

Guidance Document on Analytical Parameters for the Determination of Per- and Polyfluoroalkyl Substances (PFAS) in Food and Feed

Version 2.0

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Authorship

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Please double click on the pdf-icon to open the ANNEX

Also available online under https://eurl-pops.eu/working-groups#_pfas



1. GENERAL

1.1. Field of Application

This document on analytical parameters for the determination of selected per- and polyfluoroalkyl substances (PFAS) in food and feed was developed within the network of the European Union Reference Laboratory (EURL) for halogenated POPs in Feed and Food, the respective National Reference Laboratories (NRLs) of EU member states and international experts in the field of persistent organic pollutants (POPs) analysis. The guidance in this document is intended for laboratories involved in the official control of contaminants in food and feed and focuses on the determination of these substances in the laboratory. It is intended as general guidance for laboratories and particularly for those that do not have an existing method. It provides useful key elements in a set of analytical parameters contributing to further harmonization in the field of PFAS analysis in food and feed as part of the EURL's official mandate and scope of work.

The specific performance requirements as laid down in Commission Implementing Regulation (EU) 2022/1428 Annex Part B Table 5 are the basis for the general principles for PFAS analysis in food and feed. The present guidance document gives recommendations on the implementation of analytical methods to achieve these general principles in laboratories involved in the official control in the EU. The document is not legally binding. However, Commission Implementing Regulation (EU) 2022/1428 states that "the principles as described in the EURL Guidance Document [...] shall be followed".

NOTE: All recommendations given in this document should be considered as "Guidance for reliable analyses".

Abbreviation	Definition
br-PFOS	Branched perfluorooctane sulfonic acid
CWG	Core Working Group
EC	European Commission
ECF	Electrochemical fluorination process
EFSA	European Food Safety Authority
EURL	European Union Reference Laboratory
HDPE	High density polyethylene
HRMS	High resolution mass spectrometry
IARC	International Agency for Research on Cancer
ILIS	Isotope-labelled internal standard
IS	Internal standard
ISO	International Standardisation Organisation
IUPAC	International Union of Pure and Applied Chemistry
LDPE	Low density polyethylene

1.2. Abbreviations



L-PFOS	Linear perfluorooctane sulfonic acid
LOD	Limit of detection
LOQ	Limit of quantification
LRMS	Low resolution mass spectrometry
ME	Matrix effect
ML	Maximum level
MRM	Multiple reaction monitoring
NRL	National Reference Laboratory
OFL	Official Laboratory
PEEK	Polyether ether ketone
PFAS	Per- and polyfluoroalkyl substances
PFCA	Perfluoroalkyl carboxylic acids
PFHxA	Perfluorohexanoic acid
PFHxS	Perfluorohexane sulfonic acid
PFNA	Perfluorononanoic acid
PFOA	Perfluorooctanoic acid
PFOS	Perfluorooctane sulfonic acid
PFSA	Perfluoroalkyl sulfonic acids
POPs	Persistent organic pollutants
PTFE	Polytetrafluoroethylene
PVDF	Polyvinylidene fluoride
QC	Quality control
RS	Recovery standard
RT	Retention time
SANTE	Directorate-General for Health and Food Safety, European Commission
S/N	Signal to noise ratio
SRM	Selected reaction monitoring
TDCA	Taurodeoxycholic acid
(U)HPLC	(Ultra) High performance liquid chromatography
w.w.	Wet weight

1.3. Introduction

Information on sampling is not given in this document. The reader is referred to the Commission Implementing Regulation (EU) 2022/1428 laying down methods of sampling and analysis for the control of perfluoroalkyl substances in certain foodstuffs [1]. In addition Commission Regulation (EC) No 152/2009 for PCDD/Fs and PCBs in feed [2] can be used as the directions given therein are expected to be similar for PFAS. Additionally, all procedures used for sampling should avoid the use of equipment containing polytetrafluoroethylene (PTFE) or other fluoropolymers in order to minimize the risk of contamination.



1.4. Background

Per- and polyfluoroalkyl substances (PFAS) are a group of man-made organic compounds consisting of a hydrophobic fluorinated alkyl chain and a hydrophilic functional group. This class of compounds includes a large number of substances, namely all that contain the perfluoroalkyl moiety (C_nF_{2n+1} -) [3]. In polyfluorinated substances, one or more CF_2 moieties are replaced by the corresponding number of CH_2 groups.

PFAS have been used since the 1950s. Due to the stability of the C-F bond, many PFAS are resistant to biological, chemical and physical transformation. Some PFAS – often called 'precursors' – may undergo biological, chemical and physical transformation to a stable PFAS. PFAS are widely used as monomeric or polymeric substances in direct or indirect uses and subsequently have been found in the environment (water, air, soil, sediments, and biota) but also in food, wildlife and humans.

