

# Determination of PFAS in Biological Tissue by LC-MS/MS

- Using QuEChERS extraction followed with EMR mixed-mode passthrough cleanup

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# PFAS Analysis in Tissue

## Background Introduction

# Determination of PFAS in Biological Tissue

- Environmental monitoring of PFAS residue in wildlife, especially in fish
- EPA Method 1633
- 40 PFAS analytes
- Quantitation based on calibration curve in neat with the use of both EIS and NIS
- Reported LOQ at 0.4 – 0.5 ng/g or higher
- Relatively large and various tolerance window on targets recovery and RSD across the 40 analytes
- Food safety surveillance for PFAS residue in edible fish and meat of terrestrial animals
- AOAC SMPR 2023.003 & EC Regulation 2023/915
- 30 PFAS analytes
- No strict requirements on quantitation method
- Required LOQ at < 0.1 ng/g for core PFAS compounds, and < 1 ng/g for others
- 80-120% recovery and < 20% RSD for core PFAS; and 65-135% recovery and < 25% for others.

Targets: PFAS,

Matrix: biological tissue,

Detection: LC-MS/MS

# EPA Method 1633 Overview for PFAS Determination in Tissue



United States  
Environmental Protection  
Agency

Office of Water

[www.epa.gov](http://www.epa.gov)

December 2024

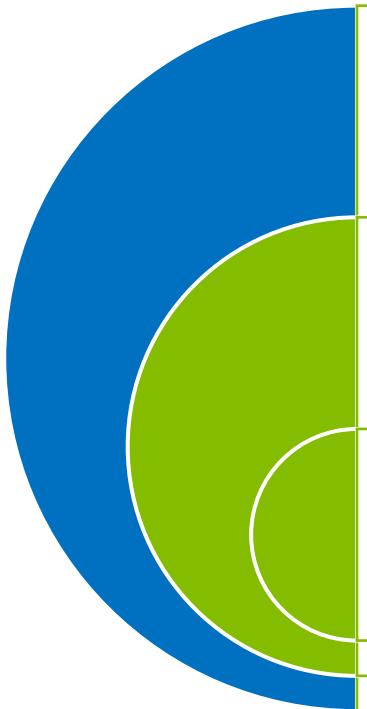
## Method 1633, Revision A

### Analysis of Per- and Polyfluoroalkyl Substances (PFAS) in Aqueous, Solid, Biosolids, and Tissue Samples by LC-MS/MS

- Performance-based EPA method
- Detection 
  - LC-MS/MS detection
- Quantitation 
  - Extraction ISTD (EIS) and non-extracted ISTD (NIS)
  - Calibration curve standards in solvent
- Sample preparation 
  - Three-step solvent extraction using alkaline MeOH and ACN with extensive incubation
  - Carbon/WAX SPE or Carbon dispersive SPE + WAX SPE
  - Long procedure taking > 20 hrs!
  - Multiple steps with high risk of contaminations and deviations
  - Challenging with complex matrices

