

Non target and suspect screening PFAS

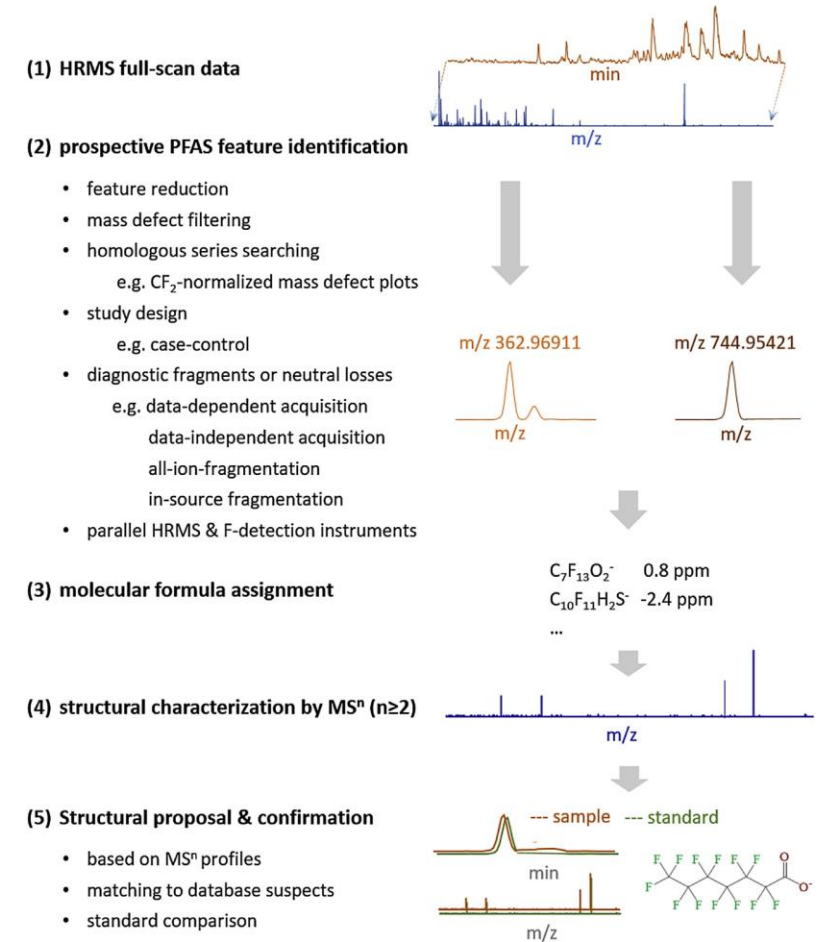
Werkgroep 28/11/2023

Griet Jacobs & Emeline Hanozin

NTS and SS PFAS

Non target screening PFAS

- Extraction and analysis according to WAC/IV/A/025
- Record full scan (pos and neg)
- Record fragmentation data (ddMS2)
- Data processing
 - Deconvolution
 - Retention time alignment (triple injections)
 - Prioritization of data
 - Mass defect and md/C – m/C plot
 - Molecular formula assignment
 - Structure proposal
 - Search in databases (fragmentation and structures)
 - Quantification or sem-quantification

