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# A novel method for quantification of decabromodiphenyl ether in plastics without sample preparation using direct insertion probemagnetic sector high resolution mass spectrometry

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## **Analytical Methods**

### Paper

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- 2 A novel method for quantification of Decabromodiphenyl ether in
- 3 plastics without sample preparation using Direct Insertion Probe -
- 4 Magnetic Sector High Resolution Mass Spectrometry.
- 5 A. Guzzonato<sup>a</sup>, H. Mehlmann<sup>b</sup>, D. Krumwiede<sup>b</sup>, S.Harrad<sup>a</sup>.

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comparable quality of results.

We report a rapid, efficient analytical method for detecting Brominated Flame Retardants (BFRs) in plastic materials using Direct Insertion Probe with double focusing magnetic sector high resolution mass spectrometry. We acquired the total ion spectrum (30 to 1000 m/z) and observed 959 and 799 m/z ions respectively for quantification and evaluation of the fragmentation reproducibility of decabromodiphenyl ether (BDE209, MW 959 amu). *Ad-hoc* prepared Acrylonitrile Butadiene Styrene (ABS) solid reference materials (RMs) containing different concentrations of BDE209 were used to develop a 5-point calibration curve that showed linearity (R²> 0.999) over a concentration range of 0.1 - 2% w/w BDE209. Relative standard deviation between triplicate determinations of BDE209 ranged from 0.32% to 0.42%. The limit of detection (LOD) obtained for BDE209 was 0.112 mg/kg, 4 orders of magnitude lower than the EU's maximum allowed concentration (MAC) in plastic. To our knowledge, this is the first method for compound specific quantification of BDE209 that does not require any sample preparation, reducing the analysis time from roughly 14 hours to 12 minutes with

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### **Analytical Methods**

#### Paper

#### Introduction

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Brominated flame retardants (BFRs) are a class of chemicals used intentionally to impart flame retardancy in a wide range of consumer goods (such as electrical and electronic equipment (EEE), furniture, and textiles, etc.). They are also found in many more goods and materials as an unintentional result of poor recycling practices (kitchen utensils, children's toys, Food Contact Materials) [1,2]. One major class of BFR are the polybrominated diphenyl ethers (PBDEs). Of the several types of PBDE formulation manufactured and used, the deca-BDE mixture was dominant (83.3% of the global market demand) [3] which to some extent, can debrominate once released into the environment, resulting in a suite of less2 brominated congeners with enhanced toxicity and ability 53 bioaccumulate relative to the parent [3]. The RoHS recase4 Directive 2011/65/EU effective from 2013 sets limit values f 55 polybrominated biphenyls (PBBs) and PBDEs (classes coverify) the most largely used additive BFRs, including deca-BDE) at 5a7 maximum level of 0.1 weight % in homogeneous materia 8 European standards give guidance on how to sample, prepar 29 extract and measure BFRs in plastics [4]. Recommend 600 methods for monitoring compliance with RoHS limits f61 PBDEs can be divided into two main approaches: orientatife2 screening and high-accuracy chemical analysis. Screening 63 preferred for in-situ evaluations as it is usually performed with solid sampling techniques like hand-held X-ray fluorescen65 spectroscopy (XRF) although this can only quantify Br as 666 proxy for the total BFR content, thereby running the risk 67 false positives. More conventional techniques recommended for high-accuracy determination of BFRs life9 PBDEs. Specifically, RoHS requires GC-MS analysis 760 determine the BFR content in styrenic polymers (preceded 51) different sample preparation steps: sub-sample grinding, cry\delta2 grinding, solvent extraction, extract filtration, selectives precipitation for oligomer removal, and chromatograph 14 purification). These traditional techniques can have a numb@5 of drawbacks aside from being time consuming and expensiv 26 Soxhlet or pressurised liquid extraction of plastics often dissolves a substantial fraction of the matrix (polyme78 together with the target compound, rendering the ensuing extract purification laborious and often leading to high 80 variable analyte recoveries. Furthermore PBDEs are prese81 across a wide range of bromination level, from the low 22 brominated tri-BDEs and tetra-BDEs with a low boiling point 🝪 the most brominated (deca-BDE) with a very high boiling point. This makes it practically difficult to use the same GC-MS5