# Project Objectives

An improved sample preparation method for PFAS analysis in biological tissue



## Improve method performance

- Sensitive and selectivity
- Accuracy (recovery) and precision
- LOQs

## Ease of use

- Simple and fast sample preparation
- Efficient matrix cleanup

## Improve lab productivity

- Saving time, cost, and effort
- Reliable quantitation results



Fish tissue



Poultry and terrestrial animal meat

# PFAS Analysis in Biological Tissue

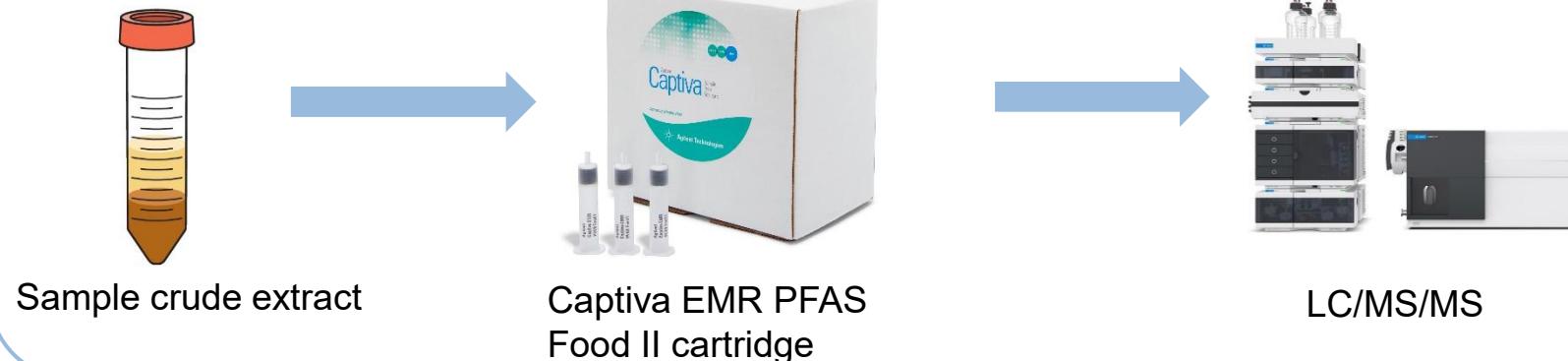
## Sample Preparation

# Sample Preparation Overview

## Sample extraction

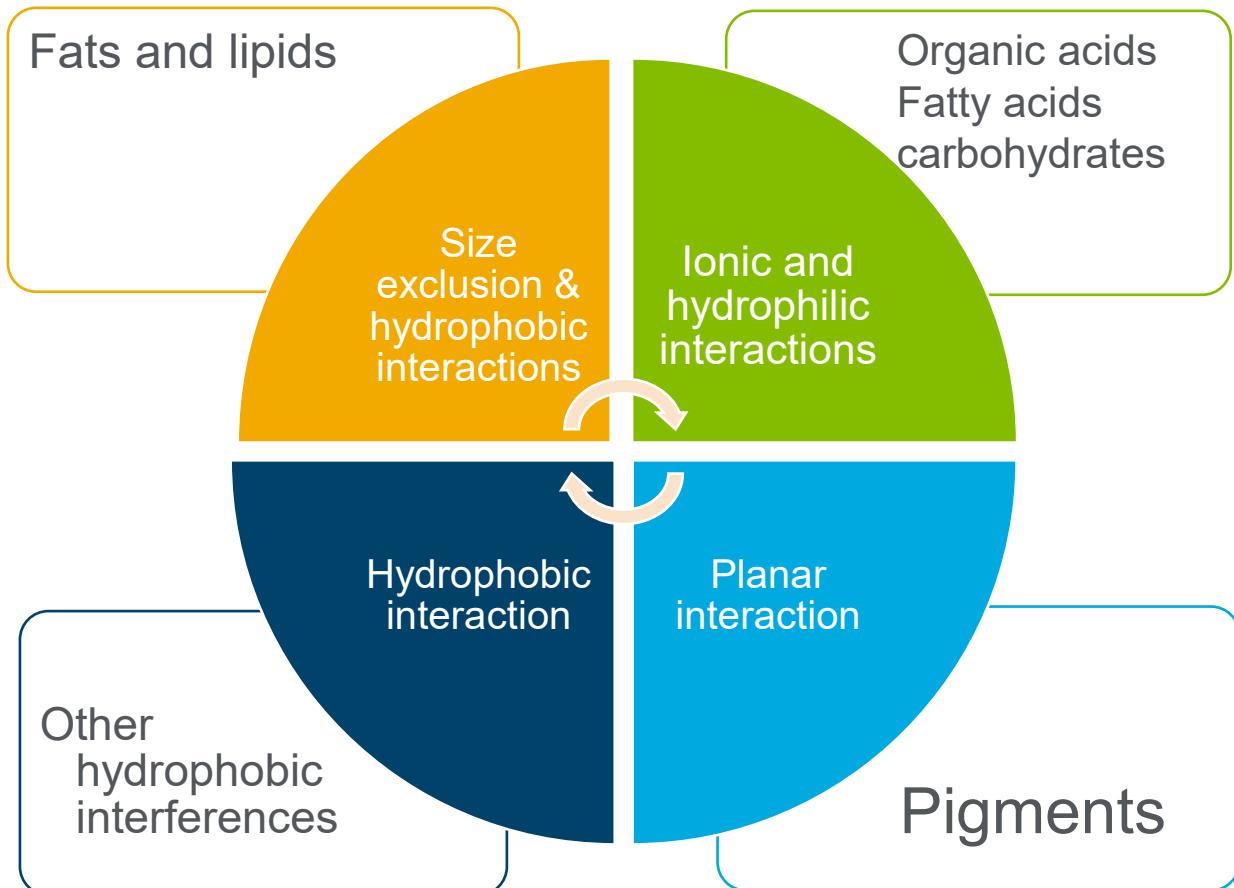


## Sample cleanup



# EMR Mixed-mode Passthrough Cleanup

## Comprehensive matrix co-extractives removal



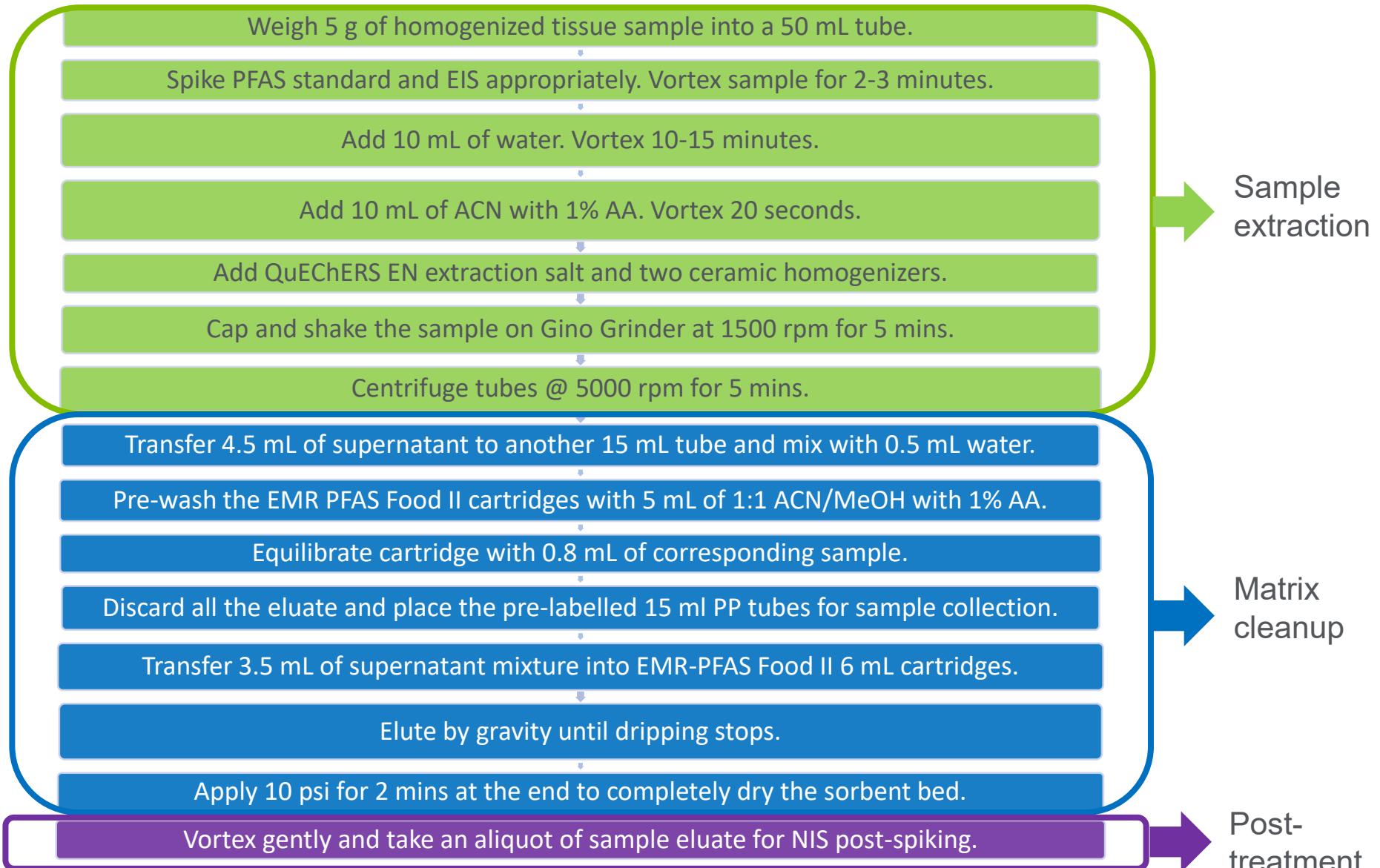
## Method features

- ✓ Matrix co-extractives targeted chemical filtration mechanism
- ✓ Minimal impact on PFAS targets recovery
- ✓ Direct compatible with QuEChERS extraction
- ✓ One step cleanup
- ✓ Comprehensive and efficient matrix removal
- ✓ Higher sample volume (>90%)
- ✓ Saving time and effort

# Sample Preparation Procedure

## Sample prep consumables:

- Bond Elut EN buffered salts and ceramic homogenizers
- Captiva EMR PFAS Food II, 6 mL cartridge
- Polypropylene centrifuge tubes, 50 mL and 15 mL
- Polypropylene vials and caps
- All consumables are either certified or pre-screened for acceptable PFAS cleanliness



# Methods Comparison

| Method            | Novel sample prep method     | Traditional EPA 1633 sample prep method   |   |
|-------------------|------------------------------|---|---|
|                   | QuEChERS <sub>ext</sub> -EMR | Solvent <sub>ext</sub> -Carbon/WAX SPE  | Solvent <sub>ext</sub> -Carbon dSPE-WAX SPE   |
| Pre-work          | Make 2 reagents              | <ul style="list-style-type: none"> <li>• Make 6 reagents</li> <li>• Pack glass wool in SPE cartridge</li> </ul>             | <ul style="list-style-type: none"> <li>• Make 6 reagents</li> <li>• Pack glass wool in SPE cartridge</li> </ul>                               |
| Sample extraction | One-step QuEChERS extraction | Three-step solvent extraction   | Three-step solvent extraction   |
| Transition step   | Dilution with 10% water      | <ul style="list-style-type: none"> <li>• Drying and redissolving</li> <li>• pH check and adjustment</li> </ul>              | <ul style="list-style-type: none"> <li>• Carbon dSPE cleanup</li> <li>• Drying and redissolving</li> <li>• pH check and adjustment</li> </ul> |
| Matrix cleanup    | EMR passthrough cleanup      | Carbon/WAX SPE extraction and cleanup   | WAX SPE extraction and cleanup  |
| Post treatment    | NIS post-spiking             | <ul style="list-style-type: none"> <li>• Sample neutralization</li> <li>• NIS post-spiking</li> <li>• Filtration</li> </ul> | <ul style="list-style-type: none"> <li>• Sample neutralization</li> <li>• NIS post-spiking</li> <li>• Filtration</li> </ul>                   |
| Total time        | 2-4 hours                    | > 20 hours  |   |
| Total cost        | Low with >50% cost saving    | High  |   |

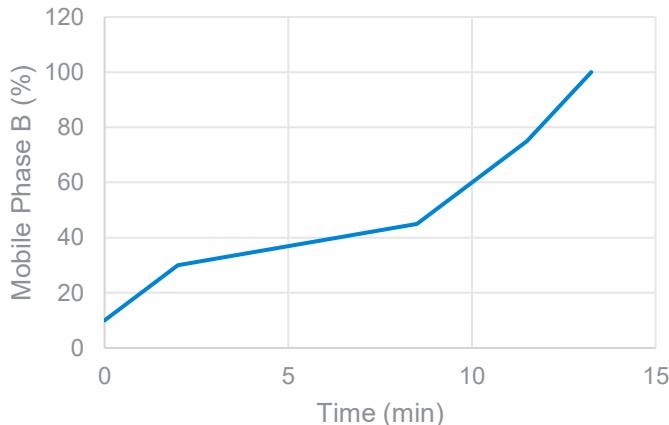
# PFAS Analysis in Biological Tissue

LC/MS/MS Instrument Detection and Quantitation

# Instrumental Analysis by LC-MS/MS

| LC method parameters                  |  | MS QQQ Parameters |                     |
|---------------------------------------|--|-------------------|---------------------|
| <b>Solvent A</b>                      | 5 mM ammonium acetate in water   | Ion Source        | ESI                 |
| <b>Solvent B</b>                      | ACN  | Acquisition       | dMRM                |
| <b>Flow</b>                           | 0.4 mL/min   | Polarity          | Negative            |
| <b>Pump Program</b>                   | $T_{0.0}$  | 10% B             |                     |
|                                       | $T_{2.0}$  | 30% B             | Source Parameters   |
|                                       | $T_{8.5}$  | 45% B             | Gas Temp 200 °C     |
|                                       | $T_{11.5}$   | 75% B             | Gas Flow 18 L/min   |
|                                       | $T_{13.25}$  | 100% B            | Nebulizer 15 psi    |
| <b>Stop Time</b>                      | 15.5 min   | Sheath Gas Heater | 300 °C              |
| <b>Post Time</b>                      | 2 min  | Sheath Gas Flow   | 11 L\min            |
| <b>Injection Volume &amp; program</b> | 5 $\mu$ L<br>15 $\mu$ L water + 5 $\mu$ L sample + 10 $\mu$ L water + 10 $\mu$ L air | Capillary         | 2500 V (-), 0 V (+) |
| <b>Needle Wash</b>                    | Multi-wash program using<br>1. IPA; 2. ACN; 3. $H_2O$                                | Nozzle Voltage    | 0 V                 |
| <b>Analytical LC column</b>           | RRHD Eclipse Plus C18, 1.8 $\mu$ m, 2.1 x 100 mm                                     |                   |                     |
| <b>Guard column</b>                   | Eclipse Plus C18, 1.8 $\mu$ m, 2.1 x 5 mm  |                   |                     |
| <b>Delay column</b>                   | InfinityLab PFC delay column, 4.6 x 30 mm  |                   |                     |
| <b>Column temperature</b>             | 55°C   |                   |                     |

LC Gradient



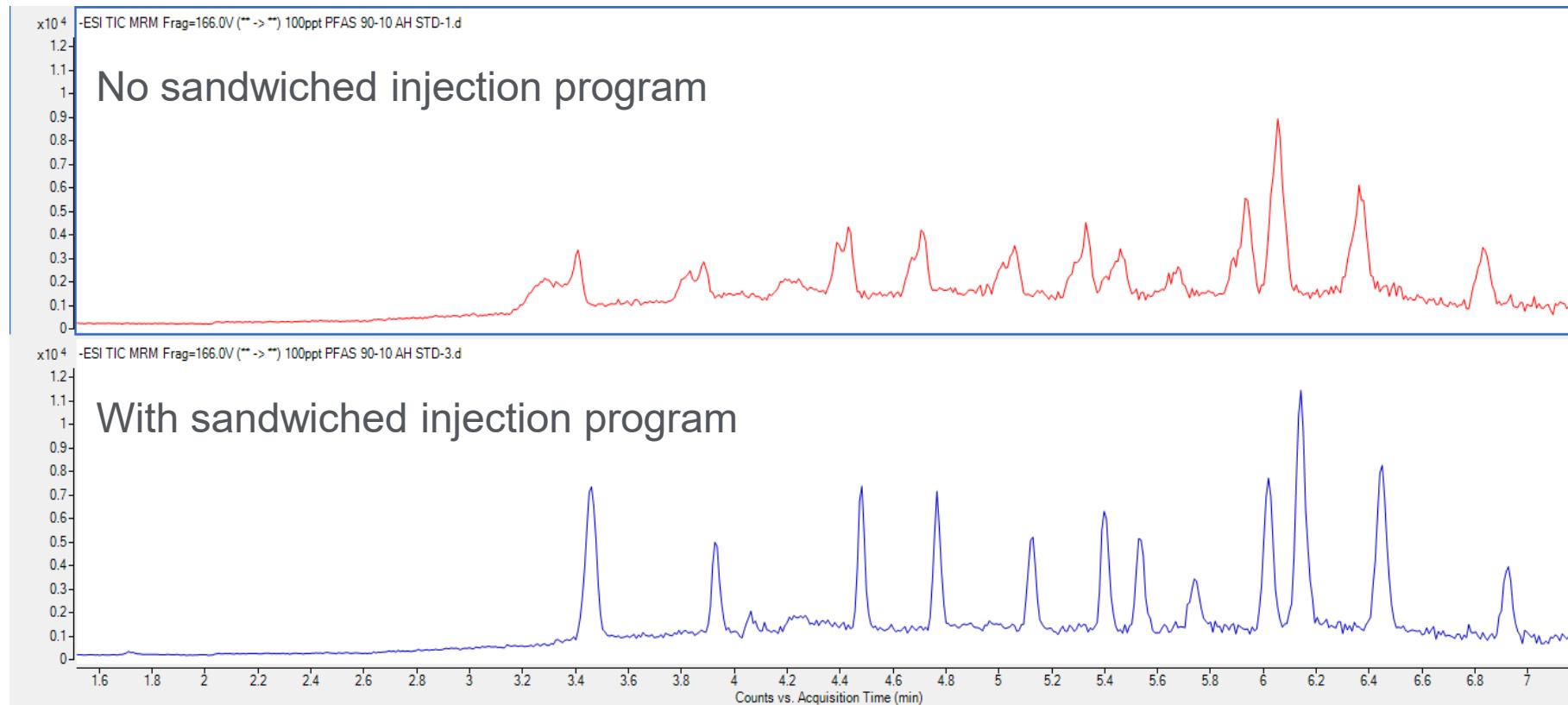
Agilent Triple quadrupole LC/MS system, 6495D mass spectrometer

