A Rapid Screening and Quantification Approach to PFAS Analysis Enabled by DART-MS

<u>William L. Fatigante¹</u>; Ronald V. Emmons²; Aghogho A. Olomukoro²; Emanuela Gionfriddo²; Brian D. Musselman¹

Bruker Applied Mass Spectrometry, Billerica, MA¹ Department of Chemistry and Biochemistry, University of Toledo, Toledo, OH²

Introduction

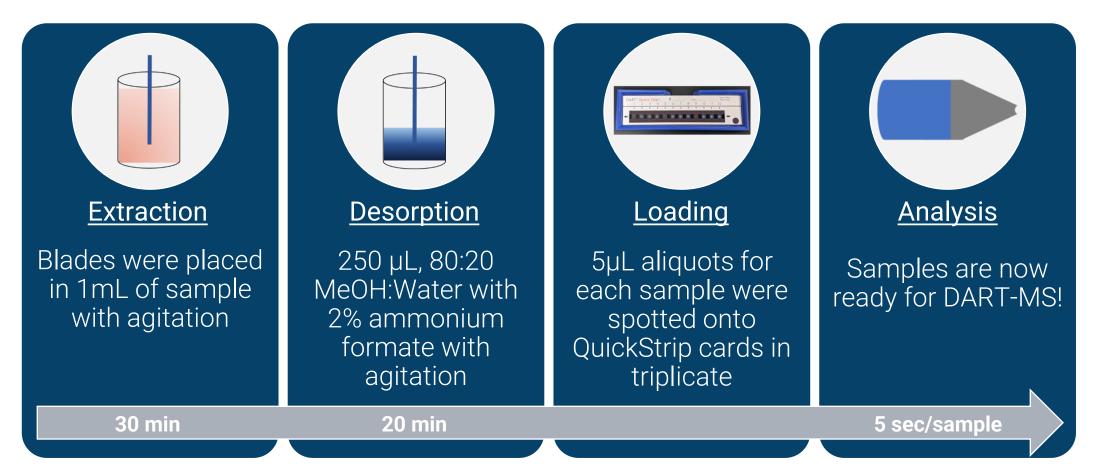
Per- and polyfluoroalkyl substances (PFAS), also known as "forever chemicals," are an emerging class of toxic anthropogenic chemicals. Due to their wide-spread contamination in the environment and human health implications, the development of rapid analytical methods for screening and quantifying these compounds is a must. Published methods require extensive sample preparation and LC-MS/MS protocols for PFAS analysis. In this work, we investigate the potential of rapid PFAS analysis performed by DART-MS coupled to ion-exchange SPME. This was enabled by the use of an HLB-WAX/PAN extraction phase and careful tuning of DART-MS parameters through a central composite design approach.¹





Figure 1. QuickStrip[™] module with JumpShot equipped DART-EVOQ-TQ

Workflow



Methods

SPME devices (fiber and blade geometry) were developed as described by Olomukoro et al.^{2,3} Hydrophilic-lipophilic balance-weak anion-exchange particles embedded in polyacrylonitrile (HLB-WAX/PAN) were used as an extraction phase. PFAS standards (see below) were spiked in ultra-pure water and extracted at a range of 10 – 5000 ppt. DART parameters included negative ionization mode with helium gas, -50 V electric grid and the Vapur interface set at 897 mbar. Plasma heater temperature was set at 250 °C for PFOA and GenX and 400 °C for PFOS and PFBS. Central-composite design (CCD) experiments were performed with a Thermo LTQ XL linear ion trap MS. Calibration and quantification were performed on a Bruker EVOQ Elite TQ-MS.

- ♦ 4 PFAS Perfluorooctanoic (PFOA), compounds: acid Perfluorooctanesulfonic acid (PFOS), Perfluorobutanesulfonic acid (PFBS) and GenX
- ♦ 3 Isotopically labeled internal standards: ${}^{13}C_8$ –PFOA, ${}^{13}C_8$ -PFOS, ¹³C₃ -GenX
- Desorption solution 80:20 (MeOH:H2O, v:v) with 2% ammonium formate

Results

100 GenX

Extracted ion chronogram of PFOA (right) using the optimized SPME-DART-MS method In-source fragmentation (below) is extensive for carboxylic acid and ether moiety PFAS

Fragmentation is plasma temperature dependent, observed with both helium and nitrogen as the plasma

185.02

[M-H-CO2-CF3-CF1]

