractions were collected. The next stage consisted of the following operations:

- evaporation of a specified extract volume to dryness; extraction with pentane (3 × 100 ml) of the dry residue in an ultrasonic bath;
- fractionation of the pentane extract in glass columns filled with freshly conditioned silica gel (8 h at 140 °C);
- collection of the fraction containing PCBs (8 ml) and evaporating it to dryness under a gentle stream of nitrogen; and
- dissolution of the dry residue in 30 ml of hexane.

Then, 2-µl aliquots of the hexane extract were injected into the chromatographic column, separated and analysed by means of GC–MS. The change in the solvent from dichloromethane to pentane allowed for a preliminary purification of the extract through the primarily separation of polar impurities.

The scheme of the procedure for determining PCBs in sediment samples is given in detail in Wolska [1].

2.4. The effect of standard addition technique on the result of PCB determinations

The aim of the study was to assess the effect of adding a solution containing an internal standard to sediment samples on the results of analyte determinations. Samples of standard were added to 1 g of sediment using three different techniques:

- (1) adding a solution containing the standard to a sediment moistened with acetone and leaving the sample for 24 h to allow the solvent to evaporate;
- (2) adding the standard directly to the extraction solvent;
- (3) adding a solution of the standard directly to the dry sediment,

and then adding 2 µl (variant A) or 20 µl (variant B) of standard solution PCB 209.

2.5. Calculation mode of the analytes amount introduced to the chromatographic column

In order to perform the calculations of the analytes quantity in the sample introduced to the chromatographic column, samples were dosed to the system in the following order:

- first—the standard solution, where the content of analytes and internal standard is known (standard solution contained analytes from PCB group at concentration of 100 ng/ml PCB 209 and 100 ng/ml of PCB mixture);
- second—the investigated sample containing internal standards (internal standard contained of 30 ng/ml of PCB 209).

The quantity of investigated analytes was calculated on the basis of formula presented below [23]:

$$\frac{pY^p/mY^p}{pC^p/mC^p} = \frac{pY^{st}/mY^{st}}{pC^{st}/mC^{st}} \quad (1)$$

where pY^p is the peak area of a determined substance Y on a chromatogram obtained after injecting extract of a sediment sample into the chromatographic system; mY^p the mass of a determined substance Y on a chromatogram obtained by dosing extract of a sediment sample into the chromatographic system; mC^p the mass of a standard C on a chromatogram obtained by dosing extract of a sediment sample into the chromatographic system; pC^p the peak area of an internal standard C on a chromatogram obtained by dosing extract of a sediment sample into the chromatographic system; pY^{st} the peak area of a determined substance Y on a chromatogram

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