# Sandwich Injection

- Injection plug sandwiched between low strength solvent plugs

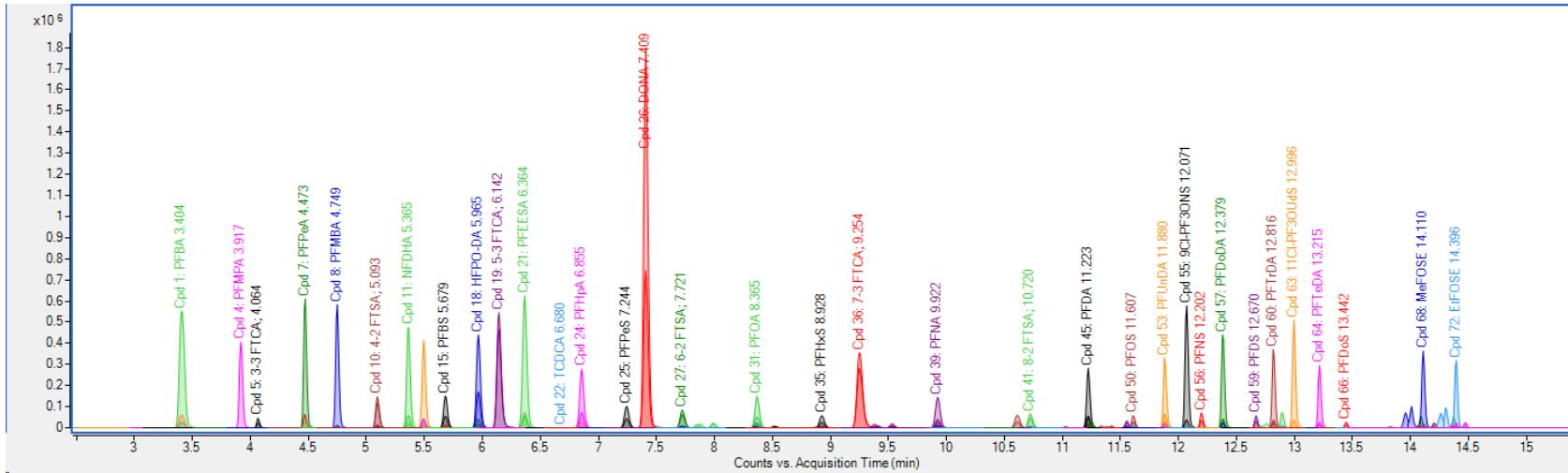


- Enable the injection of sample eluent after EMR cleanup directly

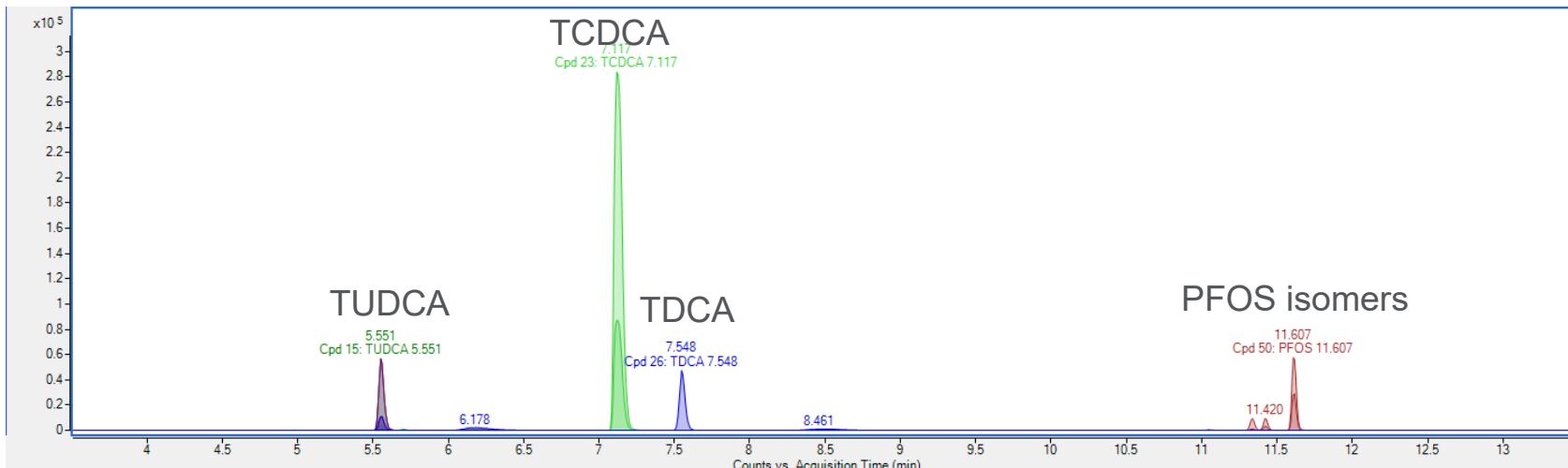


# Chromatographic Separation

PFAS analytes, EIS and NIS chromatographic separation and distribution



Baseline separation of PFOS and isobaric cholic acids



# Quantitation

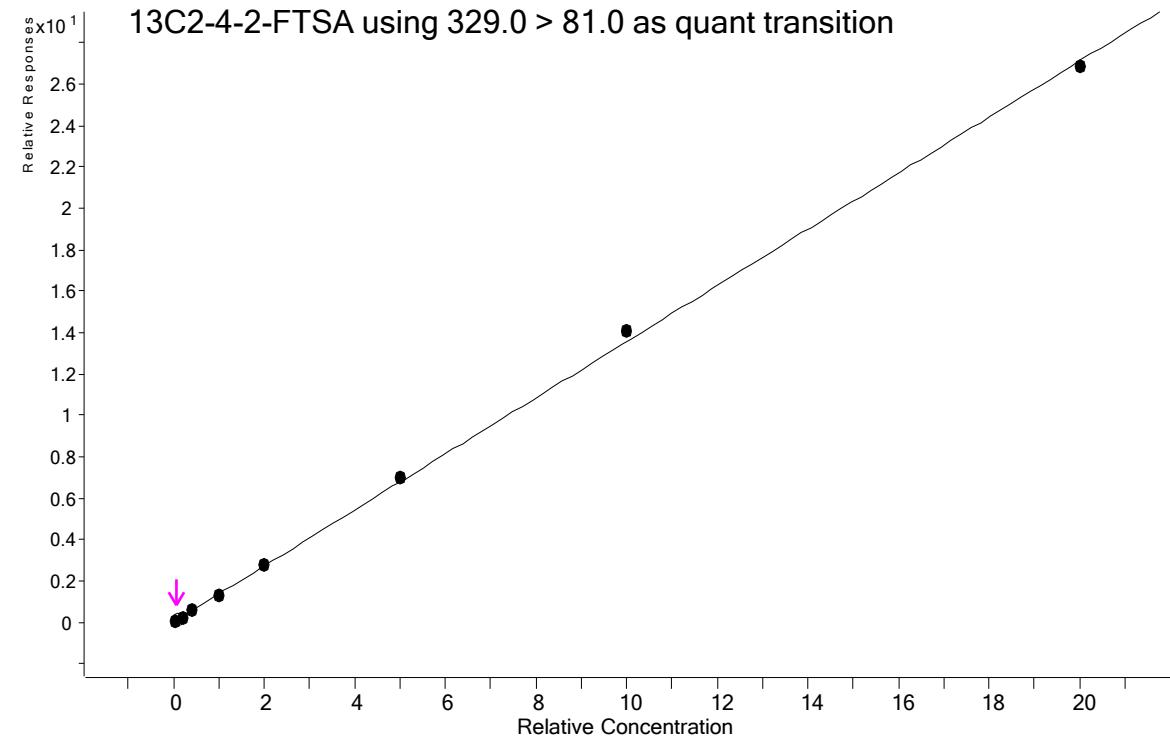
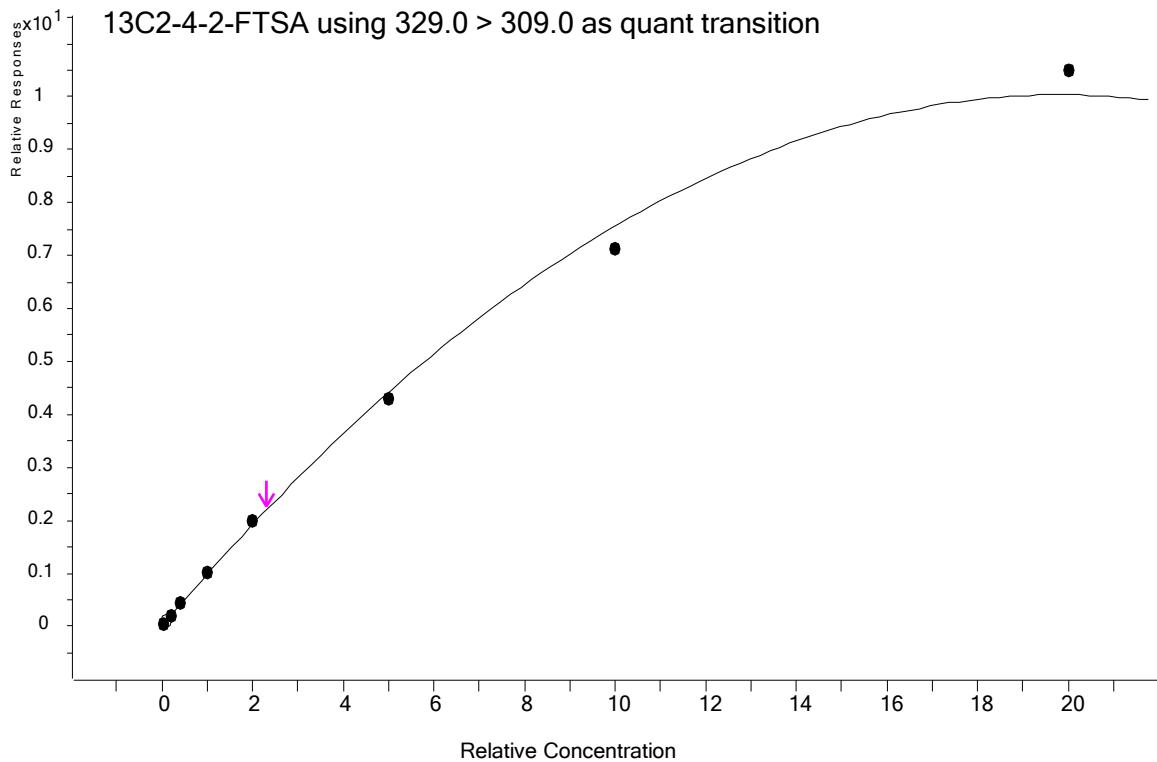
- Calibration standards made in solvent (ACN)
- 400-fold dynamic range
- PFAS analytes quantitation based on the ratio of PFAS analytes and EIS compounds
- EIS assignment was in alignment with EPA Method 1633, with few exceptions.
- Linear regression,  $1/x^2$  weight.
- $R^2 > 0.99$  across all analytes' calibration curves.