**Table 1** Composition of Reference Materials. Results obtained with Neutron Activation Analysis.

with Neutron Activation Analysis.						
RM	BDE209	BDE209	$Sb_2O_3$	TiO <sub>2</sub>	CaCo <sub>3</sub>	
	w/w %	uncertainty	w/w %	w/w %	w/w %	
		± %				
1	0	0	3	0.7	2.4	
2	0.1	0.007	5	2.4	0	
3	0.5	0.035	0	0.3	3	
4	1	0.07	0.4	3	1.2	
5	2	0.14	2.2	1.2	1.8	

ideally two different GC columns are used, causing several analytical delays (run the samples on one system set-up first, then switch columns and run them again on the second setup), although a better solution was achieved by Ballesteros-Gómez, [5] using GCxGC to resolve coeluting interferences in the second dimension, thus eliminating the need for two different GC runs. The high boiling point and its enhanced susceptibility to degradation and debromination when exposed at the elevated temperatures of the injector, column, ion source and detector make BDE209 a challenging analyte [6-16]. To face these analytical challenges novel methods for the quantification of BFRs have been developed recently combining GC, LC or GCxGC with a soft Ionization source (APCI) and a High Resolution time-of -flight mass spectrometer or for compound-specific screening using Direct Probe with HR-TOF [5]. One attempt to develop a solid sampling, compoundspecific analysis was made by exploring the potential of Direct Analysis in Real Time coupled with Time Of Flight Mass Spectrometry (DART-TOF-MS), [17] but results revealed it as constituting only a qualitative method to screen for the presence of BFRs in environmental matrices.

Against this backdrop, it is evident that a method that combines the convenience of a solid sampling technique with compound specific quantification is highly desirable.

We present here a simple, sensitive and rapid method using Direct Insertion Probe (DIP) in combination with Magnetic Sector High resolution Mass spectrometry (HRMS). This method characterizes target compounds without a chromatographic separation needed, solely via accurate mass determination combined with a traditional library search. To our knowledge, this is the first approach for compound-specific direct analysis of BFRs in polymers that does not require any sample preparation nor a GC or LC inlet. The method is validated via determination of BDE209 in

system set-up to analyze simultaneously in a single GC run: 86

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- L Acrylonitrile Butadiene Styrene (ABS) solid reference materi**ងឹ4**
- 2 (RMs), but accurate mass determination can be applied 55
- 3 unambiguously identify other PBDE congeners.

#### 1 Experimental

#### 5 Overview

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This method involves the use of matrix matched RMs for  $\overset{\smile}{\text{he}}$ 6 compound specific quantification of BFRs in polymers.  $\sqrt[4]{63}$ 7 used ABS as our polymeric matrix as it is one of the most common polymers used in Electrical and Electronic Equipment 10 (EEE) and toys. Our target BFR was BDE209 as it displays a 11 small temperature difference between evaporation and 12 thermal degradation, as a proof of concept for the DIP method 13 that samples the analytes via thermal desorption. Calibration of the method was carried out using RMs at 5 different 15 concentrations of BDE209.

#### **Reference materials**

RMs loaded with different concentrations of deca-BDE were produced by Fachhochshule Muenster Labor füß Instrumentelle Analytik (FMLIA). The method used to produced and test the RMs is described in detail elsewhere [18], but find summary, Br (in the form of deca-BDE) and Sb (in the form of Sb<sub>2</sub>O<sub>3</sub>) were added to an ABS terpolymer melt with the aid of an extruder. Sb<sub>2</sub>O<sub>3</sub> is generally used as a synergist FR in combination with BFRs [19, 20]. A set of five different reference materials was produced in the form of pellets containing different mass fractions of both Br and Sb plus typical fillers commonly used in ABS (see Table 1) in order to best simulate the matrix of the samples.