Three of the most frequently used PFAS have been listed in the annexes of the Stockholm Convention on Persistent Organic Pollutants (POPs) – perfluorooctane sulfonic acid (PFOS) in 2009, perfluorooctanoic acid (PFOA) in 2019 and perfluorohexane sulfonic acid (PFHxS) in 2022 – with the aim of elimination of production and uses [4]. In the most recent scientific opinion by the European Food Safety Authority (EFSA), four PFAS have been assessed, namely, PFOS, PFOA, PFHxS and perfluorononanoic acid (PFNA) [5]. As of 1 January 2023, maximum levels apply for PFOS, PFOA, PFNA, PFHxS and the sum of PFOS, PFOA, PFNA and PFHxS in certain foodstuffs [6].

In 2023, the International Agency for Research on Cancer (IARC) has re-evaluated the carcinogenicity of PFOS and PFOA. PFOS is now classified as possibly carcinogenic to humans (Group 2B) and PFOA as carcinogenic to humans (Group 1) [7]. In the EU, PFOA has a harmonised classification as a suspected carcinogen and presumed human reproductive toxicant [8].

PFOS and related substances have been produced by the electrochemical fluorination process (ECF) mainly; and thus, may occur as a mixture of linear (L-PFOS) and branched (br-PFOS) isomers. PFOA has been produced by either the ECF process to generate linear and branched isomers or the telomerisation process from pentafluoroethyl iodide (in the case of PFOA), which results in linear products. For PFHxS, there are currently no known direct uses. PFNA is used as a surfactant in the production of the fluoropolymer polyvinylidene fluoride (PVDF). In addition, PFNA is a by-product of the synthesis of PFOA and short-chain perfluoroalkyl carboxylic acids (PFCA) such as perfluorohexanoic acid (PFHxA).

In 2018, the European Commission, Directorate-General for Health and Food Safety (SANTE), asked the European Union Reference Laboratory (EURL) for halogenated POPs in Feed and Food to develop the analytical parameters and conditions to reach low limits of quantification in the routine analysis of food to protect human health from adverse effects of PFAS from food consumption. Subsequently, under the coordination of the EURL, a core working group on perfluoroalkyl substances ("CWG PFAS" for short) was established.



1.5. Scope

The recommendations contained within this guidance document apply to PFAS analysis, in particular to analysis of PFCA and perfluoroalkyl sulfonic acids (PFSA), perfluorooctane sulfonamide (FOSA), and selected substitutes in food and feed matrices. The recommendations may also be applied to other PFAS.

The recommendations are designed to allow the monitoring of PFAS concentrations in food and feed as part of studies on:

- the establishment and update of maximum levels of these contaminants,
- the establishment or maintenance of databases that may be used to recommend legal limits,
- the exposure assessment of populations through dietary intake and assessment of risk,
- enforcement (if limits are established).

Other purposes could include studies on time trends and patterns in order to identify the source(s) of possible contamination particularly during incidents involving such contamination.

1.6. Analytes of Interest

The requirements given in this document apply to the following PFAS (**Table 1**) in food and feed samples. PFCA und PFSA share similar physico-chemical properties, can be captured by one analytical method, and have been found to be of most concern in food and feed [5]. In addition, further emerging perfluoroalkyl substances may be considered, such as FOSA and PFAS substitutes (**Table 1**). The requirements may also be applicable to the analysis of other PFAS (e.g. PFCA/PFSA precursors) and matrices (such as human tissues, environmental samples and drinking water), but these are beyond the scope of this document.

Acronym	Description	Remark	
Perfluoroalkyl Carboxylic Acids (PFCA)			
PFBA	Perfluorobutanoic acid		
PFPeA	Perfluoropentanoic acid		
PFHxA	Perfluorohexanoic acid		
PFHpA	Perfluoroheptanoic acid		
PFOA	Perfluorooctanoic acid	Main compound [5] (reported as "total-PFOA")	
PFNA	Perfluorononanoic acid	Main compound [5] (reported as "total-PFNA")	
PFDA	Perfluorodecanoic acid		
PFUnDA	Perfluoroundecanoic acid		
PFDoDA	Perfluorododecanoic acid		
PFTrDA	Perfluorotridecanoic acid		
PFTeDA	Perfluorotetradecanoic acid		

 Table 1: Analytes of interest

Table 1 (continued)

Acronym	Description	Remark	
Perfluoroalkyl Sulfonic Acids (PFSA)			
PFBS	Perfluorobutane sulfonic acid		
PFPeS	Perfluoropentane sulfonic acid		
PFHxS	Perfluorohexane sulfonic acid	Main compound [5] (reported as "total-PFHxS")	
PFHpS	Perfluoroheptane sulfonic acid		
PFOS	Perfluorooctane sulfonic acid	Main compound [5] (reported as "total-PFOS")	
PFNS	Perfluorononane sulfonic acid		
PFDS	Perfluorodecane sulfonic acid		
PFUnDS	Perfluoroundecane sulfonic acid		
PFDoDS	Perfluorododecane sulfonic acid		
PFTrDS	Perfluorotridecane sulfonic acid		
Perfluoroalkane	Sulfonamides		
FOSA	Perfluorooctane sulfonamide		
PFAS Substitute	es		
DONA	2,2,3-Trifluoro-3-[1,1,2,2,3,3-hexa		
ConV	(trifluoromethoxy)propoxy]-propionic acid		
GenX	2,3,3,3-tetrafluoro-2-(heptafluoropropoxy)-propanoic acid		
F-53B	2-[(6-chloro-1,1,2,2,3,3,4,4,5,5,6,6-dodecafluorhexyl)oxy]-		
	1,1,2,2-tetrafluoroethansulfonic ac component of F-53B)	id (9CI-PF3ONS; major	
	2-((8-chloro-1,1,2,2,3,3,4,4,5,5,6,6	\$7788-	
	hexadecafluorooctyl)oxy)-1,1,2,2-		
	acid (11CI-PF3OUdS; minor component of F-53B)		
Capstone A	1-Propanaminium, N,N-dimethyl-N-oxide-3-		
	[[(3,3,4,4,5,5,6,6,7,7,8,8,8-		
	tridecafluorooctyl)sulfonyl]amino]-, hydroxide		
Capstone B	1-Propanaminium, N-(carboxymethyl)-N,N-dimethyl-3-		
	[[(3,3,4,4,5,5,6,6,7,7,8,8,8-		
	tridecafluorooctyl)sulfonyl]amino]-	, hydroxide	