| Target        | Assigned EIS                     | Cal. range (ng/g) | Target            | Assigned EIS                     | Cal. range (ng/g) |
|---------------|----------------------------------|-------------------|-------------------|----------------------------------|-------------------|
| PFBA          | $^{13}\text{C}_4\text{-PFBA}$    | 0.2 - 80          | PFHpS             | $^{13}\text{C}_9\text{-PFNA}$    | 0.05 - 20         |
| PFMPA         | $^{13}\text{C}_4\text{-PFBA}$    | 0.1 - 40          | 8:2 FTS           | $^{13}\text{C}_2\text{-8:2 FTS}$ | 0.2 - 80          |
| 3:3 FTCA      | $^{13}\text{C}_5\text{-PFPeA}$   | 0.25 - 100        | PFDA              | $^{13}\text{C}_6\text{-PFDA}$    | 0.05 - 20         |
| PFPeA         | $^{13}\text{C}_5\text{-PFPeA}$   | 0.1 - 40          | N-MeFOSAA isomers | D <sub>3</sub> -N-MeFOSAA        | 0.05 - 20         |
| PFMBA         | $^{13}\text{C}_5\text{-PFPeA}$   | 0.1 - 40          | N-EtFOSAA isomers | D <sub>5</sub> -N-EtFOSAA        | 0.05 - 20         |
| 4:2 FTS       | $^{13}\text{C}_2\text{-4:2 FTS}$ | 0.2 - 80          | PFOS isomers      | $^{13}\text{C}_8\text{-PFOS}$    | 0.05 - 20         |
| NFDHA         | $^{13}\text{C}_5\text{-PFHxA}$   | 0.1 - 40          | PFUnA             | $^{13}\text{C}_7\text{-PFUdA}$   | 0.05 - 20         |
| PFHxA         | $^{13}\text{C}_5\text{-PFHxA}$   | 0.05 - 20         | 9CI-PF3ONS        | $^{13}\text{C}_7\text{-PFUdA}$   | 0.2 - 80          |
| PFBS          | $^{13}\text{C}_3\text{-PFBS}$    | 0.05 - 20         | PFNS              | $^{13}\text{C}_7\text{-PFUdA}$   | 0.05 - 20         |
| HFPO-DA       | $^{13}\text{C}_2\text{-HFPO-DA}$ | 0.2 - 80          | PFDoA             | $^{13}\text{C}_2\text{-PFDoA}$   | 0.05 - 20         |
| 5:3 FTCA      | $^{13}\text{C}_4\text{-PFHpA}$   | 1.25 - 500        | PFDS              | $^{13}\text{C}_2\text{-PFDoA}$   | 0.05 - 20         |
| PFEESA        | $^{13}\text{C}_4\text{-PFHpA}$   | 0.1 - 40          | PFTrDA            | $^{13}\text{C}_2\text{-PFDoA}$   | 0.05 - 20         |
| PFHpA         | $^{13}\text{C}_4\text{-PFHpA}$   | 0.05 - 20         | PFOSA isomers     | $^{13}\text{C}_8\text{-PFOSA}$   | 0.05 - 20         |
| PFPeS         | $^{13}\text{C}_4\text{-PFHpA}$   | 0.05 - 20         | 11CI-PF3OUdS      | $^{13}\text{C}_8\text{-PFOS}$    | 0.2 - 80          |
| ADONA         | $^{13}\text{C}_8\text{-PFOA}$    | 0.2 - 80          | PFTeDA            | $^{13}\text{C}_2\text{-PFTeDA}$  | 0.05 - 20         |
| 6:2 FTS       | $^{13}\text{C}_2\text{-6:2 FTS}$ | 0.2 - 80          | PFDoS             | $^{13}\text{C}_8\text{-PFOS}$    | 0.05 - 20         |
| PFOA isomers  | $^{13}\text{C}_8\text{-PFOA}$    | 0.05 - 20         | N-MeFOSE isomers  | D <sub>7</sub> -N-MeFOSE         | 0.5 - 200         |
| PFHxS isomers | $^{13}\text{C}_3\text{-PFHxS}$   | 0.05 - 20         | N-MeFOSA isomers  | D <sub>3</sub> -N-MeFOSA         | 0.05 - 20         |
| 7:3 FTCA      | $^{13}\text{C}_3\text{-PFHxS}$   | 1.25 - 500        | N-EtFOSE isomers  | D <sub>9</sub> -N-EtFOSE         | 0.5 - 200         |
| PFNA isomers  | $^{13}\text{C}_9\text{-PFNA}$    | 0.05 - 20         | N-EtFOSA isomers  | D <sub>5</sub> -N-EtFOSA         | 0.05 - 20         |

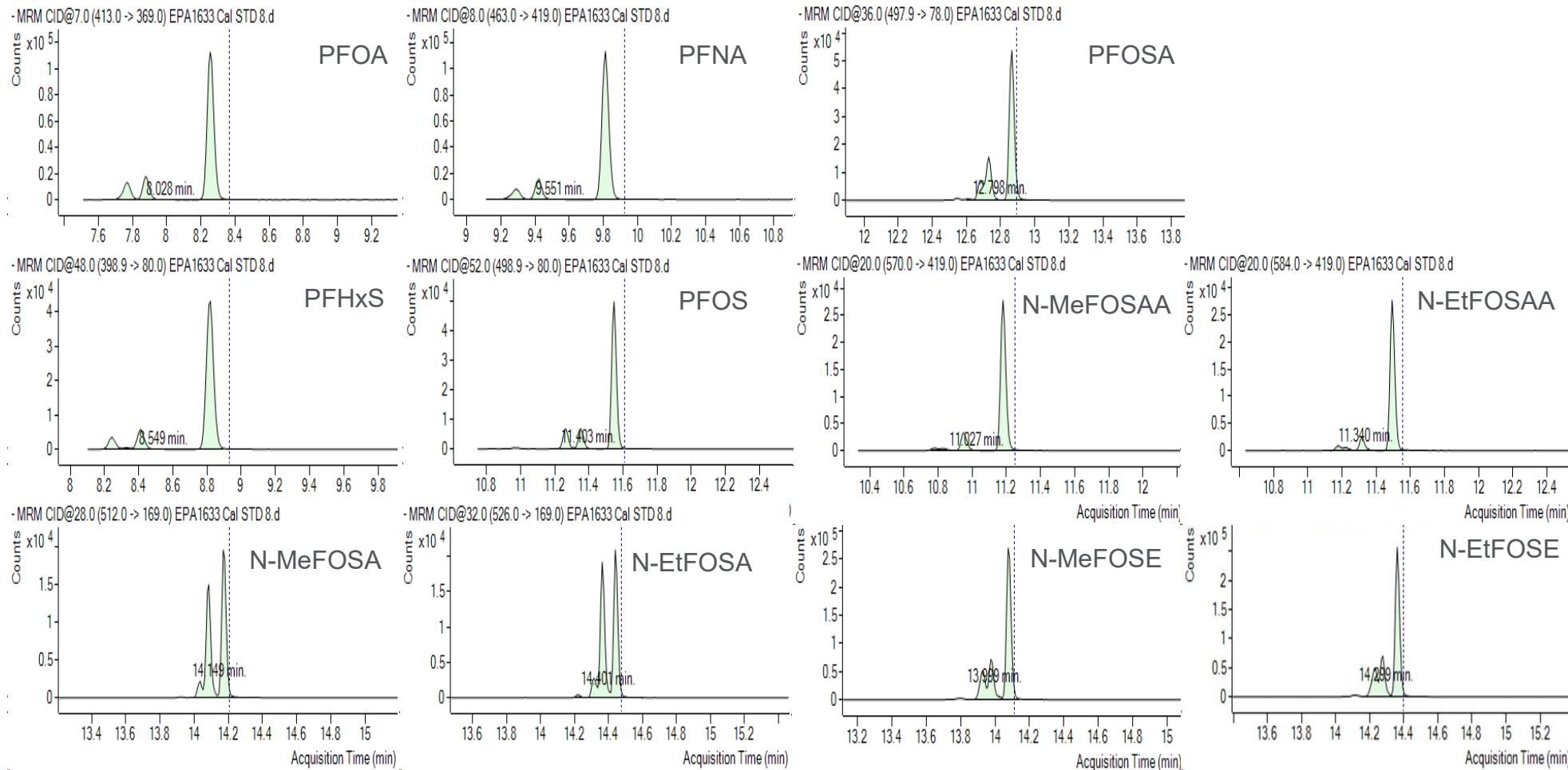
# FTS Targets Calibration

- Isotopic labelled FTS EIS compounds used the less abundant transition with product 80 or 81.
- Applied to all FTS target analytes
- Linear calibration curves were achieved.

