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Application Note

Combining Pattern Analysis and Cyclic Ion Mobility Mass Spectrometry to Research Perand Poly Fluoroalkyl Substances (PFAS) Exposure in E-waste Handlers

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Abstract

Using the SELECT SERIES[™] Cyclic[™] Ion Mobility System (IMS), data independent analysis (DIA) of human serum of electronic waste (E-waste) handlers has been performed to identify known PFAS and facilitated confirmation of PFAS environmental exposure. In this work we demonstrate the advantages of using the waters_connect[™] Pattern Analysis Application data processing workflow. We identify features that are characteristic of PFAS,

facilitating visualisation of suspected PFAS that are not incorporated in the PFAS library to enable "known" and "unknown" PFAS putative identification.

Benefits

- Streamline detection of unknown PFAS through an integrated workflow using waters_connect Software and pattern analysis
- · Efficiently identify suspected PFAS in complex samples based on distinctive mass spectral properties of PFAS
- Confidently identify "known" and "unknown" PFAS using the multiple identification criteria produced using liquid chromatography coupled with SELECT SERIES Cyclic IMS

Introduction

Stringent guidelines and legislation have been introduced because of the toxicity associated with PFAS exposure.^{1–5} The use of DIA to detect and identify PFAS compounds within the environment or to monitor environmental exposure, is driven by the lack of commercially available standards, which are conventionally used to perform targeted analysis. Adverse health conditions resulting from environmental exposure to PFAS have previously been described.^{6,7} Studies indicate that new alternative PFAS have similar toxicity to banned PFAS, further emphasising the importance of understanding the impact of environmental exposure to both known and unknown PFAS.⁸

Driven by regulation and a demand for alternative PFAS, the synthesis of new PFAS compounds continues and consequently leads to the presence of not yet known PFAS in the environment together with associated by-products and degradation products. The ubiquitous use of PFAS includes the electronics industry, to improve product quality and performance. The recovery of embedded recyclable electronic components is arduous.^{9,10} At recycling centres PFAS exposure occurs through inhalation, ingestion, and dermal routes. Atmospheric transport of particulates extends exposure risk beyond recycling centres to surrounding urban areas.

The advantage of a data independent UHPLC-cIM-MS strategy enables identification of known, unknown, emerging PFAS, PFAS precursors, and transformation products.^{11–13} The highly specific data generated for all detected analytes are analysed using holistic pattern analysis. Using the waters_connect Pattern Analysis Application, PFAS feature patterns are distinguished using information dependent data processing parameters,

these are correlated to chemical properties of analytes possessing fluorine and carbon makeup.

This application note describes the benefits of using the SELECT SERIES Cyclic IMS (see Figure 1) and the waters_connect Pattern Analysis Application to investigate a subset of data acquired from the analysis of the human serum taken from Ghanaian E-waste handlers who may have been exposed to environmental PFAS.



Figure 1. SELECT SERIES Cyclic IMS.

Experimental

Sample Description

E-waste handler anonymised human serum samples.

Anonymised human serum samples were extracted using SPE 96-well µElution plates, containing a polymeric reversed-phase, weak anion exchange mixed-mode sorbent.¹⁴

LC Conditions

| LC system: | ACQUITY $^{\!$ |
|---------------------|--|
| | and Atlantis™ Premier BEH™ C ₁₈ AX Isolator |
| | Column, 2.1 x 50 mm, 5 µm (p/n: 186010926). ¹⁵ |
| | |
| Column: | ACQUITY UPLC [™] BEH C ₁₈ Column (100 mm x 2.1 |
| | mm, 1.8 μm) |
| | |
| Column temperature: | 35 °C |
| | |
| Sample temperature: | 6 °C |
| Injection volume: | 5 |
| injection volume. | 5 με |
| Flow rate: | 0.3 mL/min |
| | |
| Mobile phase A: | 95 H ₂ O (2 mM ammonium acetate):5 MeOH |
| | |
| Mobile phase B: | MeOH (2 mM ammonium acetate) |

| Time (min) | Flow (mL/min) | %A | %В | Curve |
|---------------|------------------|----|----|---------|
| Initial | 0.30 | 95 | 5 | initial |
| 1.0 | 0.30 | 75 | 25 | 6 |
| 6.0 | 0.30 | 50 | 50 | 6 |
| 13.0 | 0.30 | 15 | 85 | 6 |
| 14.0 | 0.30 | 5 | 95 | 6 |
| 17.0 | 0.30 | 5 | 95 | 6 |
| 18.0 | 0.30 | 95 | 5 | 6 |
| 22.0 | 0.30 | 95 | 5 | 6 |