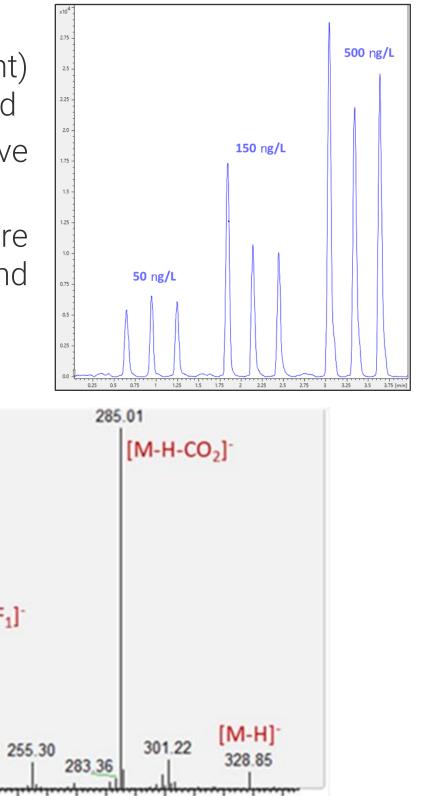
[M-H-CO2-CF3-CF1-O]

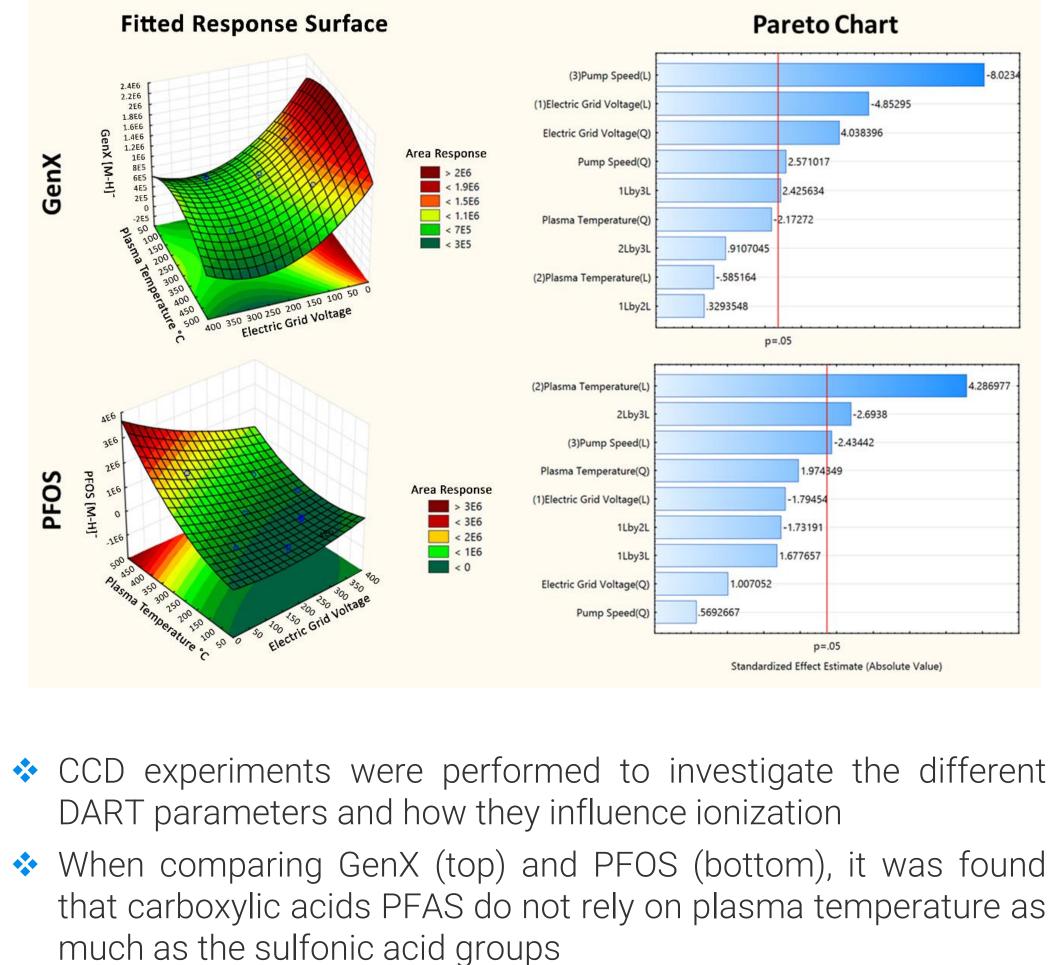
128.03

135.07 151 14

[M-H-CO₂-CF₃-CF₁-O-CF₂]⁻

82.03 97.02





Both analytes favor a lower electric grid voltage (50V) than what is commonly used for other applications (350V)