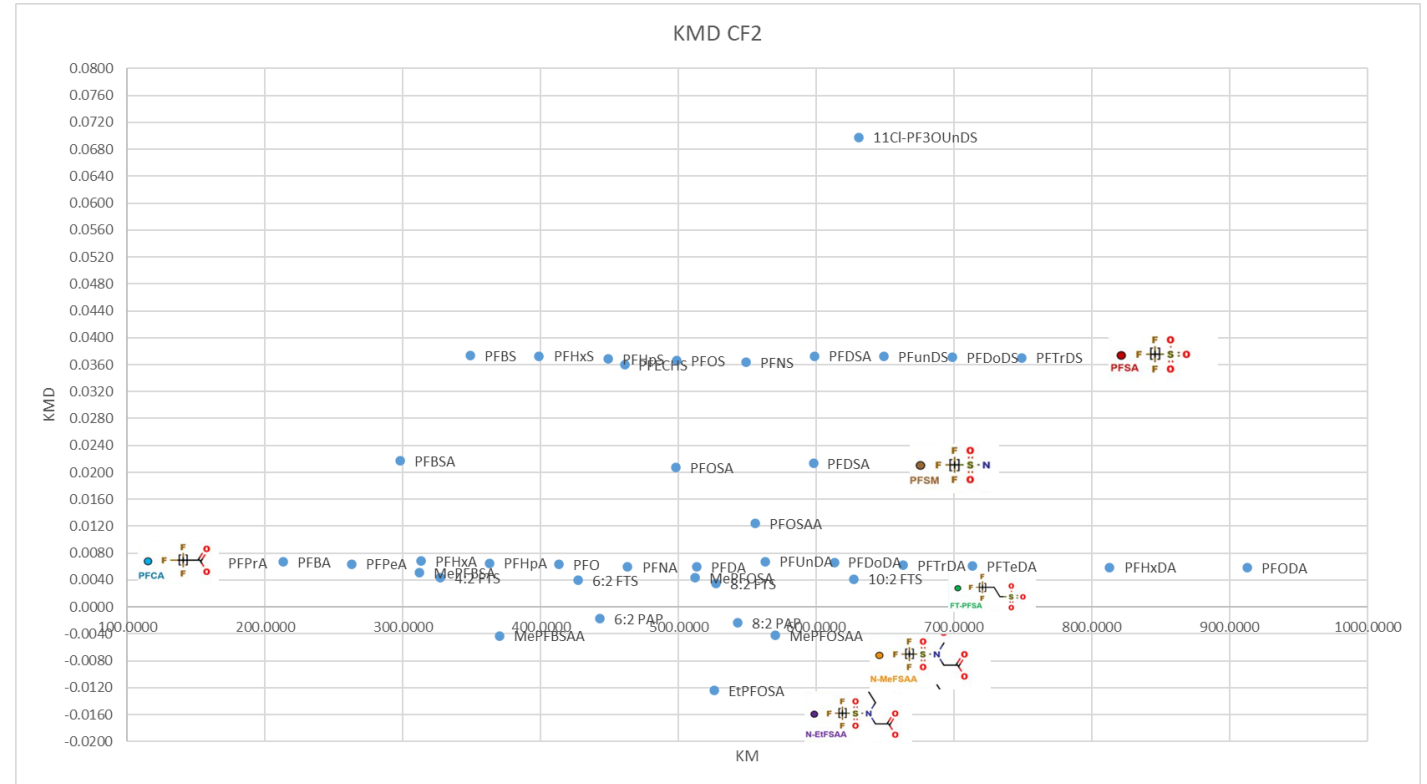
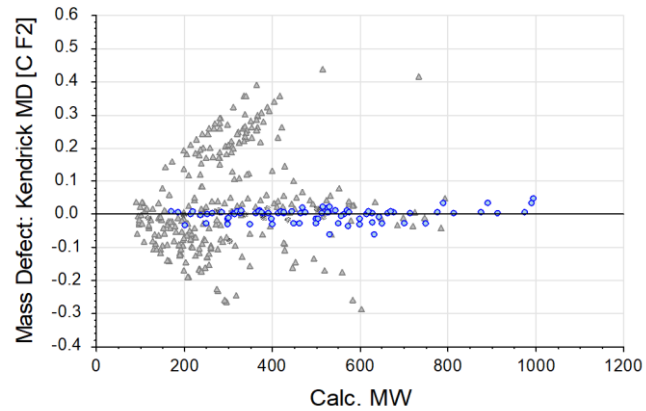


Liu *et al*, *Trends in Anal. Chem.* 2019

NTS and SS PFAS

Non target screening PFAS

- Prioritization via
 - Kendrick mass defect (CF2)



NTS and SS PFAS

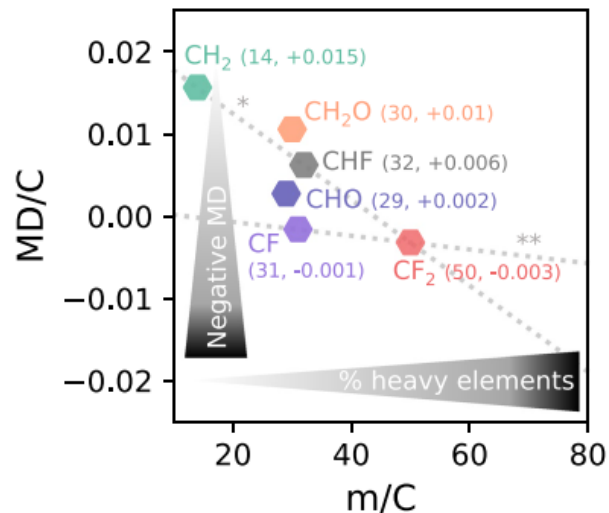
Non target screening PFAS

- Prioritization via
 - Md/C – mC approach
 - Kaufman et al. 2022 DOI: [10.1093/jaoacint/qsac071](https://doi.org/10.1093/jaoacint/qsac071)
 - Zweigle et al. 2023; DOI: [10.1007/s00216-023-04601-1](https://doi.org/10.1007/s00216-023-04601-1)