MS Conditions

Ionization mode:

ES-

| Acquisition range: | <i>m/z</i> 50–1200 (HDMS ^E) |
|--------------------------|---|
| Capillary voltage: | 0.5 kV |
| Collision energy: | 20-45 eV |
| Cone voltage: | 10 V |
| Desolvation temperature: | 350 °C |
| Source temperature: | 100 °C |
| Acquisition rate: | 0.16 seconds (HDMS ^E) R~65 |

Data Management

| Chromatography and MS: | MassLynx™ 4.2 SCN1026 Software |
|------------------------|---|
| Informatics: | waters_connect 3.1.0.243 Software |
| | waters_connect Pattern Analysis Application 1.0 |

Results and Discussion

Non-targeted UHPLC-cIM-MS was used to analyse anonymised human serum samples of Ghanaian E-waste handlers and confirmed environmental exposure to PFAS, identifying PFAS including 6:2 FTS, PFHxS, PFHpS, and PFOS isomers (L-PFOS, 5MPHpS, 6MPHpS) with Δ CCS <0.5% compared to the PFAS library values (see Figure 2 (I)). The results showed a Δ CCS of less than 0.5% when compared to PFAS library values, demonstrating how data independent analysis and post-acquisition information dependent data processing can be used to confidently determine environmental exposure to known PFAS.

Although developed countries have implemented stringent guidelines for PFAS, developing countries continue

production and often lack regulations and mechanisms to address emerging PFAS. The United States Environmental Protection Agency (USEPA) maintains a database of approximately 14,800 PFAS analytes.¹⁶ DIA affords opportunity to use a holistic approach to determine exposure to novel and emerging PFAS, in complex matrices such as human serum, soil and food, which can contain 1000's of small molecule analytes within the mass distribution range of PFAS. When performing DIA, precursor ion *m/z*, t_r, and CCS values are acquired for all analytes detected (Figure 2 (II)). Using the acquisition and data processing workflow shown in Figure 3, holistic pattern analysis can be used to characterise information and detect "known unknowns" and "unknown" PFAS within the data, including PFAS for which analytical standards are not available and those that are not present in the PFAS library.

The Pattern Analysis Application derives additional downstream outcomes from the primary data, where criteria are set to perform additional data processing of the "unknowns" data imported from waters_connect Software. Kaufmann proposed "Kaufmann plots" as a single step non-targeted strategy to detect PFAS using a plot of (m/C versus md/C), where mass (m), carbon number (C), mass defect (md) with a two-dimensional graph (m/C versus md/C), to visualise suspected PFAS.¹⁷ Kendrick mass defect is increasingly used in environmental sciences to differentiate natural organic matter, from groups of environmental contaminants and can facilitate selection of potential features to be retained for PFAS suspect screening.^{18,19} Here, we normalise the Kendrick mass defect to the CF₂ repeat unit. As a result, any features differing by this mass form a straight line, aiding the differentiation between natural compounds and anthropogenic PFAS compounds. Additionally, distinctive PFAS mass/CCS trendlines can be used to identify suspected PFAS features.²⁰ Using the waters_connect Pattern Analysis Application, holistic PFAS pattern analysis has been performed using a defined CF₂ repeat unit, designated PFAS repeat unit end groups, component spectrum, Kendrick mass defect, Kaufmann plots (m/C vs md/C), retention time vs m/z plot and m/z vs CCS plot (see Figure 4 (I)). Additional filtering parameters, such as common fragment ion, common neutral loss, CCS, and (CF₂)_n homologous series were applied. The postprocessing and filtering results are presented in Figure 4 (II). The results for a match in the "waters_connect Pattern Analysis Application PFAS library" and an "unknown" PFAS detected in E-waste handler human serum are shown in Figure 5. The waters_connect Pattern Analysis Application PFAS library has been constructed using content of the CompTox Chemical Dashboard vs2.5.2.¹⁶ It contains library hit name, elemental composition and theoretical CCS values generated using machine learning (PFASSTRUCTV5).²¹