Example: 4:2 FTS calibration curves



# Quantitation for Targets with Isomers



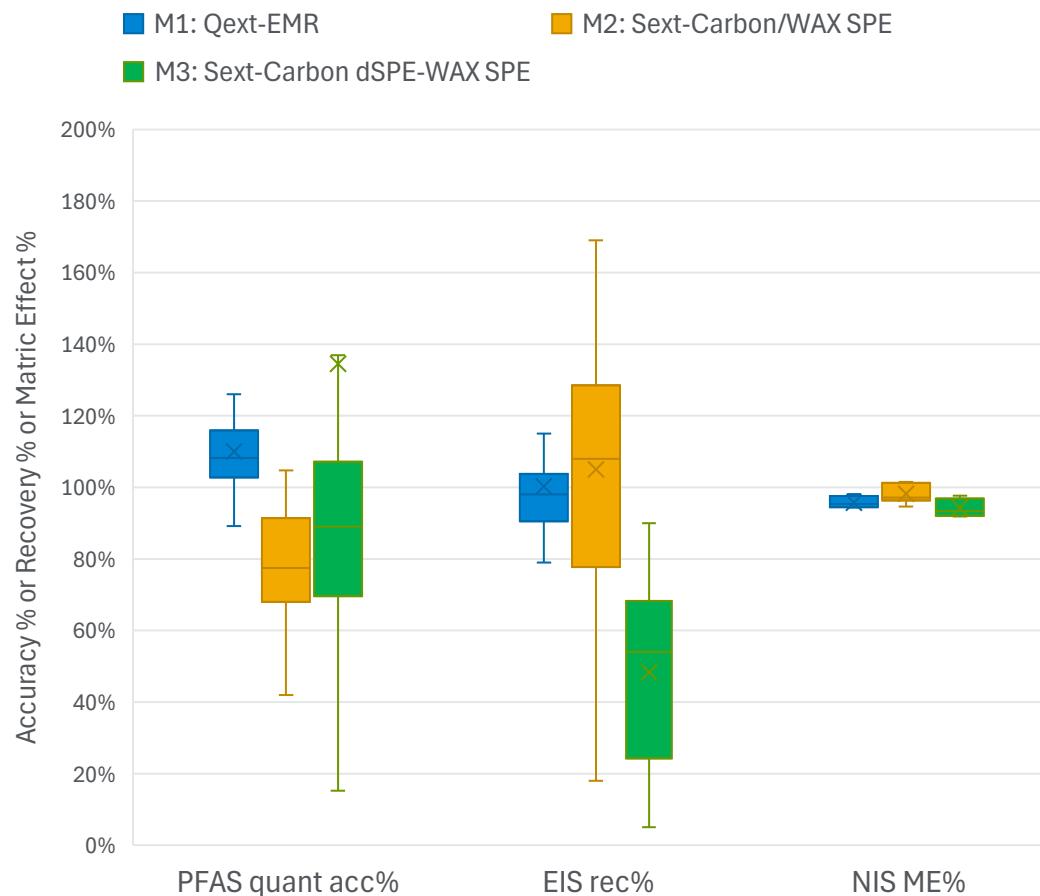
All PFAS analytes with isomers were based on summated integration of all isomers for quantitation.

# PFAS Analysis in Biologic Tissue

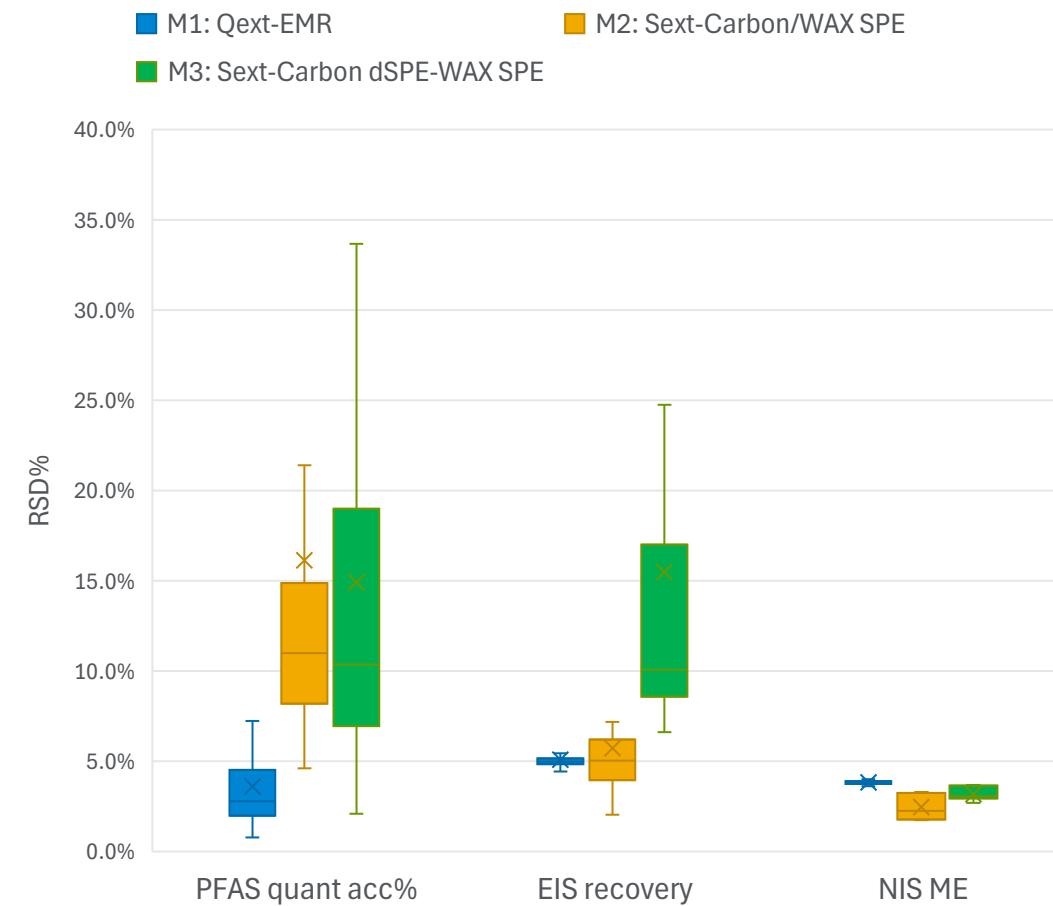
Comparison and Method Final Validation

# Methods Comparison – Quantitation Performance

**A) Method Comparison for PFAS in Tissue Quantitative Analysis**  
- Targets accuracy, EIS recovery, NIS matrix effect

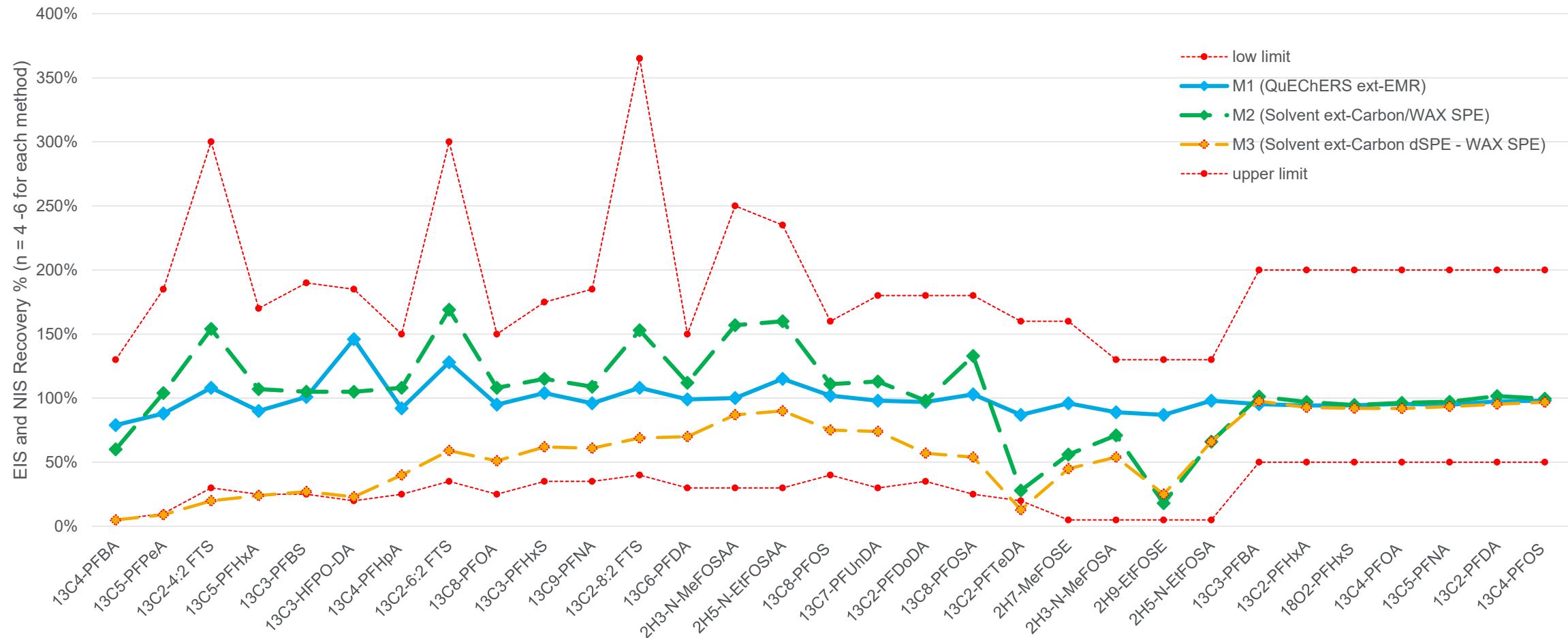


**B) Method Comparison for PFAS in Tissue Quantitative Analysis**  
- Targets, EIS, and NIS Repeatability (RSD%)



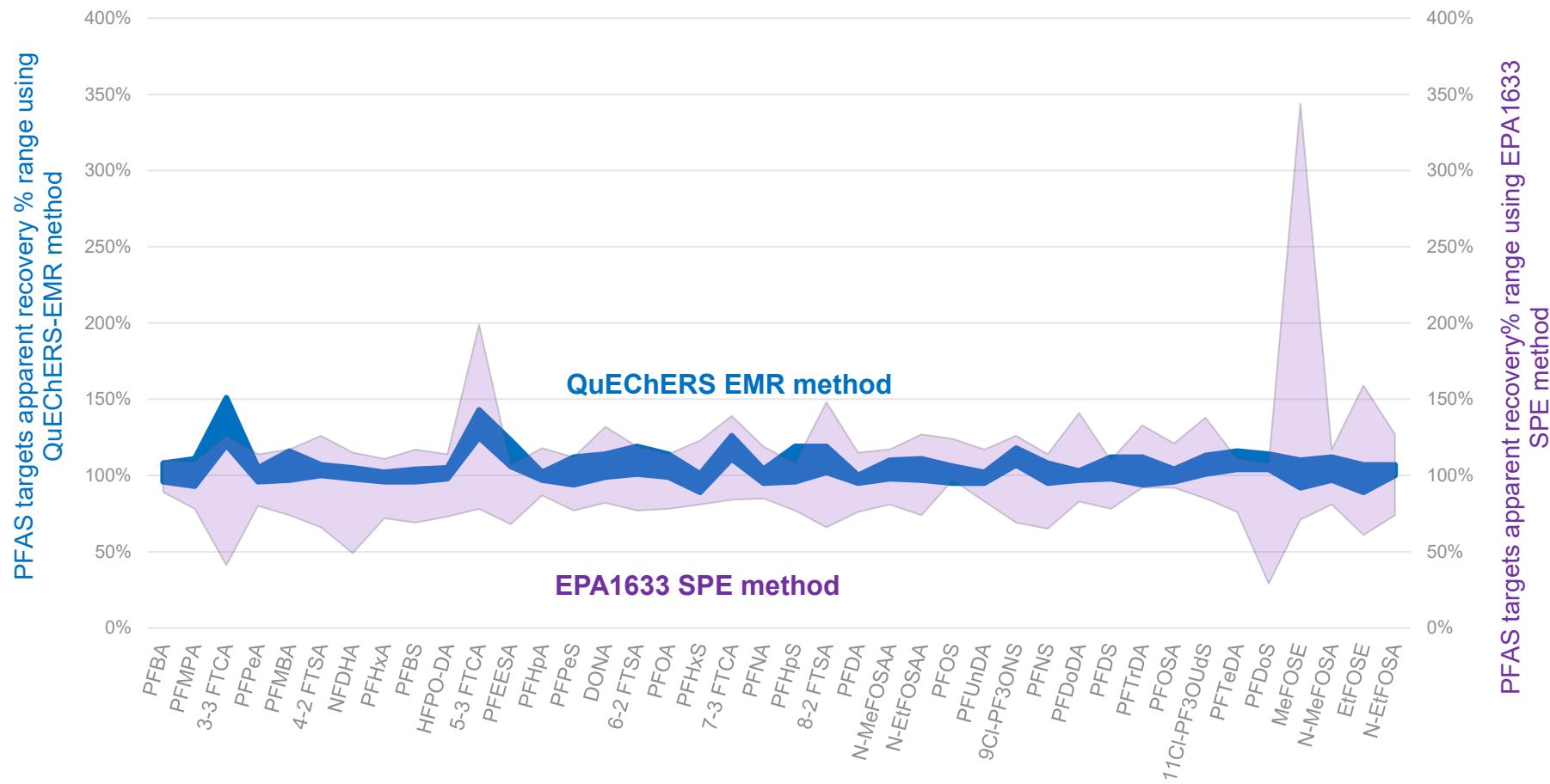
The QuEChERS-EMR method demonstrated improved quantitation performance for PFAS compound recovery and repeatability.