29 30 Mass fractions of Br in the produced materials were 31 determined (by FMLIA) via Neutron-Activation-Analysis (NAA). The uncertainty of NAA is about 7% (for exact values see Tab 76) 1). To assess macroscopic homogeneity a wavelength 33 dispersive X-ray spectrometer was used with RSD below 2% f97 34 Br. To assess microscopic homogeneity a synchrotron radiation 35  $\mu\text{-XRF}$  (SR  $\mu\text{-XRF})$  was used. The spot size of the exciting beams 36 37 was 200  $\mu$ m, the RSD for Br was 0.7%.

#### Sampling

No sample preparation was required. A very small amou88 ( $\approx 0.045$  mg) was scraped from the pellets of the RMs with 84 scalpel, accurately weighed with a precision scale ( $\pm 0.0005$  mg) and inserted in the aluminium crucibles for the DIP. T86 influence of the scale error on such a small sample is  $\pm 1.1\%$ . 87

#### Instrumentation

The Thermo Scientific™ DFS™ Magnetic Sector High Resolution 191
48 Mass Spectrometer (HRMS) was used for DIP-HRMS analysis 191
49 The probe temperature program is software controlled. The 193
50 Thermo Scientific™ ISQ™ QD Single Quadrupole GC-MS System 193
51 was used for the comparison of mass spectra obtained will 193
52 the most common GC-MS technique (and relative sample 193
53 preparation)[21].

The optimised conditions for BDE209 were obtained by varying one parameter at a time, performing a measurement and observing the influence of this variation on sensitivity, reproducibility and degree of fragmentation of the parent ion: faster DIP temperature ramps and higher electron energies were found to decrease the parent/daughter ratio and to increase the overall signal intensity, therefore a compromise between these two effects was found in the values reported in table 2:

Table 2 DIP-MS conditions	
Ion Source temperature	260 °C
Source mode	El Positive
DIP temperature programme	40(0.5°C/min)-75-400 (2 min)
Scan mode	Magnetic scan
Mass range	30-1000 m/z
Resolution FWHM	20000
Electron energy	46 eV
Emission current	1 mA
Acceleration voltage	4800 V

To set the desired resolution of 20000 FWHM the reference gas (per fluoro kerosene (PFK), indicated in the mass spectrometric determination section) was monitored on mass 792.9499 m/z and the entrance and exit slits were closed recursively until the desired resolution was reached, these parameters were stored in the measurement conditions and the instrument response was regularly checked using the same reference gas.

#### Results and discussion

#### Mass spectrometric determination

As no GC column is used, the only time difference in vaporisation is dictated by the compound's vapour pressure. The RMs we used to test this method were loaded with BDE209, although due to the process they underwent to be produced (melted and extruded several times to ensure homogeneity), some thermal decomposition is likely to have produced a small amount of decomposition products inside the polymer (Fig. 1).

Thermo Scientific™ Mass Frontier Software was used to simulate all the potential BDE209 fragments and hence identify target ions, for this method we chose the molecular ion of decaBDE (m/z 959) and its main breakdown product (m/z 799) octaBDE.

Isotopic patterns and exact masses corresponding to these two ions were simulated using the Thermo Scientific™ Xcalibur™ Software. The exact masses were used to calculate the mass measurement error (ppm) following the expression (1):

$$\Delta\Box_{\Box} = \frac{(\Box_{\Box} - \Box_{\Box})}{\Box_{\Box}} \times 10^6 \qquad (1)$$

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Where  $m_m$  is the measured accurate mass and  $m_c$  is the exall mass. The deviation of the measured masses from the exall masses was for all isotopologues of BDE209 (averaged over **240** scans) less than 1 ppm using a dedicated pre calibrated method based on Reference material PFK which uses all the exact masses contained in the PFK mixture to perform 43 polynomial correction of the measured accurate masses (selection of the measured accurate masses) (selection 2 and Table 3).