For purposes of this guidance document, the following terms and definitions apply:

Accuracy: Closeness of agreement between a test result and the accepted reference value [9]. It is determined by determining trueness and precision [10].

Apparent recovery: Observed value derived from an analytical procedure by means of a calibration graph, *expressed as percentage [11]. E.g. when using isotope-labelled internal standards the recovery is corrected for extraction/clean-up losses and matrix-effects and is about 100 %.*

Batch *also referred to as 'lot'*: A quantity of material which is known or assumed to be produced under uniform conditions [12].

Bias: Difference between the estimated value of the test result and an accepted reference value [10,9].

Blanks:

- Calibration blank: A calibration standard that does not contain the analyte(s) of interest at a detectable level [13].
- Reagent blank also referred to as 'procedural blank': Sample that does not contain the matrix that is brought through the entire measurement procedure and analysed in the same manner as a test sample [14].
- Sample blank also referred to as 'matrix blank': Matrix with no analyte present [13].
- Solvent blank: A solution which is made up from the solvent(s) contained in the solution presented to the instrument [13].

Fortified or fortification: Addition of analyte for the purpose of recovery determination [15].

Interference: A systematic error in the measure of a signal caused by the presence of concomitants in a sample [12].

Interlaboratory study: The organisation, performance and evaluation of tests on the same sample by two or more laboratories in accordance with predetermined conditions to determine testing performance. According to the purpose the study can be classified as collaborative study or proficiency study. [10]

Internal standard (IS): A substance not contained in the sample with physico-chemical properties as similar as possible to those of the analyte that has to be identified and which is added to each sample as well as to each calibration standard [16]. *Respective isotope-labelled internal standards are used as the basis for quantification of the analytes.*

Limit of quantification (LOQ): Lowest content of the analyte which can be measured with reasonable statistical certainty [17], i.e. the lowest concentration or mass of the analyte that has been validated with acceptable accuracy by applying the complete analytical method and identification criteria [15].

Lower bound: Concept which requires using zero for the contribution of each non-quantified congener or substance to a sum parameter [17].



Matrix: The material making up the sample.

Matrix effect (ME): The combined effect of all components of the sample other than the analyte on the measurement of the quantity. If a specific component can be identified as causing an effect then this is referred to as interference [12]. The matrix effect (ME) is typically expressed in % and can be calculated according to equation 1 below [18].

$$ME (\%) = \left(\frac{\overline{A}_{[matrix]}}{\overline{A}_{[solvent]}} - 1\right) \times 100$$
 (Eq. 1)

with:

ME = Matrix effect

 A_{matrix} = peak area of the analyte fortified to a blank sample extract before injection $A_{solvent}$ = peak area of the analyte in a solvent standard at same concentration ME < 0 Suppression of the ion signal ME > 0 Enhancement of the ion signal

Measurement uncertainty: A parameter, associated with the results of a measurement, that characterizes the dispersion of the values that could reasonably be attributed to the measurand (the quantity being measured) [12].

Precision: Closeness of agreement between independent test/measurement results obtained under stipulated (predetermined) conditions. The measure of precision usually is expressed in terms of imprecision and computed as standard deviation of the test result. Less precision is determined by a larger standard deviation. [10,9]

Recovery also referred to as 'extraction recovery', 'absolute recovery', or 'recovery factor': Yield of a preconcentration or extraction stage of an analytical process for an analyte divided by amount of analyte in the original sample, expressed as percentage [11].

Recovery standard (RS) also referred to as '*syringe/injection/volumetric standard*': a compound of known chemical purity that is not contained in the sample and is added to every sample, blank or standard at a known concentration, after sample processing and prior to instrument analysis. Recovery standards can be used for quantification of the IS.

Repeatability: Precision under repeatability conditions, i.e. conditions where independent test results are obtained with the same method on identical test items in the same laboratory by the same operator using the same equipment within short intervals of time [9,10]. *E.g. three replicates, same method and sample material, same operator and instruments, within a short interval of time (one day/sequence).*

Selectivity (qualitative): The extent to which other substances interfere with the determination of a substance according to a given procedure [12].