# EIS and NIS Recovery Comparison



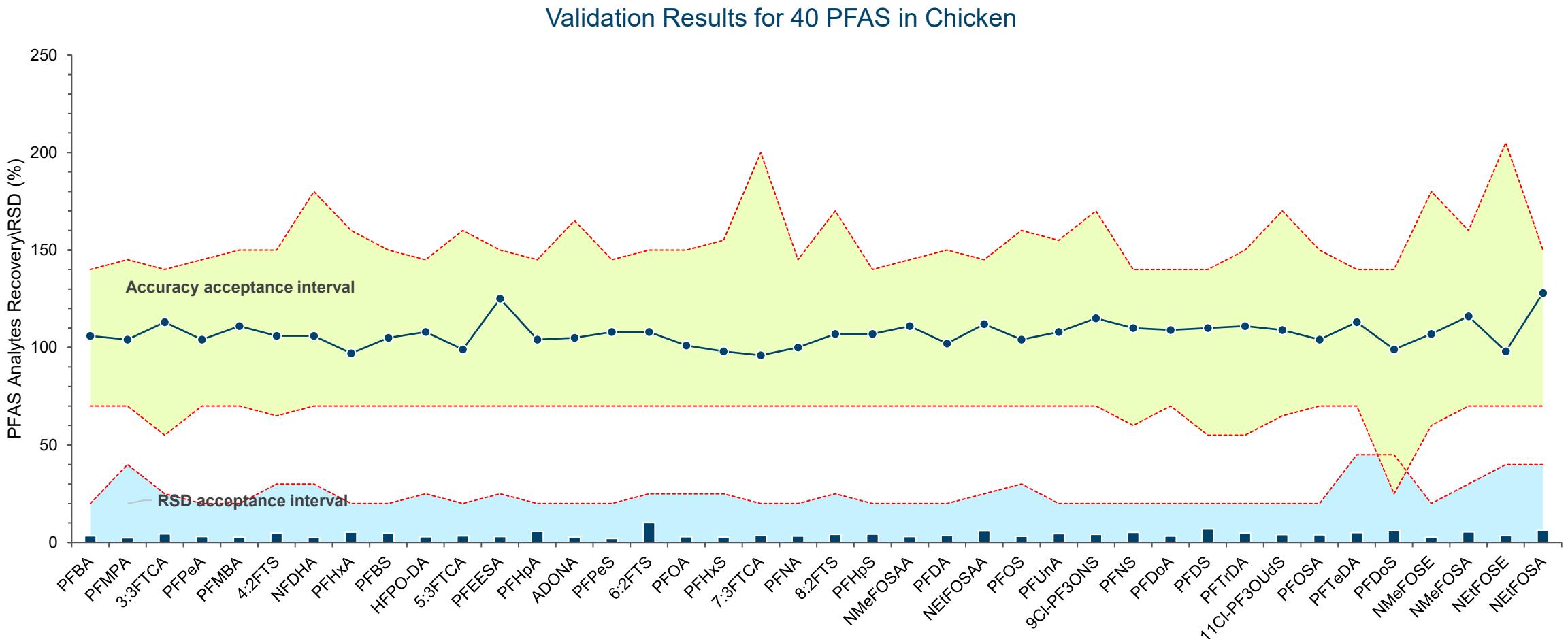
- QuEChERS-EMR protocol presented excellent recovery for all EIS compounds
- Traditional EPA 1633 protocols showed various EIS recoveries across the targets.

# Comparison 40 PFAS Targets in Fish Recovery (Single-lab validation)



- Results from QuEChERS EMR method were based on full validation results from three spiking levels with six replicates of each level.
- Results from EPA 1633 SPE method was based on single-lab study reported from *EPA draft method 1633 (EPA 1633:2021)*.
- QuEChERS-EMR method consistently produced narrower accuracy ranges close to 100% across all PFAS targets.

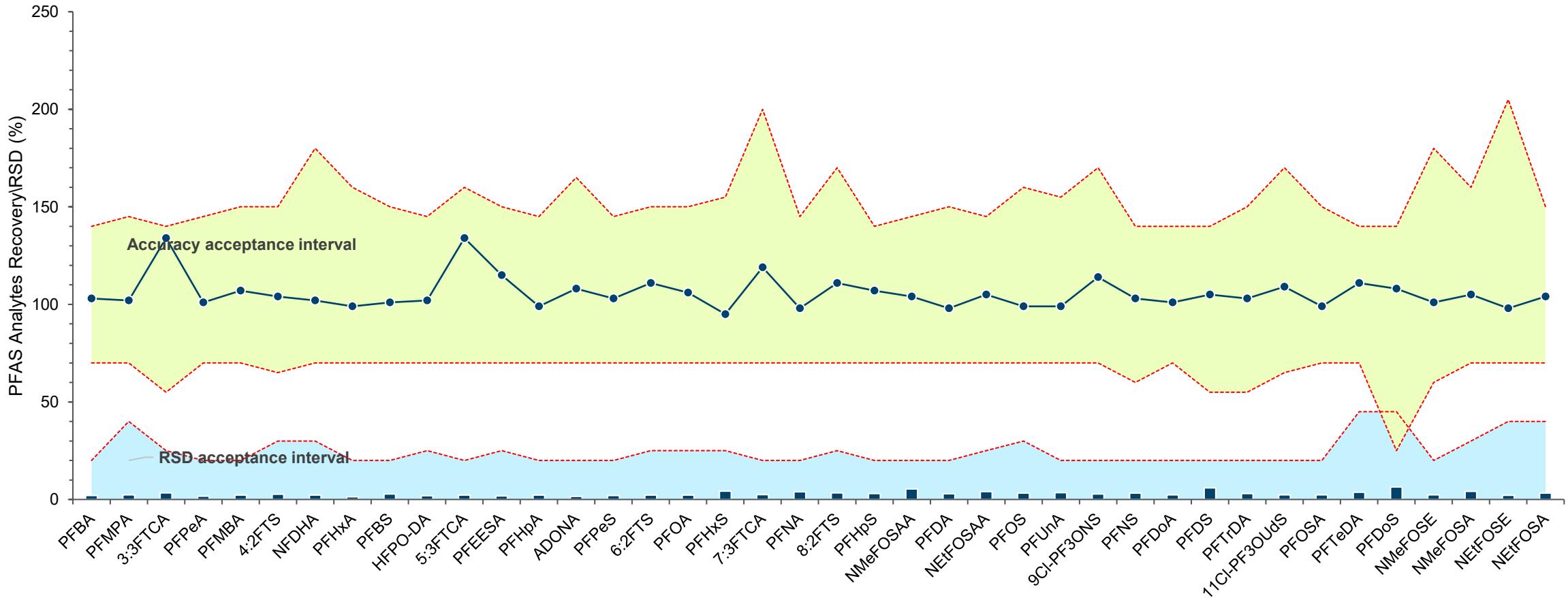
# Validation Results for PFAS in Chicken



- Summary results are based on average of three spiking levels at LOQ, mid-QC (4x of LOQ), and high-QC (40x LOQ)
- Validated LOQ (spiking): 0.05 – 1.25 µg/kg

# Validation Results for PFAS in Tilapia

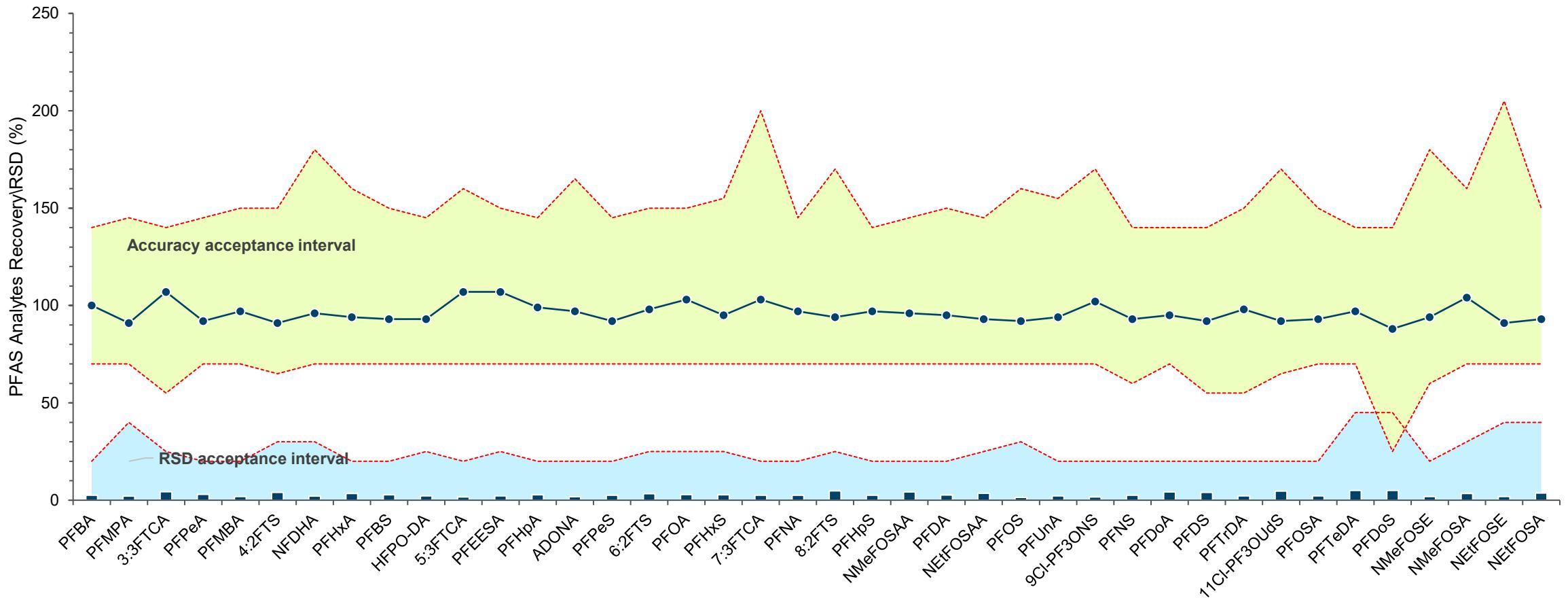
## Validation Results for 40 PFAS in Tilapia



