

## Introduction

Halogenated organic compounds (HOCs) have been used for decades as pesticides, insecticides and fire retardants. HOCs when mishandled or misused can result in contaminated food and feed These anthropogenic compounds can leach out of supplies. commercial products, enter the environment and cause harm to humans and animals. Their chemical stability and lipid solubility contribute to both their bioaccumulation and biomagnification in the environment. Novel HOCs are continually developed and introduced as replacements for banned or regulated compounds. Figure 1 shows some of the HOCs targeted in this study. They include traditional persistent organic pollutants (POPs) such as polybrominated diphenyl ethers (PBDEs), polychlorinated biphenyls (PCBs), dioxins, and emerging HOCs such as dechlorane plus (syndecabromodiphenylethane (DBDPE), and hexachlorodibromocyclooctane (HBCD).

Analyses of samples containing HOCs can be complicated due to unintentional elimination of targeted compounds (e.g., PBDEs) during sample cleanup, dirty matrices, and mass spectral interferences between coeluting analytes. These problems can be minimized by improving sample preparation and/or enhancing instrument selectivity. LECO Corporation's Pegasus<sup>®</sup> GC-HRT (Figure 2) with Folded Flight Path (FFP™) technology has the increased resolving power, sensitivity and selectivity necessary to analyze pet food samples.







**Figure 2.** LECO's Pegasus<sup>®</sup> HRT and FFP<sup>™</sup> Mass Analyzer.

Dog and cat food samples were spiked with surrogate standards (Accustandard PCBs WHO/NIST/NOAA: 28 PCBs at 10  $\mu$ g/mL) and extracted with acetonitrile using Qsep Q150 extraction salts (Restek #26213). The samples were analyzed right after extraction or treated with dispersive Solid Phase Extraction (dSPE) sorbents before analysis.

### Sample Preparation

1.0 gram of pet food was placed in a 50 mL centrifuge tube, spiked with congener standard (50 or 100 ng standard/ g pet food) and vortexed at 2200 rpm for 30 seconds. Water (9 g) and acetonitrile (10 mL) were added to the food and the mixture was homogenized by vortexing at 2200 for one min. Extraction salts were added, mixed vigorously for one min., vortexed at 2200 for an additional min., and then centrifuged at 3000 rpm for 5 mins. A 1.0 mL aliquot of the acetonitrile layer was spiked with internal standards (30  $\mu$ L of BDE-118 and BB-209, conc. = 50  $\mu$ g/mL) and transferred to an AS vial for analysis. Additional 1.0 mL portions of samples were transferred to dSPE sample clean-up tubes (Restek # 26125 red, 26215 yellow, 26219 orange, 26218 grey), and vortexed for 2 mins., centrifuged at 3000 rpm for 5 mins., and 0.4 mL portions were transferred to AS vials for analysis.

## **Instrument Parameters GC** Parameters

GC Column Typ Injection Oven **Carrier Gas** 

## **MS Parameters**

Spectrom Ion Source: Polarity: Flight Path: **Spectral Ac** m/z Range: m/z Calibro



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## Experimental

Gas Chromatography high performance time-of-flight mass spectrometry (TOFMS) was used for characterization of HOCs in four different varieties of cat and dog food. Spectral data were collected using two of three possible TOFMS operating modes (Figure 3): high resolution (R = 25,000, FWHM at 218.985080 m/z) and ultra-high resolution mode (R = 50,000, FWHM at 218.985080 m/z).

	Agilent 7890
pe:	Restek Rtx-1614 (15m, 0.25 mm ID, 0.18 mm df)
	Splitless, 2 µL , 250 °C
	70 °C (1 min.) to 320 °C at 20 °C/min (1.5 min.)
S	He, Constant Flow (1.00 mL/min.)

eter: LEC	O PEGASUS GC-HRT®
•	EI
	Positive
•	HR, UHR ( $R = 25K$ and 50K, FWHM)
equisition:	6 spectra / second
•	50 – 1000 (HR Mode); 200 – 700 (UHR Mode)
ation:	PFTBA