Table 3. Mass Table of measured isotopes

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m/z	Relative intensity	Delta (ppm)	Composition
951.1749	4.34	-0.88	C <sub>12</sub> O Br <sub>9</sub> [81]Br
953.1735	19.34	-0.16	C <sub>12</sub> O Br <sub>8</sub> [81]Br <sub>2</sub>
954.1772	2.5	0.24	C <sub>11</sub> [13]C O Br <sub>8</sub> [81]Br <sub>2</sub>
955.1718	51.57	0.16	C <sub>12</sub> O Br <sub>7</sub> [81]Br <sub>3</sub>
956.1756	6.25	-0.41	C <sub>12</sub> [13]C Br <sub>5</sub> [81]Br <sub>5</sub>
957.1696	86.22	-0.01	C <sub>12</sub> O Br <sub>6</sub> [81]Br <sub>4</sub>
958.1725	10.78	-0.46	C <sub>11</sub> [13]C O Br <sub>6</sub> [81]Br <sub>4</sub>
959.1677	100	0.16	C <sub>12</sub> O Br <sub>5</sub> [81]Br <sub>5</sub>
960.1711	12.92	0.22	C <sub>11</sub> [13]C O Br <sub>5</sub> [81]Br <sub>5</sub>
961.1658	81.39	0.38	C <sub>12</sub> O Br <sub>4</sub> [81]Br <sub>6</sub>
962.1689	10.16	0.07	C <sub>11</sub> [13]C O Br <sub>4</sub> [81]Br <sub>6</sub>
963.1634	45	0.00	C <sub>12</sub> O Br <sub>3</sub> [81]Br <sub>7</sub>
964.1666	5.79	-0.13	C <sub>11</sub> [13]C O Br <sub>3</sub> [81]Br <sub>7</sub>
965.1613	16.71	-0.05	C <sub>12</sub> O Br <sub>2</sub> [81]Br <sub>8</sub>
964.1646	1.76	-0.16	C <sub>11</sub> [13]C O Br <sub>2</sub> [81]Br <sub>8</sub>
967.1596	3.52	0.33	C <sub>12</sub> O Br [81]Br <sub>9</sub>

#### Verification of the DIP-HRMS method

evaluated to test the reproducibility of the fragmentation. Although selected ion mode analysis can provide better sensitivity and transient signals that are easier to interpret, we decided to acquire in the complete mass range (m/z 30-1000) for BDE209 for three reasons: (a) when BDE209 is present in consumer goods, whether it is added voluntarily or not, its concentration is usually orders of magnitude higher than the detection limits of the DFS Magnetic Sector GC-HRMS; moreover the regulatory limits set a relatively high concentration threshold 0.1% in homogeneous material; (b) for quantitative purposes it is very important to include in the calculation every fragment (including molecular Br) deriving from the parent ions present in the samples. This approach allowed us to understand whether it was reasonable to assume that - granted a very stable fragmentation yield - the total Br would have been linearly proportional to any of the main fragments produced (octa-BDE and nona-BDE); and (c) such a wide mass range - covering the vast majority of commonly used BFRs - delivers the flexibility to identify and quantitate different compounds simultaneously.

Octa-BDE and deca-BDE were measured and their ratio

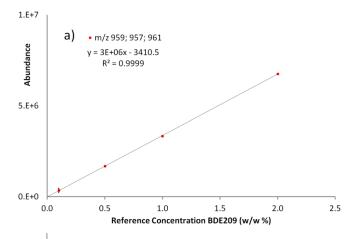
The calibration curve was determined by analyzing each of the five solid RMs (0 %, 0.1 %, 0.5 %, 1 %, 2% w/w of BDE209) in triplicate. Intensities were considered selecting the 3 most

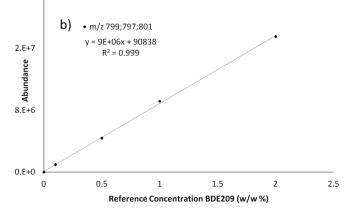
intense m/z values from the isotopic pattern and averaging the intensities of the time signals corresponding to those 3 masses. Scans from the tails of the transient signal were excluded when their relative intensity was less than 5% of the most intense scan (this corresponds to ca. 40 scans for each "peak").