$C = 100 \cdot \text{abundance of the first isotopic peak} / \text{abundance of the corresponding monoisotopic peak} / 1.1145$

Mass over carbon “m/C” was calculated by dividing the measured mass of the monoisotopic peak over the estimated number carbon atoms “C.”

Mass defect over carbon “md/C” was calculated by dividing the measured mass defect “md” of the monoisotopic ion over the estimated number of carbon atoms “C.”



$$MD/C_{CH_xF_{2-x}} \approx -5.24 \times 10^{-4} \cdot m/C + 0.023 \quad (1)$$

$$MD/C_{CF_x} \approx -8.406 \times 10^{-5} \cdot m/C + 0.001 \quad (2)$$

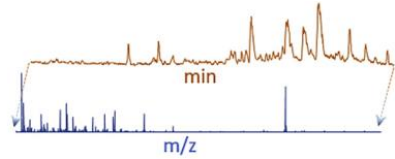
NTS and SS PFAS

Suspect screening

- After identification of unknown PFAS
 - How much is present in the sample?
- Not always reference standards commercially available
- How to quantify?

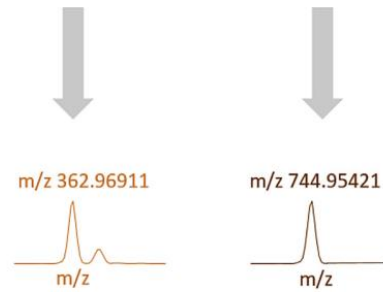
LC-HRMS Workflow for nontarget discovery of PFAS

(1) HRMS full-scan data



(2) prospective PFAS feature identification

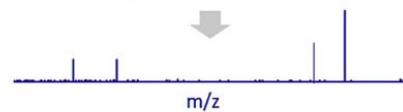
- feature reduction
- mass defect filtering
- homologous series searching
e.g. CF_2 -normalized mass defect plots
- study design
e.g. case-control
- diagnostic fragments or neutral losses
e.g. data-dependent acquisition
data-independent acquisition
all-ion-fragmentation
in-source fragmentation
- parallel HRMS & F-detection instruments



(3) molecular formula assignment

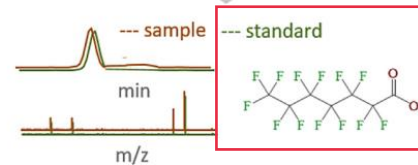
$\text{C}_7\text{F}_{13}\text{O}_2^-$ 0.8 ppm
 $\text{C}_{10}\text{F}_{11}\text{H}_2\text{S}^-$ -2.4 ppm
 ...

(4) structural characterization by MS^n ($n \geq 2$)



(5) Structural proposal & confirmation

- based on MS^n profiles
- matching to database suspects
- standard comparison



Not always available!

Current NIST PFAS list (v1.5)

> 4500 PFAS

EPA master list

>10000 PFAS

LC-ESI(-)-HRMS

50 target PFAS standards commercially available

30 PFAS labeled standards (int. std. or surrogate)

LC-ESI(+)-HRMS

5 target PFAS standards commercially available

0 PFAS labeled standards

How to estimate suspect PFAS concentration without each individual standards being commercially available?

How to estimate suspect PFAS concentration?

Guidelines to choose the best target-surrogate pair:

Same number of F

Same head group

Same class

Same chain length

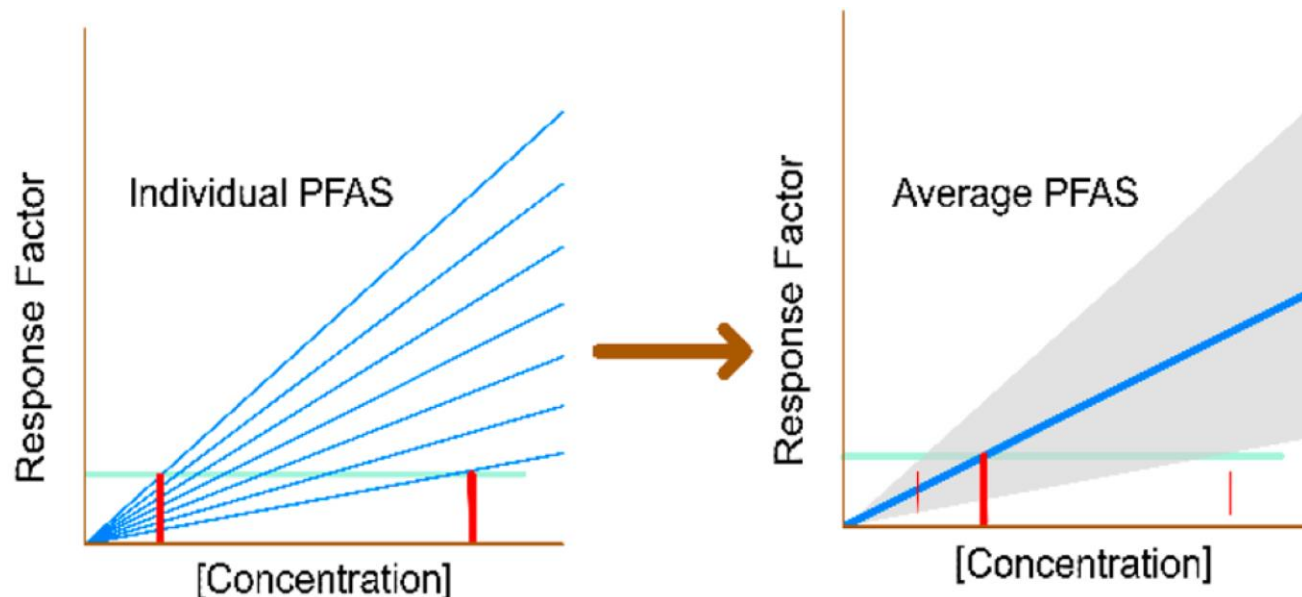
Similar retention time

1:1 matching strategies
Currently used
but no universal selection

Molar response factors vary within a given PFAS class!

Alternative approach:

Use average target PFAS response factor



The concept of “average calibration curve”

→ 50 anionic target PFAS

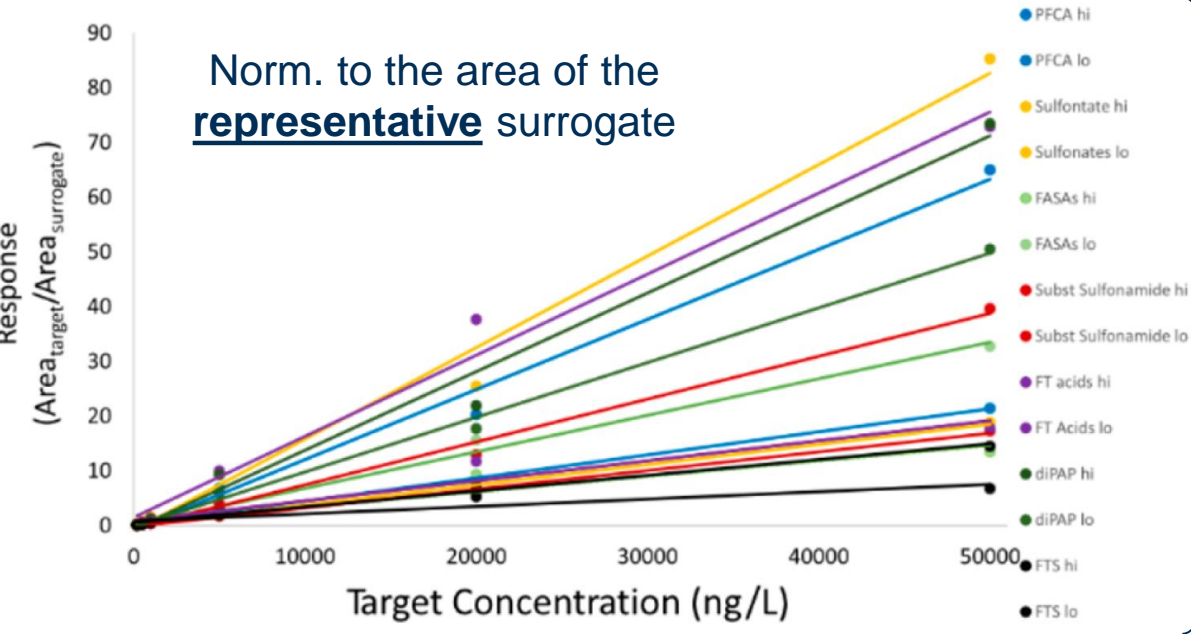
→ 30 labeled anionic (non)matched surrogate standards