Quechers extraction of cat food sample A, followed by dSPE cleanup using either Resprep Q213 (150mg MgSO<sub>4</sub>, 25mg PSA, 7.5mg Operation of the GC-HRT in ultra-high resolution mode GCB) or Resprep Q252 (150mg MgSO<sub>4</sub>, 50mg PSA, 50mg C18, 50mg GCB) resulted in the most effective removal of potential interfering matrix components (Figure 3). The high resolving power of the Pegasus HRT is very useful for the analysis of extracted food samples with or without dSPE cleanup as shown in Figure 4. High quality spectral data were obtained for targeted HOCs, as well as, major matrix components such as fatty acids and sterols. Mass spectral data for octadecanoic acid and cholesta-3,5-diene were searched against commercially available libraries (e.g., NIST, Wiley 9) and resulted in matches of 814 and 874 out of a possible 1000. In addition, mass accuracy values for the acid and cholestadiene were 0.16 and 0.008 ppm respectively. All surrogate (28 PCBs) and internal standards (BDE-118 and BB-209) were easily extracted from the data as shown in Figure 5.



Figure 3. TICs -- Cat Food A

A TIC and extracted ion chromatogram (XIC) of cat food sample B is shown in Figure 6. Further evidence of the high quality data produced by the GC-HRT can be seen in Figure 7 where mass spectra for CB-209, BDE-118 and BB-209 are displayed. Mass accuracy values for these HOCs were -0.36, 0.52 and 0.17 ppm. An average of 0.64 ppm was obtained for all HOCs in this sample (Table 1)



**Figure 6.** TIC & XIC – Cat Food B

Figure 8 shows a TIC and XIC for dog food sample A. Five penta-chlorinated biphenyls, one which coelutes with BDE-118 are visible in the XIC of the sample. The mass spectrum in Figure 9 illustrates the GC-HRT's ability for the separation of low abundance isobaric ions with nominal mass 324. BDE-118 mass accuracy values for the molecular ion and [M-Br<sub>2</sub>]<sup>+•</sup> fragment were 0.049 and -0.64 ppm.





# Results (HR Mode)

Figure 4. MS Data – Cat Food A

**Figure 5.** TIC & XIC – Cat Food A

Figure 7. MS Data – Cat Food B

**Table 1.** Mass Accuracies – Cat Food B.



Observed Ion m/z	Mass Accuracy (ppm)
221.99982	0.30
255.96081	0.10
255.96076	-0.09
289.92154	-0.93
289.92158	-0.79
289.92233	1.80
289.92181	-1.18
289.92178	-0.10
323.88297	0.39
323.88336	1.59
323.88303	0.60
323.88242	-1.29
323.88325	1.26
323.88284	0.77
357.84418	0.86
357.84393	0.18
357.84401	0.39
357.84394	0.19
357.84369	-0.48
357.84337	-1.38
357.84387	-0.81
391.80472	-0.45
391.80503	0.34
391.80485	-0.12
391.80498	0.22
425.76592	-1.14
459.72716	0.45
493.68780	-0.36
559.62547	0.52
933.18282	0.17

## System Performance (UHR Mode)

significantly reduces HOC interferences as shown in Figure 10. Robust analysis of the mass spectrum for CB-180 and BDE-47 would be difficult without a high performance mass spectrometer to isobaric fragment ions such as the ones shown in Figure 11. Operation of the GC-HRT in ultra-high resolution mode resulted in a calculated resolution of 23,937 (m/ $\Delta$ m) for the [M-Cl<sub>2</sub>+2]<sup>+•</sup> and [M-Br<sub>2</sub>]<sup>+•</sup> ions of CB-180 and BDE-47. ChromaTOF-HRT software provided deconvoluted spectra for each of the compounds (Figure 11).



Figure 10. MS data for coeluting HOCs – Environmental Sample



Figure 11. Peak True MS Data for Coeluting HOCs – **Environmental Sample** 

### Summary

The data in this poster demonstrates the selectivity and performance of LECO Corporation's Pegasus GC-HRT. The instrument minimizes background interference problems caused by coeluting compounds. The GC-HRT provided the high resolution and enhanced selectivity required for the analysis of complex matrices such as pet food. Dispersive clean-up was not necessary for these samples since targeted and untargeted analysis can be easily accomplished with LECO's Pegasus HRT and ChromaTOF HRT software. The high performance GC-HRT can be successfully employed for analysis of complex matrices.