Trueness: Closeness of agreement between the average value obtained from a large series of test results and an accepted reference value [9,10]. Trueness is usually expressed as bias [19].

Validation: Provision of objective evidence that a given item fulfils specified requirements, where the specified requirements are adequate for an intended use [20].



Within-laboratory reproducibility *also referred to as 'intermediate precision*': Precision obtained in the same laboratory under a set of within-laboratory conditions [10], i.e. conditions where test results are obtained with the same method, the same test sample, under some different operating conditions. E.g. different operators, different instruments, three replicates, three concentrations across three days/sequences in a specific laboratory.

NOTE: "Method" as used in this document can be considered synonymous with the term "procedure".

1.8. Guidance for Analytical Quality Assurance

1.8.1. Recommendations for Laboratories

Laboratories should be accredited by a recognised body operating in accordance with ISO/IEC 17011 [21] to ensure that they are applying analytical quality assurance. Methods shall be accredited following the ISO/IEC 17025 [20] standard.

Laboratories should demonstrate proficiency in the analysis of PFAS at the concentrations of interest (i.e. between the LOQ and e.g. 100x the LOQ and/or the range of legal limits) by validation, ongoing internal quality control and continuous successful participation in interlaboratory studies conducted by accredited bodies according to EN ISO/IEC 17043 [22], e.g. the EURL.

1.8.2. General Aspects regarding Sample Pre-Treatment and Storage

The samples must be stored and transported in containers that can be demonstrated to be free from the relevant PFAS (e.g. polypropylene/polyethylene containers) while preserving the integrity of the sample.

Sample quantity used for the extraction should be sufficient to fulfil the requirements with respect to a sufficiently low working range including the concentrations at the suggested LOQs.

The specific sample preparation procedures used for the products under consideration shall follow Commission Implementing Regulation (EU) 2022/1428 [1] for food. For feedingstuff no legal documents are available, but general aspects on sample preparation procedures can be found in e.g. Commission Regulation (EC) No 152/2009 [2].

1.8.3. General Aspects to Avoid Contamination / High blank Levels

Measures must be taken to avoid cross-contamination at each stage of the sampling and analysis procedure in the laboratory.

In the course of sampling and the preparation of the samples, precautions shall be taken to avoid any changes which would affect the content of PFAS, adversely affect the analytical determination or make the aggregate samples unrepresentative.



The person responsible for sampling should take the following precautions into account: do not wear clothing or gloves that contain fluoropolymer linings or that are treated with PFAS to improve water and stain repellence; do not use PFAS containing moisturizers, cosmetics, hand cream, sunscreens and related products at the sampling day.

Materials used during sampling, sample storage and sample transmission should be free of PFAS. Specifically, avoid the sample to be in contact with any fluoropolymer materials (e.g. PTFE, PVDF and others), such as fluoropolymer cutting boards, sampling containers, linings of caps of sampling containers. Avoid contact with other PFAS containing materials.

The analyst shall ensure that samples do not become contaminated during sample preparation by following the precautions described above. Furthermore wherever possible, the apparatus and equipment coming into contact with the sample shall not contain PFAS and shall be replaced by e.g. stainless steel, high density polyethylene (HDPE) or polypropylene parts. These should be cleaned with PFAS-free water and/or PFAS-free solvents and detergents to minimise the risk of contamination. [1]

The following (not exhaustive) list gives an overview of materials/consumables that may cause cross-contamination in the laboratory:

- PTFE products (e.g. PTFE lined vial caps)
- Aluminium foil
- TeflonTM and other fluoropolymer-containing materials
- Low density polyethylene (LDPE)
- Decon 90
- Gore-Tex®
- Lubricants during instrument maintenance

Reagents and other equipment used for analysis and sampling should be controlled to avoid possible introduction or loss of PFAS.

A reagent blank analysis should be performed by carrying out the entire analytical procedure without replacing the matrix by water. The levels in the reagent blanks should be monitored in each sequence of samples and over time by running a control chart. Further blanks (e.g. solvent blank, calibration blank) can optionally be monitored. They may help to check for potential contamination at different stages of the analysis.

1.8.4. General Quality Control Measures

Regular analysis of reagent blanks, and fortification experiments or control samples (preferably, using (certified) reference materials or in the absence of these, materials from successful interlaboratory studies) should be performed as internal quality control measures. Levels in reagent blanks, fortification experiments or control samples should be recorded in quality control (QC) charts and checked to make sure that the analytical performance is in accordance with the requirements. [23]



1.9. Validation Parameters

For routine analysis of PFAS in food and feed samples, laboratories should demonstrate the performance of the methodology during the validation procedure and during routine analysis. Performance should be demonstrated in a range from 1x the LOQ to e.g. 100x the LOQ and should cover legal limits (if available).

Table 2 provides recommendations of selected parameters for validation studies that should be carried out before routine analysis. Definitions can be found in section 1.7.