The signal intensity of BDE209 (average intensity between m/z 959, 957, 961) was plotted against the reference value of the RMs. In the same way, the signal intensity of the -2Br fragment (average intensity between m/z 799, 797, 801) was plotted against the reference value. The correlation factor R<sup>2</sup> was >0.999 for both BDE209 and his octabrominated breakdown product, showing linearity over the selected range. The calibration curve for BDE209 was obtained by averaging the signal intensity of the three most abundant isotopologues of BDE209, m/z 959, 957, 961 for each calibration level (Fig. 3a). The calibration curve for the main fragmentation product of BDE209 -which is octaBDE- was obtained by averaging the signal intensity of the three most abundant isotopologues of octaBDE, m/z 799, 797, 801 for each calibration level (Fig. 3b). The LOD was defined as in the ICH1 Guidance (Q2,R1: Validation of Analytical Procedures) as:

$$\Box\Box\Box=3.3\sigma/S$$

where  $\sigma$  = the standard deviation of the response on the triplicate measurement of blank samples (RM BDE209) and S = the slope of the calibration curve. The noise – defined as the





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Fig. 3 Calibration curves and linearity for the analysis of (a) [BDE209]+, (b) [BDE209-57] Br2]+ obtained by DIP. Error bars are the SD between triplicate measurements of ea  $\frac{58}{8}$  RM

intensity of the signal given by the target mass on a bland measurement- was below 1/3 of the instrument detection limits. This result was foreseeable, considering that each sample, and the crucible containing it, was removed from the probe before inserting a new sealed crucible containing different sample, therefore no physical residues of the previous sample could be left on the one following (unlike a traditional GC analysis, where polymeric residues might build up in the injector liner and in the column and create a memory effect). The calculated LOD with this method was 0.112 mg/kg, the LOQ was 1.120 mg/kg for BDE209, slightly lower than a similar study performed with Direct Exposure Probe (DEP) [22]2 and with the advantage of no sample preparation needed.

The memory effect was evaluated by calculating the RSD% between triplicate measurements of the most concentrated RMs: the percent variation was 0.47% and no increasing trend was observed.

Intraday stability was evaluated by performing control runs of RM3 at the beginning, in the middle and at the end of each day of analysis. Over 3 days the intraday RSD of the signal intensity for BDE209 averaged at 1.96%, while interday RSD was 0.51%

The method was applied to 21 real polymeric samples (children's toys and food contact articles, Table 1, SI) for which the BDE209 concentration was measured. BDE209 data and total elemental Br measured with an X-Ray Fluorescence spectrometer were plotted to evaluate if a correlation existed between the two metrics. BDE209 was detected in a concentration ranging from 8.8 mg/kg to 4327 mg/kg. Considering that these data refer to real samples, containing a suite of different BFRs, each potentially contributing to the total elemental Br concentration; the correlation (R²=0.86) between our BDE209 concentration measurements and those for total Br is striking. Moreover, our measurements of BDE209 -which is likely to be a fraction of the total BFR content-never

#### Reproducibility of the fragmentation ratios

exceeded those of total detected Br (Fig.1, SI).

The ionisation behaviour was tested for reproducibility by selecting m/z 799 and m/z 959 from the time signal and measuring the intensity for these masses over the selected time interval. DIP offers a specific advantage with respect to GC-MS analysis: as there is no column or injector between the sample induction system and the ionization volume, it is possible to differentiate between breakdown products (caused by thermal degradation) and ionization fragments (produced by the El ionization process). This is easily done by comparing the time signals for the molecular ion and for its possible moieties as shown in Fig.1: a) and b) show the overlap in intensities of the time signal respectively for the decabrominated ion and the octabrominated ion, meaning that the latter was formed simultaneously in the source, as a fragment of the former. Following this approch we are also

able to say that as the pentabromophenate ion (d)) was detected at the same time of molecular bromine (g)), the debromination happened in the ion source and not as a thermal process in the sample; moreover, time signals d) and g) are both detected before the deca- and octa brominated fragments meaning that their parent ion was already present in the reference material before the insertion in the source.