- Summary results are based on average of three spiking levels at LOQ, mid-QC (4x of LOQ), and high-QC (40x LOQ)
- Validated LOQ (spiking): 0.05 – 1.25 µg/kg

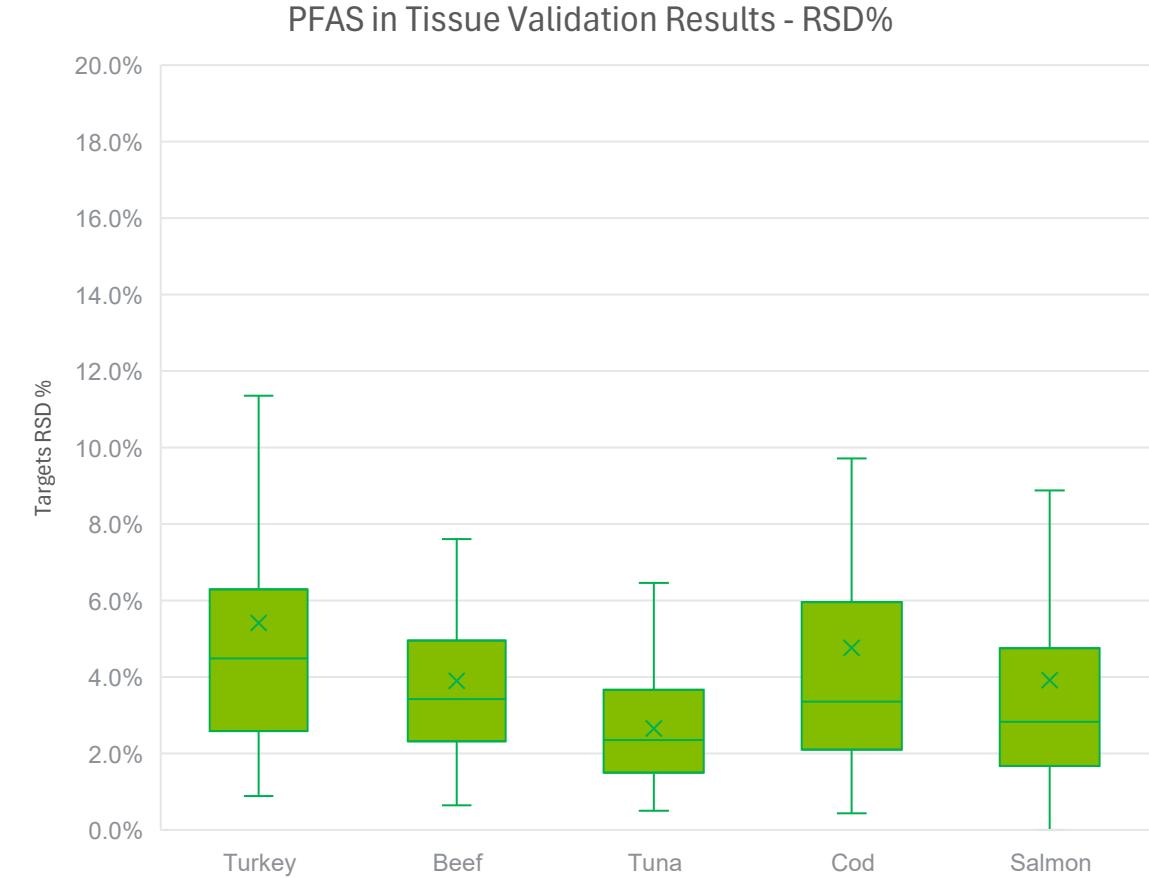
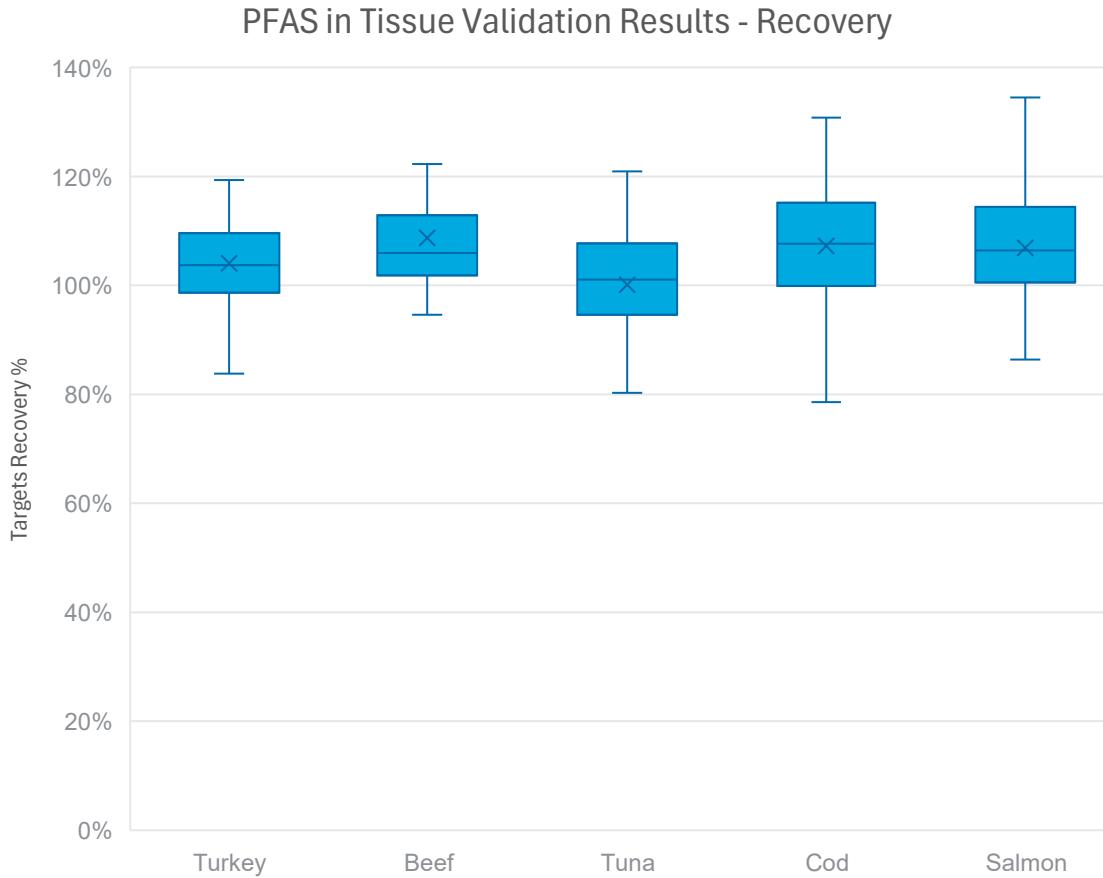
# Validation Results for PFAS in Pork

## Validation Results for 40 PFAS in Tilapia



- Summary results are based on average of three spiking levels at LOQ, mid-QC (4x of LOQ), and high-QC (40x LOQ)
- Validated LOQ (spiking): 0.05 – 1.25 µg/kg

# Cross-validation Results for PFAS in Cod, Tuna, Salmon, Turkey and Beef



- Summary results are based on LOQ spiking levels in additional five more tissue matrices
- LOQ between 0.05 – 0.2 µg/kg, except 1.25 µg/kg for 5:3 FTCA and 7:3 FTCA