Table 2: Recommendations for validation studies and routine quality control measures

Grouping of matrices	 Use of different food or feed matrices to represent a matrix group if the matrices in the group share similar physico-chemical properties.
	Examples for matrix groups are given in Annex A of document No SANTE/12682/2019 for pesticides: e.g. milk and dairy products; meat (muscle) and seafood; etc. [15]
	 Further sub-grouping might be necessary for analytes and/or methods if significant matrix effects are observed.
Selectivity of the analytical procedure	Similarly, analytical methods should demonstrate the ability to reliably and consistently separate the analytes of interest from other co-extracted and possibly interfering compounds that may be present.
Trueness	The measurement process used must provide a valid estimate of the true concentration in a sample.
	Trueness can be estimated from regular analysis of certified reference materials, fortification experiments or participation in interlaboratory studies.
Precision	 Precision can be calculated from results generated under repeatability and within-laboratory reproducibility conditions.
Limit of quantification	Specific LOQs are given in Commission Recommendation (EU) 2022/1431 [24]. These may be revised in the future according to evaluations resulting from new toxicological studies and risk assessments.



1.10. Instrumentation

(Ultra) High performance liquid chromatography ((U)HPLC) coupled to low resolution or high resolution mass spectrometry (LRMS or HRMS) is recommended for analysis of the analytes under the scope.

1.10.1. LC-System

The LC-system must provide consistent sample injection volumes and be capable of performing binary linear gradients at a constant flow rate. PFAS may build up in PTFE transfer lines when the system is idle for more than one day. To prevent long delays in purging high levels of PFAS from the LC solvent lines, it may be useful to replace PTFE tubings with polyether ether ketone (PEEK) tubings and PTFE solvent frits with stainless steel frits. In addition a delay column can be installed before the injection valve to prevent the co-elution of PFAS originating from sources prior to the sample loop (e.g. mobile phase, fittings, tubes). Thorough rinse of the injection needle can reduce the co-elution of PFAS accumulated in the sample loop and valves.

1.10.2. Analytical Column

The laboratory may select the LC column. Based on previous experience a C₁₈ liquid chromatography column packed with solid phase particles is recommended (see Annex).

1.10.3. Mass Spectrometer

The mass spectrometer must be capable of electrospray ionization in the positive and negative ion mode. The system must be capable of producing specific product ions for analytes within specified retention time segments.



2. SPECIFIC REQUIREMENTS

2.1. Analytical Performance Criteria

Analytical performance criteria are an important element of quality assurance. These parameters provide information about the suitability of a method and the quality of the results. Common criteria are the basis for the comparability of results and methods between laboratories.

However, different control purposes require different performance criteria; i.e. substance/matrix combinations for which maximum levels (ML) are legally prescribed require more stringent criteria than substance/matrix combinations without existing ML. **Table 3** defines the criteria for the methods for PFAS analysis that shall be verified. Further explanation of each parameter is given in the following sections.

Table 3: Validation parameters and performance characteristics for PFAS analysed in samples for compliance testing of maximum levels or monitoring purposes

Parameter	Compliance testing of maximum levels ^{a)}	Monitoring purposes ^{b)}
Trueness ^{c)}	± 20 % [1]	± 35 %
Within-laboratory reproducibility (intermediate precision)	≤ 20 % [1]	≤ 25 %
LOQ	See 2.1.5 and Commission Recommendation (EU) 2022/1431 [24]	

^{a)} only for substance/matrix combinations <u>with</u> legally defined maximum levels

b) for substance/matrix combinations <u>without</u> legally defined maximum levels and for substance/matrix combinations <u>with</u> legally defined maximum levels in order to achieve the LOQs in section 2.1.5 for collecting occurrence data

^{c)} or expressed as apparent recovery (80-120% for compliance testing and 65-135% for monitoring purposes)

2.1.1. Trueness

Trueness can be estimated from a large series of analysis of certified reference materials, fortification experiments or participation in interlaboratory studies and shall be between -20 % and +20 % for compliance testing of maximum levels and should be between -35 % and +35 % for monitoring purposes (**Table 3**).

2.1.2. Apparent Recovery

QC samples should frequently be analysed as internal QC measures. The apparent recovery of PFAS in QC samples should be in the range of 80-120 % (compliance testing) and 65-135 % (monitoring purposes). Higher deviations for individual results might be accepted, if the criterion for trueness can be fulfilled.

2.1.3. Recovery

If a RS is available, the recovery of the added IS may conveniently be measured. For PFAS, the recoveries of the individual IS should be in the range of 30-140 %, reflecting what is currently achieved.



2.1.4. Precision

Precision can be calculated from results generated under repeatability and within-laboratory reproducibility conditions, e.g. derived from routine QC samples. Within-laboratory reproducibility should be ≤ 20 % for analysis of PFAS for compliance testing of maximum levels and ≤ 25 % for monitoring purposes (**Table 3**).

2.1.5. Limit of Quantification

For the LOQ estimation in PFAS analysis, the lowest validated level approach is recommended [1,25]. This means that the LOQ is the lowest successfully validated level of an analyte, for which it has been demonstrated that the respective criteria for identification (see 2.4.3), trueness and precision (see 2.1.1 and 2.1.4) are met.

According to Commission Implementing Regulation (EU) 2022/1428 Annex Part B Table 5 [1] the LOQ for PFOS, PFOA, PFNA and PFHxS shall be \leq the maximum level for the respective individual PFAS.