The ratio between the molecular ion and its main fragmentation product was – for all measured concentrations - 3.1±0.04 (see table 4), showing it to be independent of the sample concentration and suggesting very reproducible fractionation behaviour. This is important as it allows subtraction of the contribution made by the BDE209-2Br fragment to the signal for m/z 799, thereby facilitating quantification of any octa-BDEs present.

A comparison between the mass spectrum of the sample RM obtained using our DIP-HRMS method and that obtained via GC-MS following traditional sample preparation methods and liquid sampling in Fig. 4 shows how the ratio between m/z 799 and m/z 959 is almost two times higher for the traditional GC-MS technique. The thermal decomposition is reduced in the DIP method because the sample is introduced in a chamber under vacuum (instead of under pressure as it would be in a GC injector) therefore the sufficient vapor pressure is reached at lower temperatures.

Table 4. Fragmentation ratio between the two main BDE209 ions.

Reference BDE209 w/w %	Ratio m/z 799 and m/z959	SD
0.1	3.002	0.04
0.5	3.079	
1	3.051	
2	3.002	

#### **Conclusions**

The method reported here represents a rapid, accurate way of performing compound specific quantification of BDE209 in polymers, that avoids completely the labour intensive, time consuming preparation of the samples.

Because of the conveniently small sample size required for our analysis (~0.045 mg), this virtually non-destructive method is designed to be used on articles still in use as domestic appliances (therefore allowing application in studies requiring identification of putative source items in human exposure studies) as well as future waste items. With a linear range covering a concentration span of 19,000 mg/kg which for new and recycled plastics represents the full range of detected concentrations (a considerable improvement with respect to a recent DEP study [22], where the calibration span was from 0.5 to 16 mg/kg) this technique can be a valid, easier, alternative to existing analytical methods for monitoring RoHS compliance in consumer goods. These articles belong now to a second and

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third generation of recycling; and thus generally contain lower 2 concentrations of BFRs (compared to the concentrations of the intentionally added BFRs in older items) as the contaminated 3 polymeric fractions have been mixed with new polymers 4 5 [4][23]. 50 This is illustrated by a recent study where a Direct Injection 6 7 Probe coupled with a HR-TOF-MS was used to screen BFRs plastics, in which the concentration of FRs in the analyzed 8 9 samples never exceeded 1.6% in WEEE items [5]. 10 Our method is tested here for BDE209 in ABS as a proof 56

concept, but given suitable solid RMs, quantification of low 67 brominated compounds in other polymers and over wide calibration ranges will be feasible. DIP-MS optimised for PBDန်ဉ် 13 14 in plastics is able to give results that are as accurate as GC-MS

[24] but are at least 50 times faster to achieve. Considering the 16 burgeoning need for quantification of BFRs in waste samples [IEC 62321], we believe our method will be of significant value 65 17

#### **Future developments**

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This solid sampling technique has the advantage of bei completely solvent-free and hence a "greener" alternative  $\frac{70}{12}$ techniques that involve sample preparation. It would desirable - in order to make it routinely available - to have 78 more complete set of solid reference materials that represent the 3 most widely used FRs (pentaBDE, octaBDE and decaBDD) commercial mixtures) in their respectively most common  $\psi_{\overline{z}}$ used polymers (mainly ABS, PS, and PP/PE). The resources required for production of such suitable reference materials for matrix-matched calibrations with this method would 80 significantly less than the multitude of sample preparations

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- 2013/ under REA grant agreement n° 606857 (ELUTE).

this technique would render redundant.

