

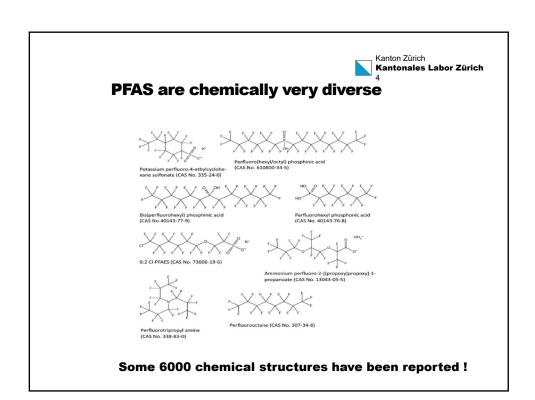






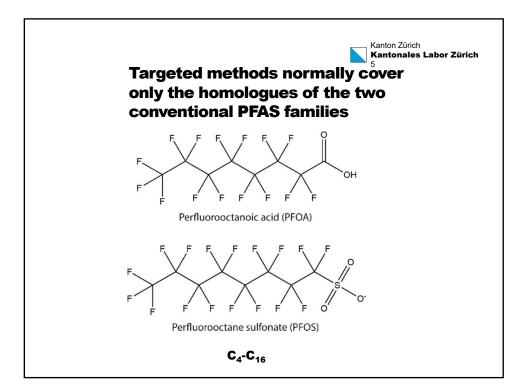
PFAS

- Are everywhere (textile impregnation; firefighting foam etc.)
- Are chemically inert
- Can accumulate in animal and human tissue
- Are suspected to have various negative toxicological effects











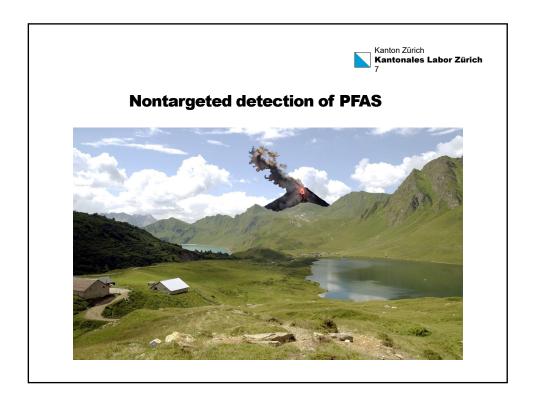
Lack of commercial reference substances

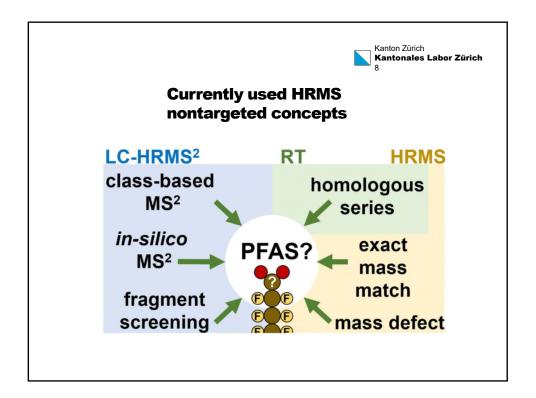
Detection and quantification is only possible for a limited set of PFAS

Does monitoring a limited set of compounds answer the question, how humans are exposed to PFAS originating from the environment or food?















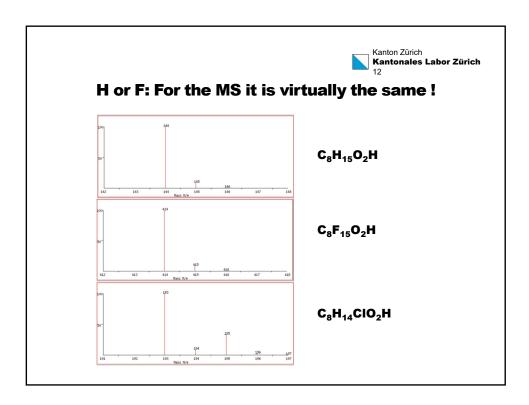




Nontargeted PFAS is hindered by the fact that fluorine has no natural isotopes

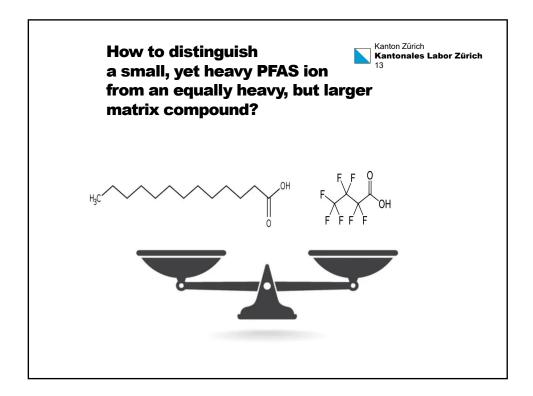
Atoms like S, Cl, Br, I show characteristic isotopes which facilitates MS based detection