For the four individual PFAS (PFOS, PFOA, PFNA, PFHxS), indicative levels given in Commission Recommendation (EU) 2022/1431 Number 7 reflect maximum LOQs which should be achieved for certain matrices [24].

However, given that some food samples show concentrations even below these levels, target LOQs in the range of $0.001 - 0.050 \mu g/kg$ w.w. for the four individual PFAS (PFOS, PFOA, PFNA, PFHxS) are desirable (Commission Recommendation (EU) 2022/1431 Number 6) [24].

NOTE: Specific aims of the analytical method (as defined under 1.5) may call for even lower LOQs, particularly for exposure assessment.

The estimation of LOQs requires consideration of the reagent blanks as follows:

- For compliance testing of maximum levels: Contribution of reagent blank levels should be ≤ 30% of the target LOQ. Higher contribution of more than 30% requires the inclusion of reagent blank levels in the estimation of LOQs. As mentioned in sections 1.8.3 and 1.8.4 reagent blank levels should be monitored in each sequence of sample analysis and should be recorded in a QC chart.
- For monitoring purposes: In the case where due to elevated reagent blank levels the lowest possible target LOQ cannot be achieved by using the lowest validated level approach, the LOQ can be estimated using 3.3x the average reagent blank concentration (see **Table 4**).



	Imaginary reagent blank values* (µg/kg)	LOQ calculation	Resulting LOQ (µg/kg)
Reagent blank 1	0.028		
Reagent blank 2	0.033	3.3x average	
Reagent blank 3	0.048	reagent blank	0.109
Reagent blank 4	0.026	reagent blank	
Reagent blank 5	0.029		
Average (µg/kg)	0.033		
Standard deviation (µg/kg)	0.009		

Table 4: Estimation of the limit of quantification (LOQ) to achieve low LOQs for e.g. monitoring purposes

*It is recommended to use at least five reagent blanks to calculate the average reagent blank value. These values can either be determined during validation (to estimate the LOQ) or from the QC chart (to monitor the LOQ in routine analysis).

2.1.5.1. Branched Isomers

The LOQs for all components that are separately measured (i.e. sum of branched isomers, linear isomer) should be determined. In practice, there are two cases (see 2.3.1 and 2.3.2):

- a) Quantification of branched isomers using an analytical standard containing branched and linear isomers: individual LOQs for sum of branched isomers and linear isomer should be determined;
- b) Quantification of branched isomers against linear isomer: the same LOQ as for the linear isomer can be applied.

2.1.5.2. Sum Parameters

For sum parameters, such as the sum of PFOS, PFOA, PFNA, and PFHxS, a sum LOQ is not needed for compliance testing [1]. This is also applicable to total-PFOS, total-PFNA, total-PFHxS and total-PFOA.

2.2. Method Validation

As described in section 1.9, prior to routine analysis of PFAS in food and feed samples, laboratories should demonstrate the performance of their methodology during the validation procedure. Within-laboratory method validation is essential to provide evidence that the method is fit for the intended purpose. **Table 5** summarizes the parameters and criteria which shall be verified during method validation. An example of a practical approach to the validation procedure (minimum requirements) is given below.

Table 5: Validation parameters and criteria

Parameter	Description	Criterion	Cross ref.
Linearity	Linearity check from five calibration levels	Deviation of back- calculated concentration from true concentration ≤ 20 %	-
Trueness	Average apparent recovery for each fortification level tested; expressed as 'bias'	Table 3	Section 2.1
Precision	Within-laboratory reproducibility for each fortification level tested	Table 3	Section 2.1
LOQ	Lowest fortification level meeting the identification requirements and analytical performance criteria for trueness and precision	-	Section 2.1.5
Ion ratio, retention time	Check compliance with identification requirements for MS techniques	Table 8	Section 2.4.3

Validation needs to be performed for all analytes within the scope of the method. Validation can be performed for a single matrix (see 2.2.1) or for a matrix group (see 2.2.2) within the scope of the method.

2.2.1. Example Approach: Single Matrix

General overview:

- Duration: ≥ 2 days
- Operator: \geq 1 technician
- Matrix: 1 matrix (e.g. pork meat) with \geq 5 different batches
- Reagent blank: ≥ 1 per sequence
- Sample set per matrix batch:
 - o 1 sample blank
 - o 1 fortified sample at 1x the target LOQ
 - $_{\odot}$ 1 fortified sample at one other higher level e.g. in the range 2-50x the target LOQ
 - $\circ~$ 1 fortified sample at one other higher level e.g. in the range 50-100x the target LOQ
 - o Additional samples and fortification levels optional

NOTE: The range of fortification levels should cover achievable LOQs and, if available MLs, of all analytes within the scope of the method.



