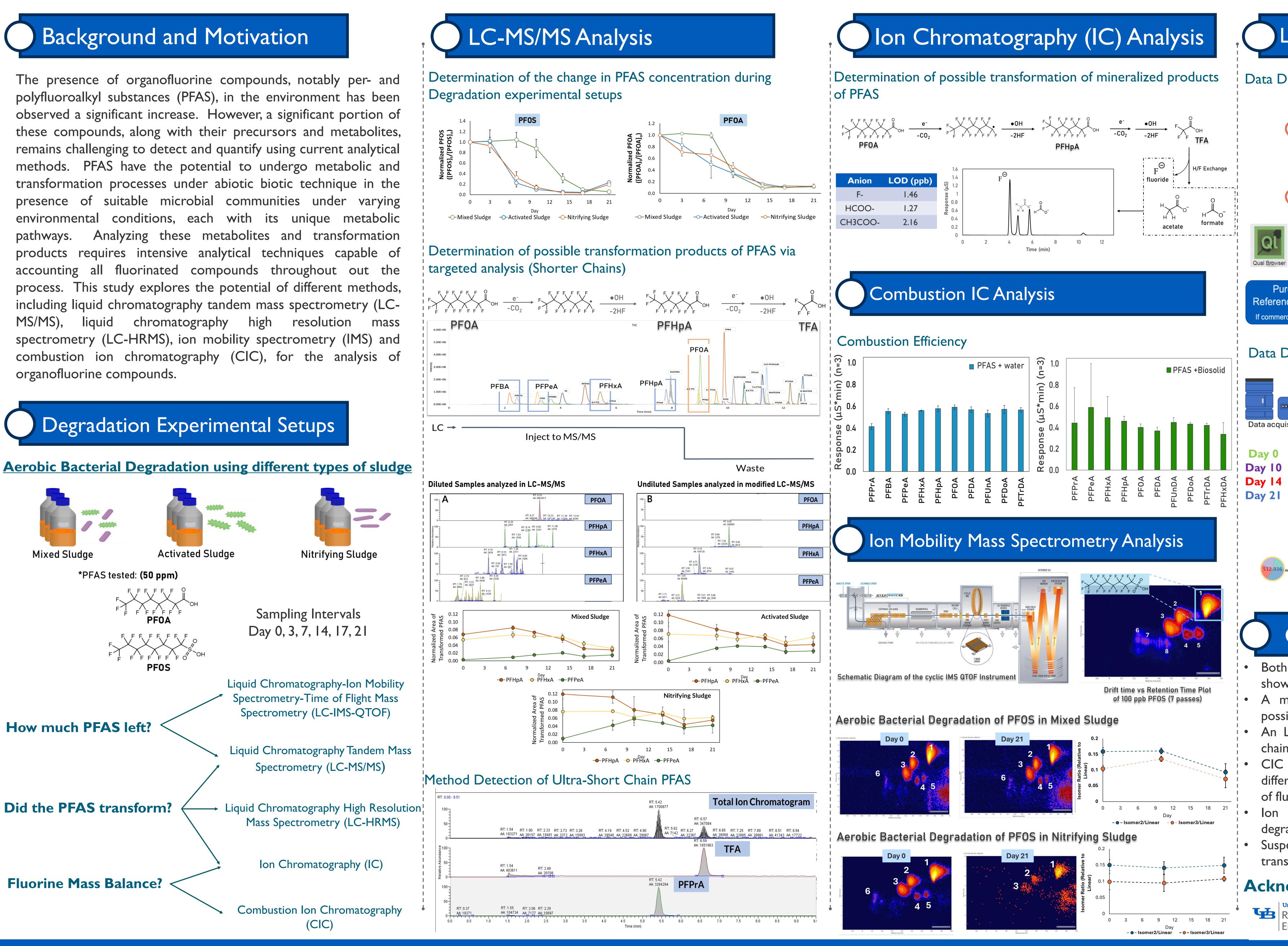
A Comprehensive Analytical Approach and Potential Solutions in Understanding Microbial Transformation of Legacy PFAS in Wastewater

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Analyzing these metabolites and transformation