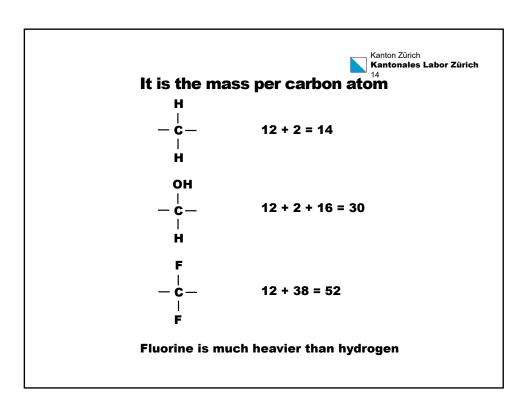
Fluorine has a slight mass defect, yet this is smaller than that of Cl, S or even O















Turning a weakness into a strength

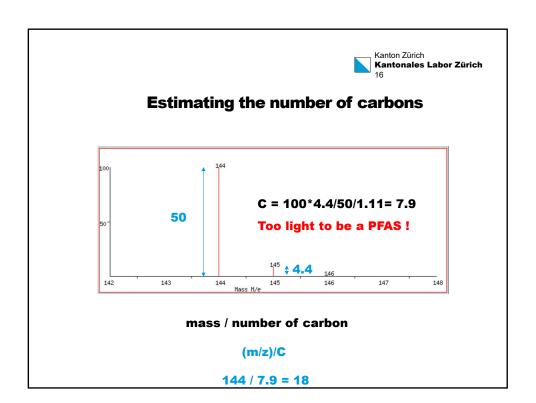
The absence of F isotopes means that the isotopic profile remains unchanged when H is substituted by F

This means that the ion abundance of the first isotopic ion of a PFAS is primarily defined by the number of carbon atoms present within the chemical formula

This permit us to estimate the number of carbons and with this a PFAS marker parameter:

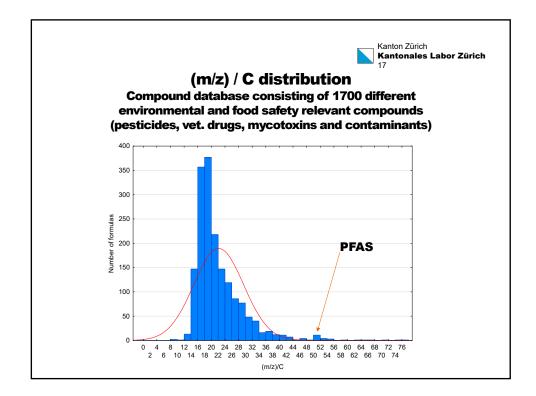
mass of unknown/ estimated number of carbon within the unknown:

(*m/z*) / C











Deconvolution and componentization

- Ten thousands of chromatographic peaks are extracted
- Based on retention time and peak shape extracted mass traces are grouped to produce a "feature"
- A feature represents a chromatographically resolved compound

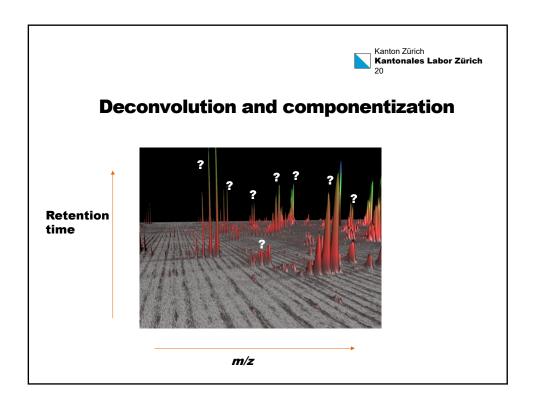






Analyzing the spectrum of each feature

- The spectrum of a feature contains the monoisotopic ion, the isotopic peaks and possible adducts or fragments
- Ratios of abundances of isotopic peaks versus the monoisotopic peak and the accurate mass of the monoisotopic peak for each extracted feature are exported into Excel