A homologous series of unsaturated PFAS anions have been observed (C_4F_7 , C_5F_9 , C_6F_7 , and C_7F_{13}); they diverge from the waters_connect Software PFAS library matches for detected known PFAS, excluding potential false assignment of fragment ions of labile PFAS. Despite the absence of fragmentation information at low intensity responses, a characteristic CCS fingerprint of a labile PFCA species was obtained. By combining

elemental composition, information from the waters_connect Pattern Analysis Application PFAS library, and the observed CCS fingerprint, it is possible to putatively assign the four unknown PFAS species as unsaturated PFCAs, see Figure 6. Barola *et al.* have previously investigated unknown PFAS and putatively assigned unsaturated sulphonic acid, carboxylic and PFAS structures.²² The mechanistic degradation of PFECAs, including Hexafluoro-4-(trifluoromethyl) pent-2-enoic acid (PFMeUPA) have also been investigated.^{23,24} PFMeUPA is used as a fluorinated solvent and cleaning agent in the semiconductor industry. Semiconductor electrical properties are foundational for computers, electronic devices, and everyday electrical appliances; it is possible we have identified a perfluoroalkyl homologous series containing perfluoro-(4-methyl-2-pentenoic acid). Although branched unsaturated PFCAs, such as perfluoro-(4,4-dimethylpent-2-enoic acid), may have been observed, the retention time information from the waters_connect Pattern Analysis Application indicates the observation of an unsaturated PFCA homologous series. Additional confirmation may be demonstrated using concentrated samples or analytical standards.



Figure 2. I) waters_connect Software component summary for "known" identified PFAS.

| Obse | erved m/z | Observed RT (m Observe | ed CCS (Å ²) Res | ponse | Observed drift (ms) | | |
|----------|-----------|------------------------|------------------------------|-------|---------------------|---|----------|
| | 323.2226 | 11.42 | 187.9 | 827 | 3.47 | | |
| | 311.2228 | 11.42 | 179.0 | 7849 | 3.07 | | |
| | 323.2223 | 11.42 | 203.7 | 148 | 4.21 | | |
| 1.2 | 448.9356 | 11.43 | 158.6 | 517 | 2.31 | | |
| | 299.2243 | 11.43 | 180.0 | 103 | 3.10 | | |
| | 267.1970 | 11.43 | 173.5 | 806 | 2.78 | | |
| | 307.1896 | 11.43 | 176.3 | 664 | 2.95 | | |
| | 299.2242 | 11.44 | 174.1 | 99 | 2.84 | | |
| | 439.2095 | 11.44 | 205.0 | 227 | 4.38 | | |
| lobility | Traces 👻 | T | | 1V2 - | | Spectra - 🗼 🔊 🗠 🖽 🗠 🛣 | <u> </u> |
| 500- | | | | | | 25000 | |
| - | | | | | | 293.2124 -312.2261 | |
| 000- | | | | | (* | 100 200 300 400 500 600 700 800 900 1000 11 | 100 1. |
| | | | | | | -171.1033 | |
| 500- | | | | | | 1000- | |
| | | | | | | 201.1145 | |
| | | | | | | 0 | |

Figure 2. II) waters_connect Software component summary for "unknown" detected components.



Figure 3. UHPLC-cIM-MS acquisition and data processing workflow using waters_connect Software and the waters_connect Pattern Analysis Application.