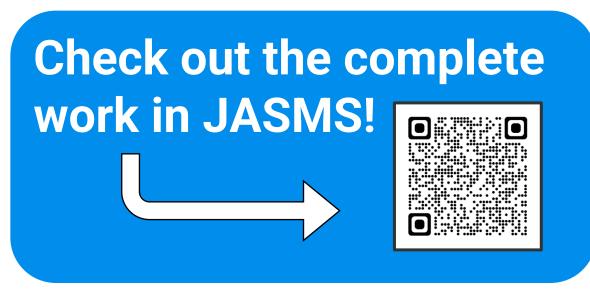
1) Emmons, R. V.; Fatigante, W.; Olomukoro, A. A.; Musselman, B. D.; Gionfriddo, E. J. Am. Soc. Mass Spectrom. 2023.

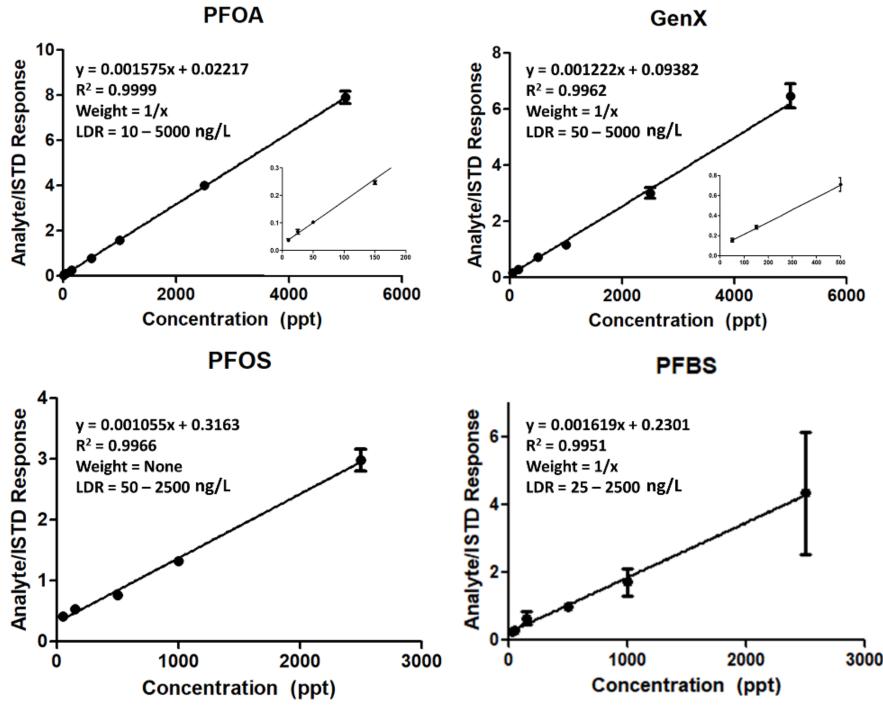
References

2) Olomukoro, A. A.; Emmons, R. V.; Godage, N. H.; Cudjoe, E.; Gionfriddo, E. J. Chromatogr A. 2021, 1651, 462335.

3) Olomukoro, A. A.; DeRosa, C.; Gionfriddo, E. Anal. Chim. Acta. 2023, 1260.

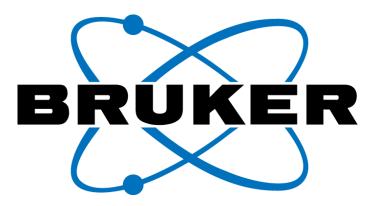
Special thanks to the lab of Emanuela Gionfriddo (University of Toledo) for providing the samples and statistical analysis of results.





Conclusions

- The hyphenation of ion-exchange SPME and DART-MS analysis enabled rapid quantification of PFAS
- When compared to traditional LCMS testing, using DART-MS reduces:
- Method has potential for screening of >1000 samples per hour against large databases with reduced solvent or chromatographic materials



Each concentration was analyzed in triplicate. 12 samples were ionized in ~1.8 min

Area counts of the 4 PFAS analytes were normalized against the corresponding internal standard

✤ PFBS, shown corrected with 13C3 -GenX, was found to require a different internal standard for more robust quantification

- The time required for individual sample analysis
- The cost and amount of organic consumables needed
- By adjusting DART parameters, we can promote greater
- ionization of different PFAS classes (i.e., carboxylic acid v.
- sulfonic acid groups), creating optimal DART methods for each.