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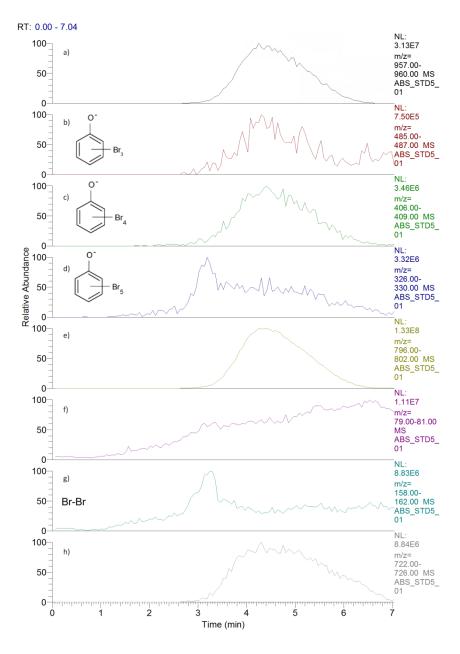


Fig. 1. Time signal obtained with DIP-HRMS (selected over the entire time signal for a) m/z 957-960; b) m/z 485-487; c)m/z 406-409; d) m/z 326-330; e) 796-802; f) m/z 79-81; g) m/z 158-162; h) m/z 722-726. It is possible to notice that the formation of the penta brominated ions (d)) corresponds to a simultaneous release of  $Br_2$  molecules (g)).

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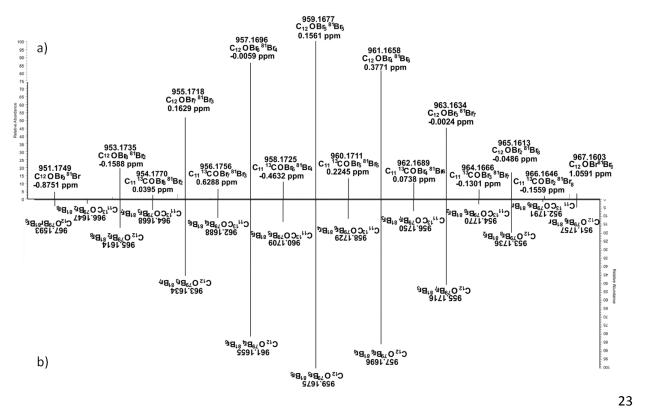
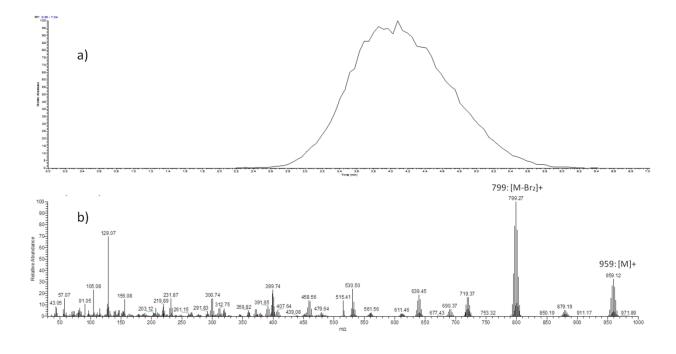


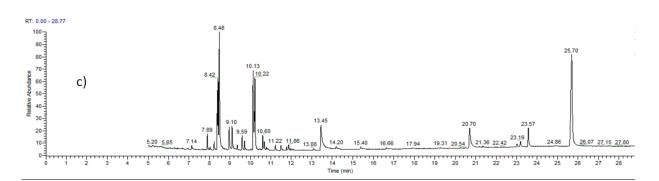
Fig. 2. Comparison of a) accurate masses measured over 20 scans for BDE209 (reported in table 3), and b) their calculated exact value.

1	25 26
2	27
3	28 29 30
4	31
5	32 33 34 35
6	36 37 38
7	38 39 40
8	.0
9	

10

11





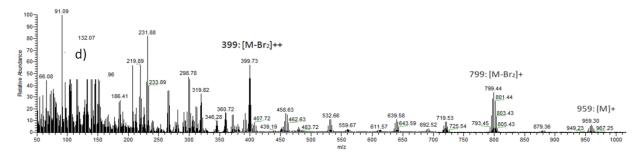


Fig. 4 TIC for the same RM (ABS with 0.5% BDE209) obtained with a)DIP-HRMLA:)GC-MS. Mass spectra obtained with b) DIP-HRMS (selected over the entire time signal for m/z 799; 959), and d)GC-MS (selected over the chromatographic peak RT:25.105 corresponding to BDE209)