| Pattern Analysis Applica | tion | | | | | | | | | | (2) Hel | p 🗐 Feedb | eck 🏮 Sett | args Uce | nee (j) Alb | out 🔒 | waters | connect A |
|--------------------------|--|----------------|--|---------------------|--|--|--|--|--|---------------|------------------------------|----------------------------|--------------------------|----------------------|--------------------------------|----------------------------|------------|-------------|
| | Summary Calculations | San A | iple: PBVMMCC, | JULY2ND2024HDMSE_06 | 5 Status : Data Pro | ocessed: 22 Assignments | | | | | | | | Q Search | | 8 | 15 | 医侧 |
| Select Data | PBVMMCC_JULY2ND2024HDMSE | | te T | Banked 😤 Target Nar | ie target | Larget Mass Error (ppm) T Ubrary Hit Name | + Library Hit + Mass Beron 7 (mDa) 3 | Library Hit + Mass Error (ppm) 7 | CCS Error (%) 7 | Match 7 Brays | Rt R Library Hi 7 Formula | R Library Hit 7 C/S No. | * Library Hi 7 Adduct | * m/z | e Retention Time 7 (min) | + Drift Tansa T Imsi | ₹ 555 7 | * Intensity |
| | | 5 | Unknown | | | | | | | | | | | 113.09758 | 13.66 | 5.41 | 199.9 | 87 |
| Define Method | | 2 | Unknown | | | | | | | | | | | 114.95671 | 15.32 | 1.01 | 152.7 | 62 |
| | | 3 | Unknown | 0 | | | | | | | | | | 114.93674 | 15.33 | 0.61 | 118.4 | 246 |
| View Results | | 4 | Unknown | 0 | | | | | | | | | | 115.04046 | 14.33 | 7.67 | 286.1 | 82 |
| | | 5 | Unknown | | | | | | | | | | | 115,04046 | 14.32 | 3.32 | 197.4 | 1434 |
| | | 6 | Unknown | | | | | | | | | | | 119.05082 | 4.07 | 0.72 | 122.0 | 58 |
| Multi-plot view | | 7 | Unknown | | | | | | | | | | | 119.05085 | 4.06 | 1.36 | 143.5 | 2621 |
| Sequence Detection | | | Unknown | | | | | | | | | | | 130.94479 | 15.67 | 8.80 | 301.9 | 116 |
| | | 9 | Unknown | | | | | | | D | | | | 130.94503 | 15.97 | 8.92 | 303.8 | 93 |
| | Drif Auto | Mots Defect | 14-00-02-02-02-02-02-02-02-02-02-02-02-02- | 10 40 m/2 Plet | 60 60 70 Mess | 60 60 90 90 90 90 60 0 0 0 0 0 90 60 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 | 1000 1100 Man Defett - | - + 0 | 0.1- 0.05- 0.05- 0.1- -0.15- 0 m/z vs Cl 350- | 20 CS Plot | 80000 800 0 * | • • | 0 0 0 0 0 0 | 0 0 00 120 mrc | 140 | MDA | ło | - 4 |
| | Ideoxinal PRG Carboylic Adda PRG Stationeric Adda Unanageed x:2 Subpenic add seconer | 10 5/m 2 | 00- 00- 00- 00- 00- 00- 00- 00- 00- 00- | | | | ú | đ | 300 - 0 5 5 5 5 5 5 5 5 5 5 5 5 5 | | | | | | | | • 6 | 0 |

Figure 4. I) waters_connect Pattern Analysis Application visualisation of the component summary of unknown components and known PFAS detected using data independent analysis.