Specific overview of sample set:

- Prepare a set of aliquots (n ≥ 4) of the selected batches of test material, i.e. ≥ 5 different batches of e.g. pork meat (see **Table 6**).
- Fortify half of the test material batches (in case of an uneven number, chose a number just below or above the average) with the analytes at 1x the target LOQ and at least two other higher levels e.g. in the range of 2-100x the target LOQ. One aliquot per batch remains unfortified (= blank).
- Perform the analysis.
- Calculate the concentration detected in each sample.
- Repeat these steps on at least one other day with the rest of test material batches, different operators (if possible) and as many different environmental conditions as possible, e.g. different batches of reagents, solvents or a variation of other parameters (Table 6).
- Determine the mean concentration, standard deviation and the coefficient of variation (%) of the fortified samples for each fortification level tested.
- Evaluate the parameters from **Table 5** and verify them against the criteria.

	Batch 1	Batch 2	Batch 3	Batch 4	Batch 5
Test material	Pork meat 1	Pork meat 2	Pork meat 3	Pork meat 4	Pork meat 5
Operator	Technician A	Technician A	Technician A (or B)	Technician A (or B)	Technician A (or B)
Day	1	1	1 or 2	2	2
Fortification levels	1x target LOQ				
	E.g. 5x target LOQ				
	E.g. 50x target LOQ				
Number of replicates per fortification level and operator	1	1	1	1	1
Number of sample blanks per operator	1	1	1	1	1
Number of reagent blanks per sequence			≥ 1		

Table 6: Example of a validation sample set if validation is performed for one matrix (e.g. pork meat)

NOTE: Additional batches of test material, replicates, days, operators and fortification levels optional.



Instrumental sample sequence:

- Calibration standards
- Reagent blanks
- Sample blanks
- Fortified samples
- Calibration standards

2.2.2. Example Approach: Matrix Group

General overview:

- Duration: ≥ 2 days
- Operator: ≥ 2 technicians (if possible)
- Matrix: ≥ 5 different matrices of one matrix group (see **Table 2** in section 1.9)
- Reagent blank: ≥ 1 per sequence

Sample set **per matrix batch**:

- o 2 sample blanks
- 2 fortified samples at 1x the target LOQ
- 2 fortified samples at one other higher level e.g. in the range 2-50x the target LOQ
- 2 fortified samples at one other higher level e.g. in the range 50-100x the target LOQ
- Additional samples and fortification levels optional

NOTE: The range of fortification levels should cover achievable LOQs and, if available MLs, of all analytes within the scope of the method.

Specific overview of sample set:

- Prepare a set of aliquots (n ≥ 4) of the selected batches of test material, i.e. ≥ 5 different batches of test material (e.g. pork meat, lamb meat, salmon muscle, plaice muscle, and bovine meat) (see
- **Table 7**).
- Fortify all test material batches with the analytes at 1x the target LOQ and at least two other higher levels e.g. in the range of 2-100x the target LOQ. One aliquot per batch remains unfortified (= blank).
- Perform the analysis.
- Calculate the concentration detected in each sample.
- Repeat these steps for all (n ≥ 5) batches of test material on at least one other day and with a different operator (if possible) and as many different environmental



conditions as possible, e.g. different batches of reagents, solvents or a variation of other parameters (

- **Table 7**).
- Determine the mean concentration, standard deviation and the coefficient of variation (%) of the fortified samples for each fortification level tested.
- Evaluate the parameters from **Table 5** and verify them against the criteria.

Table 7: Example of a validation sample set if validation is performed for a matrix group (e.g. meat (muscle) and seafood)

	Batch 1	Batch 2	Batch 3	Batch 4	Batch 5
Test material	Pork meat	Lamb meat	Salmon muscle	Plaice muscle	Bovine meat
Operator	Technician A and B				
Day	1 and 2				
Fortification levels	1x target LOQ				
	E.g. 5x target LOQ				
	E.g. 50x target LOQ				
Number of replicates per fortification level and operator	1	1	1	1	1
Number of sample blanks per operator	1	1	1	1	1
Number of reagent blanks per sequence			≥ 1		

NOTE: Additional batches of test material, replicates, days, operators and fortification levels optional.

Instrumental sample sequence:

- Calibration standards
- Reagent blanks
- Sample blanks
- Fortified samples
- Calibration standards



2.3. Quantification

In order to validate the complete analytical procedure the addition of IS should be carried out at the very beginning of the analytical method e.g. prior to extraction.

It is preferable that isotope-labelled IS (ILIS) of at least the four main compounds (**Table 1**) should be used as this would improve the reliability of the quantitation.

If additional PFAS (for which no ILIS are available) are being simultaneously determined, relative response factors shall be determined using appropriate¹ isotope-labelled PFAS. The validity of these can be confirmed either by using appropriate calibration solutions or by reference materials.

2.3.1. PFOS

1-PFOS often co-elutes with L-PFOS on C_{18} columns and contributes only 1% to the br-PFOSK² standard. The concentration of di-methyl-substituted branched isomers is believed to be negligible low. For identification of the retention time of br-PFOS it is recommended to measure a native PFOS standard that contains a mixture of br-PFOS and linear PFOS (e.g. br-PFOSK) with each sequence.

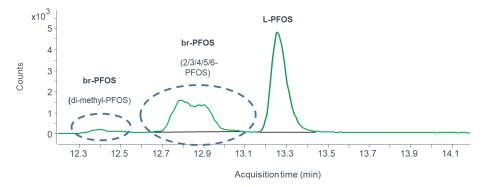


Figure 1: Extracted-ion chromatogram (mass transition $m/z 499 \rightarrow 80$) of linear PFOS (L-PFOS) and its branched isomers (br-PFOS) of a technical PFOS standard.