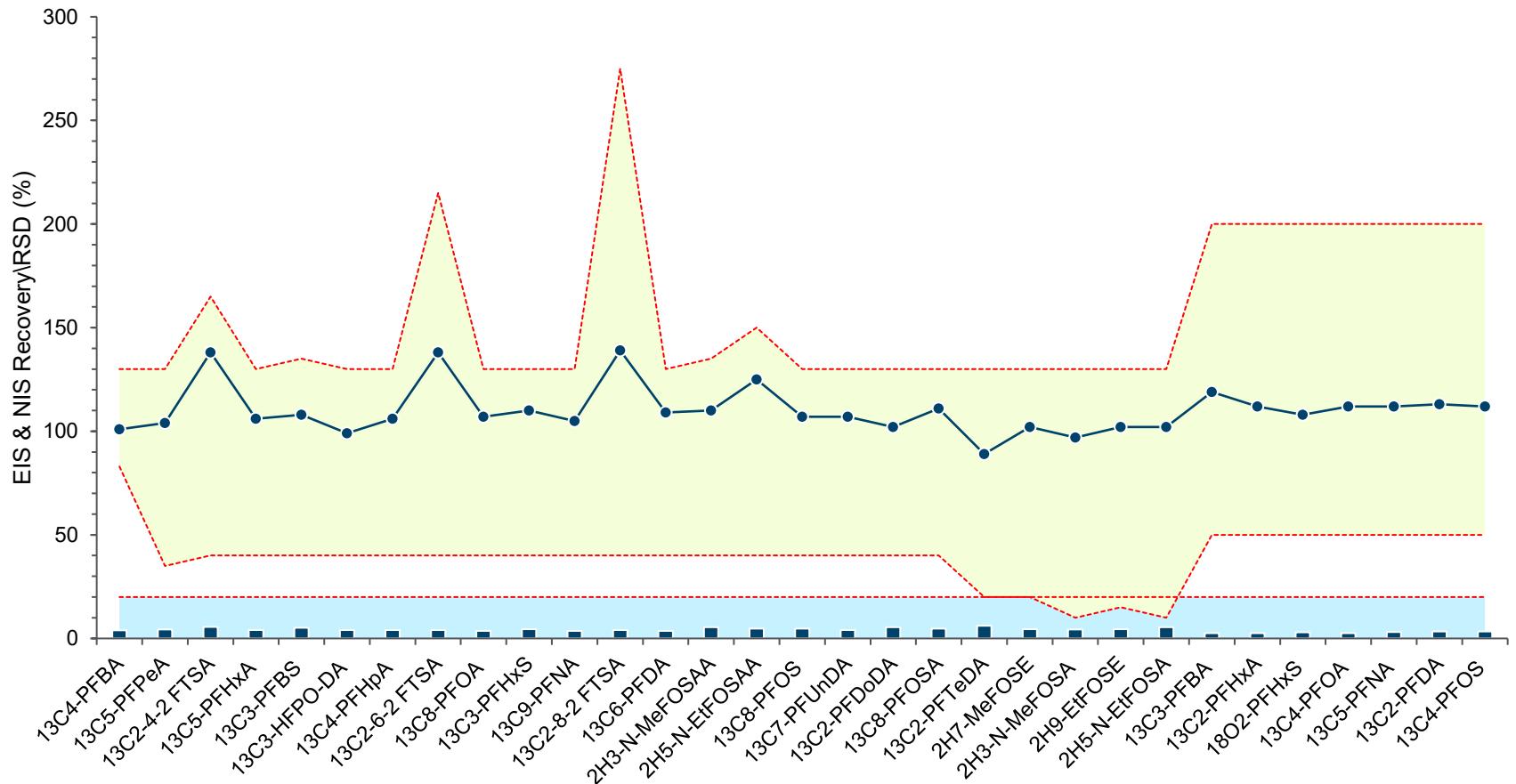
# PFAS Analysis in Biosolid and Soil

Extended applications

# Method Extension to Soil

- Soil is a moderate complex matrix without too much fatty components
- Same instrument method
- Modified sample prep method
  - Captiva EMR PFAS I, 680 mg cartridge
  - Direct loading of QuEChERS crude extract
- LOQs (spiking validated): 0.05 – 1.25 µg/kg

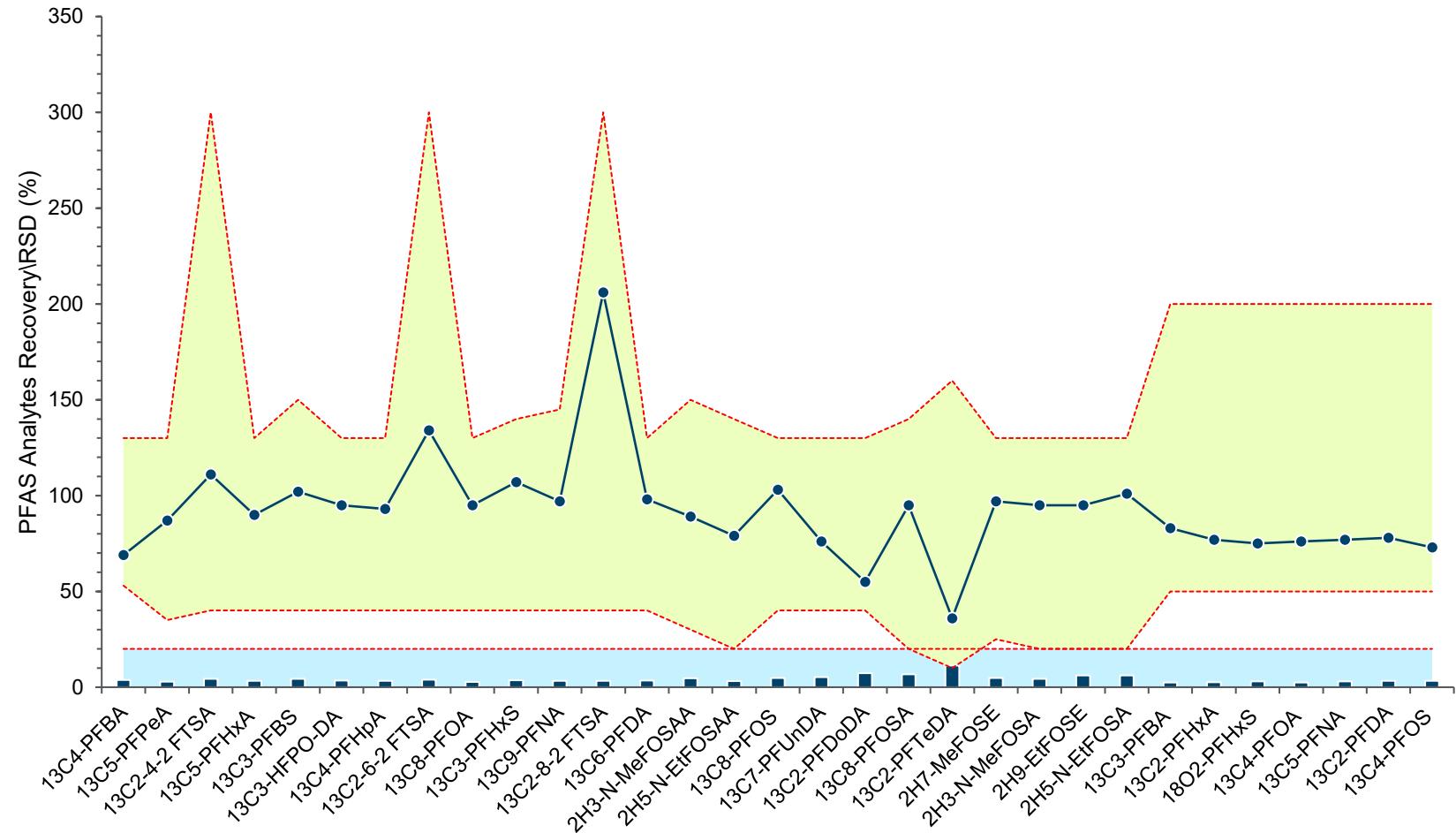
Validation Results for EIS and NIS in Soil



# Method Extension to Biosolid

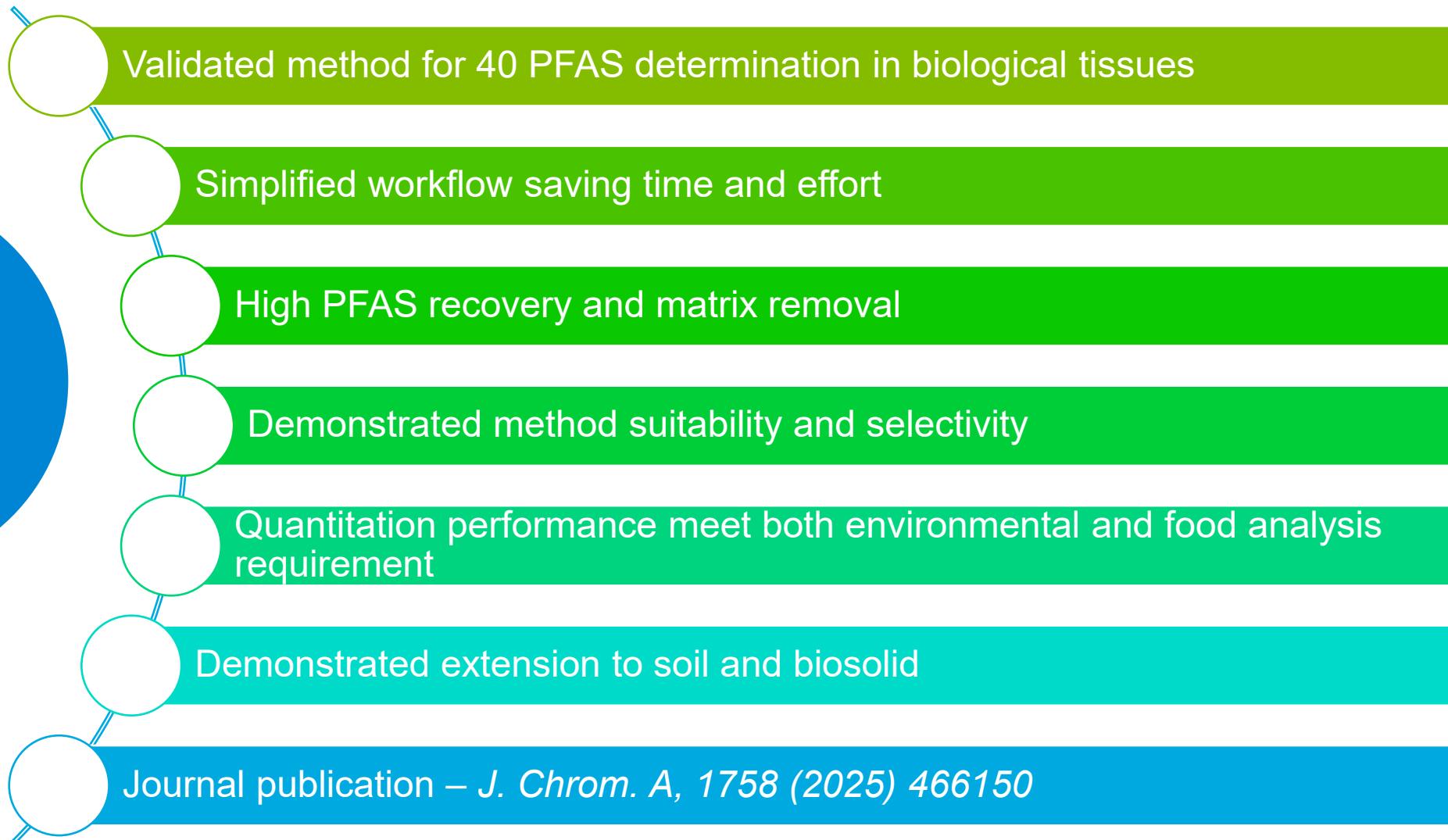
- Biosolid matrix is significantly complex with high positive background
- Same instrument method
- Modified sample prep method
  - Smaller sample size: 0.5 g
  - Captiva EMR PFAS II cleanup
  - Reduced loading volume: 2 mL
- LOQ (calculated and spiking validated): 0.03 – 136.3 µg/kg

Validation Results for EIS and NIS in Biosolid



# Summary

## QuEChERS - EMR



# Acknowledgement

Carrie Xu from New Jersey Department of Health

Matthew Giardina, Megan Juck, and other colleagues from  
Agilent

[Limian\\_zhao8@agilent.com](mailto:Limian_zhao8@agilent.com)

Additional related talks:

1. A Semi-Automated Workflow for the Extraction and Analysis of 40 PFAS Targets in Biosolids, Emily Parry, Wednesday 2pm
2. The Evaluation of Novel Weak Anion Exchange and Graphic Carbon Sorbent Blends for PFAS Extraction and Matrix Reduction in Environmental Extracts, Mattew Giardina, Thursday 11 am.