→ 5 zwitterionic/cationic target PFAS

→ Prometon-d3, metolachlor-d6 as nonmatched surrogate standards

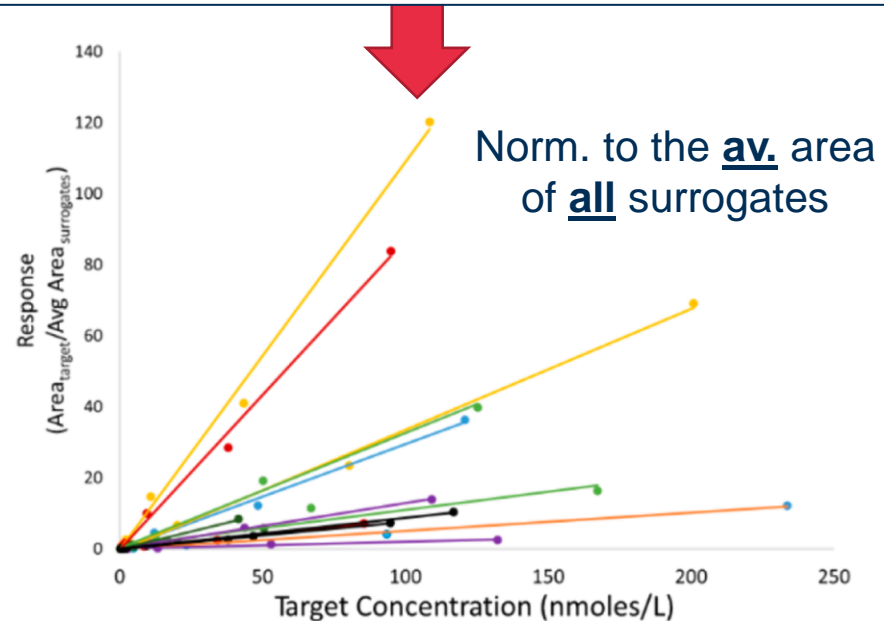
Response of PFAS by classes

High and lowest response within each PFAS class



7 classes

→ Nanomolar response for all classes
 → Response vary by head group and F tail length
 → Overlapping response ranges
 → PFAS classes can be also treated together



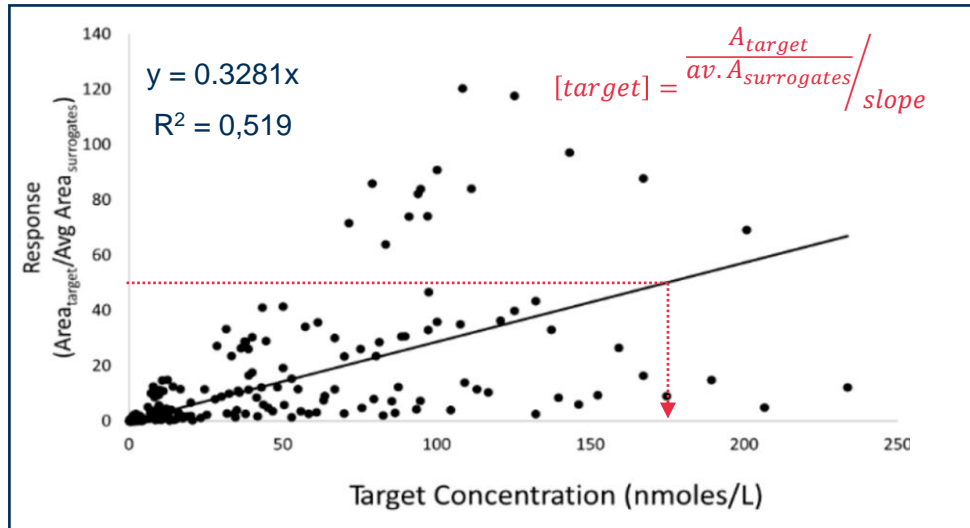
All other homologues within each class give calibration curves that fell within the maximum and minimum of the class



The concept of “average calibration curve” – Performance

Average calibration curve

1/ x weighted linear regression (forced to 0)