| Pattern Analysis Applica | tion | | | | | | | | | | | | | | | ⑦ Help | (ji) Feet | sbeck 🤨 | Settings | Licen | × 0 | About | • | aters_co | nnect |
|--------------------------|---|--|----------------------------------|----------------------------|-----------------|-------------------------------|------------------------------|----------|-------------------------------|--|--------------------------------|-------------------------|--|--------------------------------|---------------------------------|---|--------------------------------|-----------------------------|------------|--------------|-------|-----------------------|---------------|----------|-------|
| | Summery Celculations | Samp * Elle | ile: PBVMMCC | JULY2ND20 | 4HDMSE_06 | 5 Støti | us : Deta | Processe | d: 22 Assigni | ments | | | | | | | | | Q, See | ndh | | | R : | JI 8 | ۹ |
| Select Data | PBVMMCC_JULY2ND2024HDMSE | | * Library Hit 7 CAS No. | * Ubrary Hit 7 Adduct | * m/2 | n Retentio Timo 7 (min) | e e Drift Title T (ms) | , 1B | * Inconsity | C12/C13 Intensity Ratio | * Carbon Estimate 7 (CE) | e m/s: 15 7 ratio | "Charge T | * Common Progment T Tons | SCommon Tragments T Count | Common Neutral T Losses | * Neutral T Losses Count | ⁿ in τ Series | * Assigned | * cri | T CNL | τ ^Φ . T | miner 7 | ct vs | s r |
| | | 3. | | | 180.98991 | 4.26 | 1.00 | 127.4 | 162 | | | | -1 | None Fou | D | None Fou | 0 | 2 | | | | | | | 2 |
| Define Method | | 2 | | | 180.99000 | 4.26 | 0.59 | 113.4 | 3007 | 28.2 | 3 | 60.3 | -1 | None Fou | 0 | None Fou | 0 | | | | | | | 1 🖬 | 4 |
| | | 3 | | | 230.98717 | 6.61 | 0.85 | 120.9 | 1283 | | | | -1 | None Fou | 0 | None Fou | 0 | | | | | | | | 3 |
| | | κ. | | | 230.98720 | 6.62 | 1.26 | 133.7 | 122 | | | | -1 | None Fou. | 0 | None Fou | 0 | | | | | | | | 3 |
| | | | | | 242.98734 | 7.34 | 1.08 | 127.9 | 537 | | | | -1 | None Fou | D | None Fou | 0 | | | | | | | | 2 |
| | | 6 | | | 242.98738 | 6.98 | 1.49 | 140.2 | 463 | | | | -1 | None Fou | D | None Fou | 0 | 5 | | | | | | | 2 |
| Multi-plot View | | 7 | | | 257.22773 | 13.65 | 3.38 | 187.8 | 37273 | 4.5 | 20 | 12.9 | -1 | None Fou | 0 | None Fou | 0 | | | | | | | | 2 |
| | | ÷ | | | 259.24312 | 15.54 | 2.79 | 173.9 | 1457 | 5.3 | 17 | 15.2 | -1 | None Fou | 0 | None Fou | 0 | | | | | | | | 2 |
| Sequence Detection | | 3 | | | 271.22821 | 15.11 | 3.78 | 196.1 | 456 | 5.5 | 16 | 17.0 | -1 | None Fou | 0 | None Fou | 0 | | | | | | | | |
| Compare | | Kenc 0.1 100 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1 | trick Plot | ක ගෙ ග ක ලෙදිව ා | 0 0 0 828000 | 0 0 00 6 M | 00 · · · · | 700 | sóo s ó Mass: - | CF2 CF2 0 1006 Mas | * Tibo | ન વ લ • | m/C 0.0 0.0 0.0 0.0 0.0 0.15 0.15 | AS MD/C Plo | 1 2009 of 20 | . © | 0 | so m/C | 100 | 120 m/C - | | 140 | 160 MD/C - | | |
| | DHP Acids Identified PR45 Suppone Acids PR45 Suppone Acids Unsagenet x3 Suppone Acid suppone | 1204 1004 7/E 504 404 204 | 0- 0- 0- 0- 0- 0- | 0 | 0.0 | 0 0 | 00 | 0 00 | & & | | |] े ब | 350 900 230 200 150 | | 999 8 8 | 600000 88 8 | 9 ⁰ °0 8 8 | | -4- | | | | | | |

Figure 4. II) waters_connect Pattern Analysis Application visualisation of suspected PFAS for investigation, meeting applied information dependent filtering criteria.



Figure 5. Selected "unknown" PFAS with "waters_connect Pattern Analysis Application PFAS library" match.



Figure 6. Putative identification of "unknown" PFAS detected in human serum of E-waste handlers.

Conclusion

Known PFAS have been identified in anonymised human serum of E-waste handlers using highly specific nontargeted UHPLC-cIM-MS data, incorporating retention time, CCS values, and accurate mass measurements. Applying the waters_connect Pattern Analysis Application facilitated a holistic approach to discriminate unknown PFAS related analytes from thousands of matrix components. This approach uses the distinctive mass spectral properties of perfluorinated analytes to disclose suspected PFAS with high certainty. It has enabled the putative assignment of a homologous PFAS series not present in the PFAS library. Identification of unknown PFAS further confirms the advantage of non-targeted analysis, where highly specific data is generated for all analytes detected. UHPLC-cIM-MS and holistic pattern analysis using the waters_connect Pattern Analysis Application streamlines data analysis for monitoring known and unknown PFAS resulting from environmental exposure.

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ACQUITY Premier System <

https://www.waters.com/nextgen/global/products/chromatography/chromatography-systems/acquity-premiersystem.html>

SELECT SERIES Cyclic IMS <https://www.waters.com/nextgen/global/products/mass-spectrometry/massspectrometry-systems/select-series-cyclic-ims.html>

Combining Pattern Analysis and Cyclic Ion Mobility Mass Spectrometry to Research Per-and Poly Fluoroalkyl Substances (PFAS) Exposure in E-waste Handlers

waters_connect Informatics <https://www.waters.com/nextgen/global/products/informatics-andsoftware/waters_connect.html>

MassLynx Mass Spectrometry Software <https://www.waters.com/nextgen/global/products/informatics-andsoftware/mass-spectrometry-software/masslynx-mass-spectrometry-software.html>

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