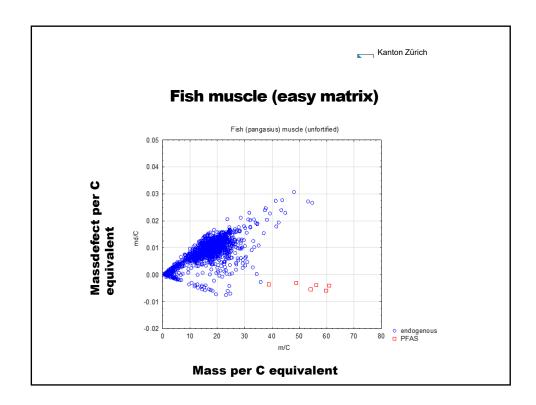






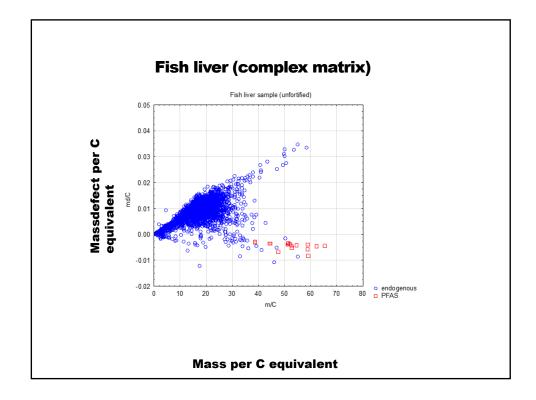
How well does it work?

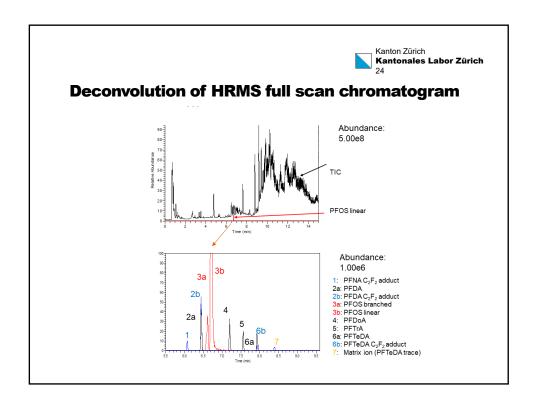
- Processing HRMS data from extracts which were produced by a validated, targeted PFAS method
- Anion exchange based clean-up of fish matrix samples





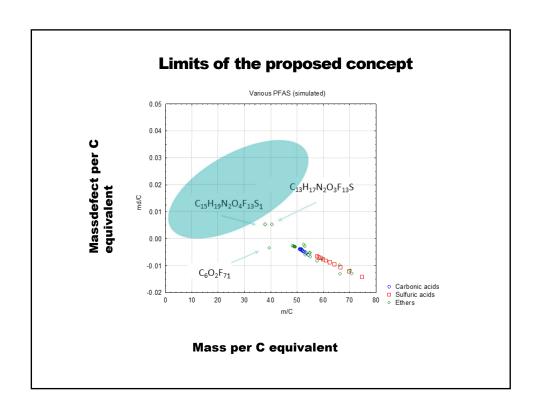












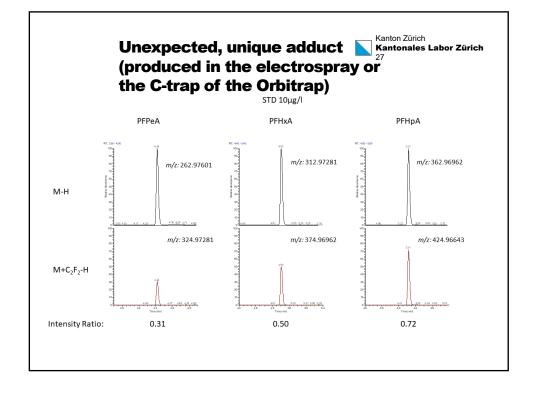
How did we perform?



- The nontargeted approach successfully detected the PFAS covered by the targeted method
- Perfluorotetradecanoic acid was only detected by the nontargeted methodology (the compound was not present in the standard mix)
- A set of previously unreported MS adducts was elucidated







Recent developments



A. Kaufmann: Simplifying Nontargeted Analysis of PFAS in complex Food matrices; J. AOAC int. 2022, Sep. 6 (105), 1280-1287

- The commercial software "Compound discoverer" integrated the proposed algorithm.
- J. Zweigle: Efficient PFAS prioritization in Nontarget HRMS data: systematic evaluation of the novel MD/C-m/C approach;
 J. ABC, 2023, 415, 1791-1801









Nontargeted works if we can clearly describe a common property of our compounds of interest

This property has to be significantly different from the properties of our matrix compounds

If this is not the case, the exogenous analytes will be hidden within the dense cluster of endogenous compounds