University at Buffalo The State University of New York

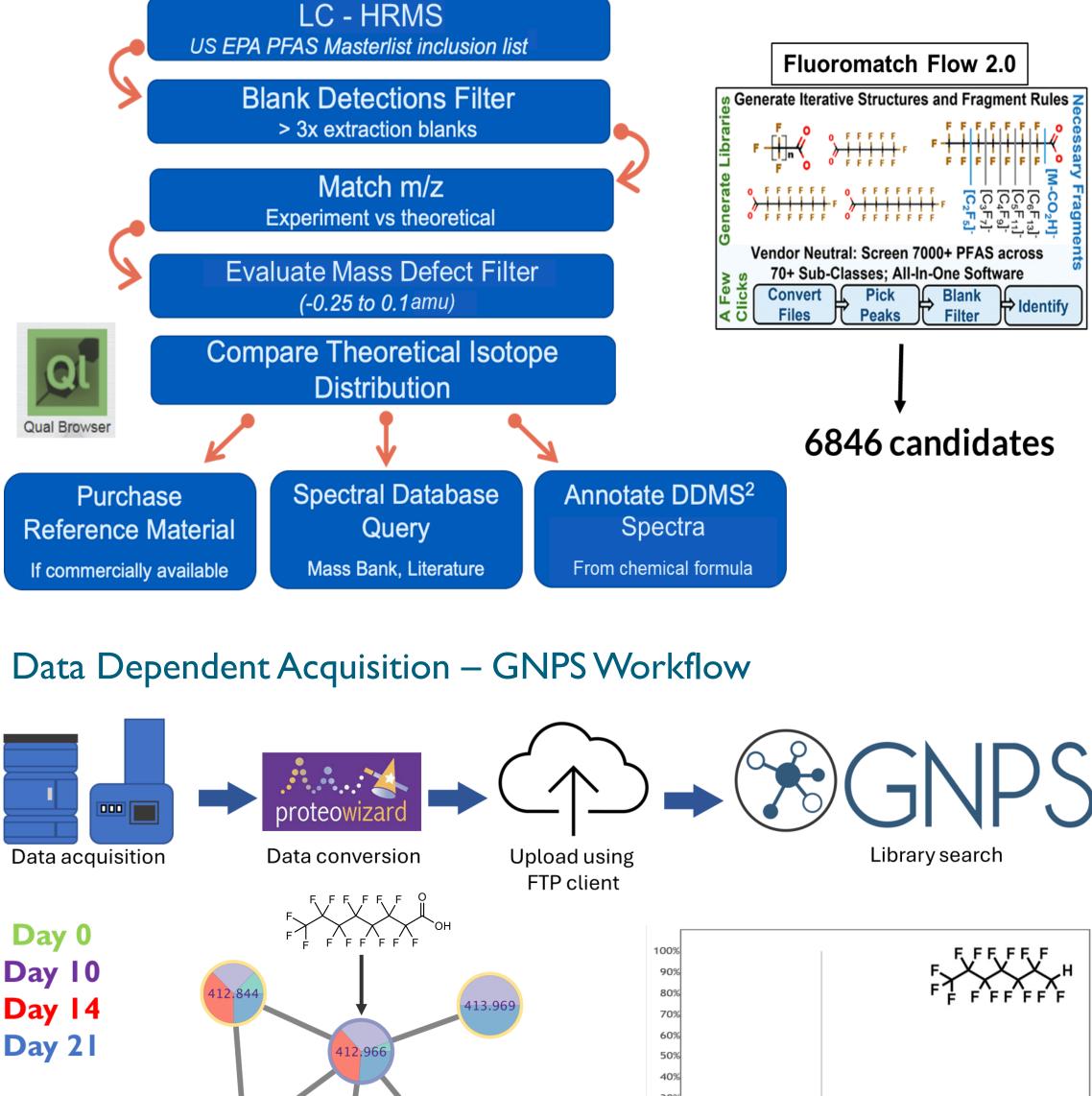
Purchase Reference Materia If commercially available Data acquisition

degradation



LC-HRMS Analysis (Suspect Screening)

Data Dependent Acquisition – Fluoromatch Workflow



Conclusions

Both abiotic and biotic degradation/removal experimental setups showed efficient removal of PFAS.

A modified targeted LC-MS/MS analysis was done to determine possible transformed products of shorter chain homologues.

An LC-MS/MS method was developed to determine the ultra-short chain PFAS (TFA and PFPrA) as possible degradation product.

CIC analysis showed no variability in the combustion efficiencies of different PFAS homologues and can be used to quantify different forms of fluorine that can be used for mass balance approach.

Ion Mobility separation exhibited the regioselectivity of PFOS

Suspect screening using different workflows showed possible transformed products

Acknowledgements

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