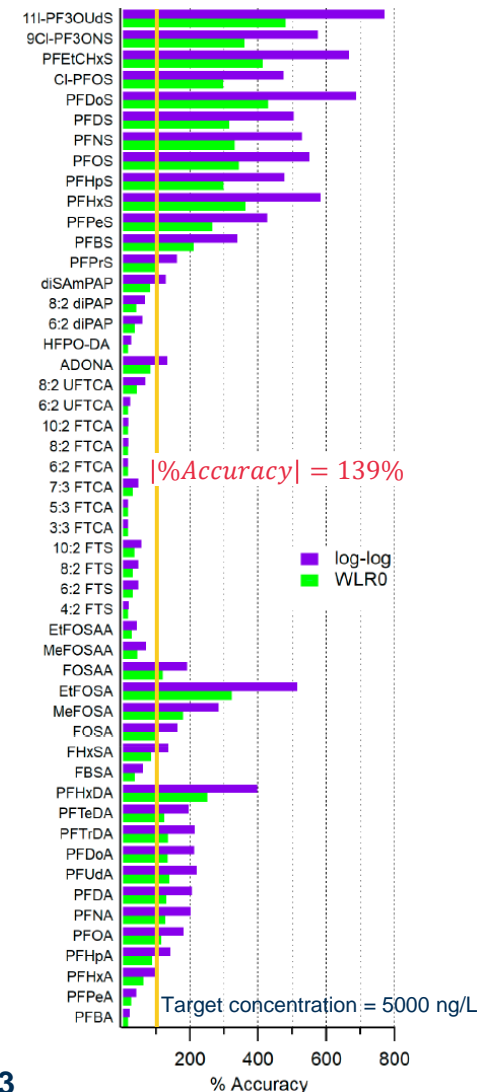
overestimated

underestimated

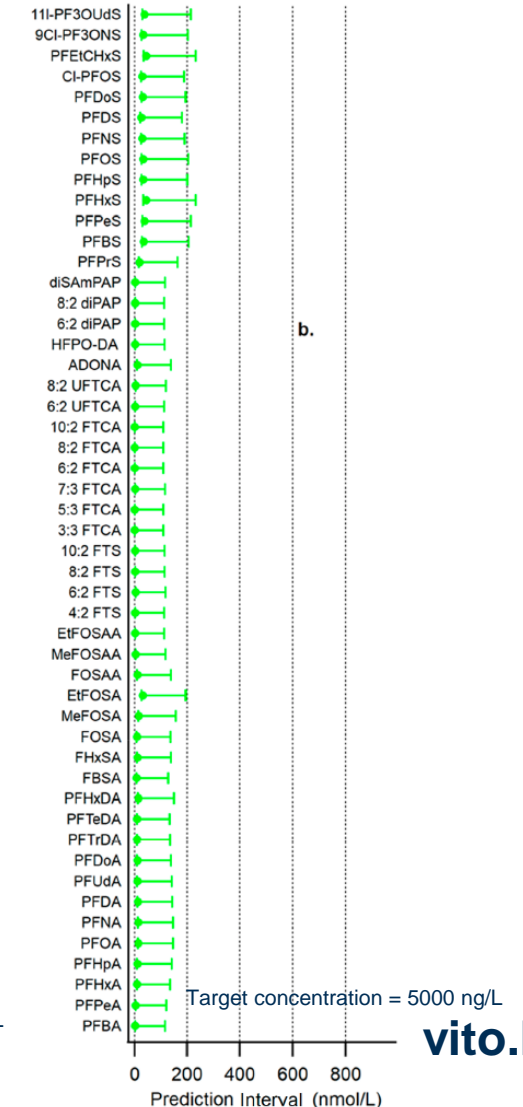
Advantage:

- No preliminary knowledge of the suspect response or structure are required to estimate the concentration.
- Continuous expansion of the average calibration curve to include new standards.
- Average curve can be constructed for each PFAS class.
- Fast, reproducible over time.
- More uniform reporting of suspect concentrations.

$$\%Accuracy = \frac{[target]_{fit}}{[target]_{expected}} * 100$$



Upper and lower bounds from 95% prediction intervals



Practical aspects

- Next steps:
 - Apply the approach to WAC compounds and samples
- New WAC method will be available in 2024 for suspect screening
 - that include semi-quantification via the average calibration curve
- Ring trial needed for NTS and/or SS?