Quantification of total-PFOS should include L-PFOS and br-PFOS (**Figure 1**). Total-PFOS may be determined using L-PFOS if br-PFOS is not separated chromatographically from L-PFOS. If br-PFOS is separated chromatographically from L-PFOS, br-PFOS may be quantified using one of the following approaches:

- a) Quantification of br-PFOS against br-PFOS by using an analytical PFOS standard that contains a mixture of br-PFOS and linear PFOS (e.g. br-PFOSK) in a known ratio,
- b) Quantification of br-PFOS against L-PFOS by using a linear PFOS standard and calculation of the average of the two mass transitions $m/z 499 \rightarrow 99$ and $m/z 499 \rightarrow 80$ if fragment ions are used for quantification (typically LRMS),
- c) Quantification of br-PFOS against L-PFOS using a linear PFOS standard if the ion m/z 499 is used for quantification (typically HRMS).

¹ e.g. the chromatographically closest available ILIS

² Potassium perfluorooctanesulfonate (linear and branched isomers), CAS-number not available



These quantification procedures are regarded by the authors as the most pragmatic and accurate approaches in view of challenges in method development. In addition, a comparative study among the members of the CWG PFAS has shown that the results of these approaches do not statistically differ from each other [26]. Furthermore, results obtained by Orbitrap have shown that the ion m/z 499 has an equal response for L-PFOS and br-PFOS and can thus be used for quantification by Orbitrap. Users of other HRMS systems are advised to test if their instruments provide similar results.

2.3.2. PFOA, PFHxS, PFNA

Quantification of PFOA, PFHxS, and PFNA should include linear and branched isomers. The approaches used for quantification of PFOS (see 2.3.1) also apply to PFOA, PFHxS and PFNA. Analytical standards containing linear and branched isomers of PFOA, PFHxS and PFNA are commercially available.

NOTE: Branched isomers of PFOA, PFHxS and PFNA are less prevalent and can be less frequently detected in food samples than branched isomers of PFOS.

2.4. Measurement

2.4.1. Exclusion of Interfering Substances

Separation of PFAS from interfering (e.g. taurodeoxycholic acid) or other possible coeluting interfering substances should be carried out by suitable sample preparation methods and/or chromatography/mass spectrometry techniques.

NOTE: Taurodeoxycholic acid (TDCA) is an endogenous compound which is formed in liver cells and normally found in matrices of animal origin (mainly eggs and offal). Under typical C₁₈ LC column separation conditions it elutes at the same retention time and shares the same mass transition as PFOS (499 \rightarrow 80). This may lead to false positive identification or over-reporting of the PFOS concentration. Therefore, TDCA should be removed by suitable sample preparation methods using ENVI-Carb or suitable LC separation methods (e.g. FluoroSep RP Octyl column). [27,28] Alternatively, the interference-free 499 \rightarrow 99 mass transition can be used for quantification of PFOS. This mass transition is, however, less sensitive than the 499 \rightarrow 80 transition and thus leading to a higher LOQ for PFOS [28].

In addition, interfering substances (i.e. fatty acids) have been reported for PFBA in e.g. tissue and PFPeA in e.g. shellfish and hot cocoa [29,30].

2.4.2. Analytical Calibration Curve

The lower range of the calibration curve is indicated by the LOQ (or target LOQ) for PFAS. This should extend to between 5.0 and 50 μ g/kg at the higher end of range, reflecting the concentrations for PFAS that are reported in the current literature or established legal limits. At least five calibration concentrations are required to prepare the initial calibration curve spanning the expected concentration range. If the calibration curve spans several orders of magnitude the use of weighting factors (e.g. 1/x) is recommended.



2.4.3. Identification Requirements

In **Table 8**, mass spectrometric performance and peak identification criteria for reliable analysis of PFAS are listed. Further identification and confirmation criteria are described in internationally standardized methods for e.g. PFAS in drinking water [31] and pesticides [15].

Table 8: Mass spectrometric performance and peak identification criteria for different LC-MS techniques

(U)HPLC-LRMS	Unit mass resolution		
Typical systems (examples)	MS/MS triple quadrupole, ion trap, Q-trap		
Acquisition	E.g. selected or multiple reaction monitoring (SRM, MRM)		
Minimum number of ions	2 product ions		
Ion ratio	Ion ratio from sample extracts should be within \pm 30 % (relative) of		
1011 1410	average of calibration standards from same sequence ^{e)} .		
Signal to noise (S/N) ratio	≥ 3		
	The ratio of the chromatographic RT of the analyte to that of the IS (i.e.		
Retention time (RT)	relative RT of the analyte) shall correspond to that of the calibration		
	standard with a maximum deviation of 1 %. (NOTE: Only applicable for		
	analytes with an isotopically labelled analogue.)		
Other	Analyte peaks from both transitions in the extracted ion		
Other	chromatograms must fully overlap.		
(U)HPLC-HRMS	Accurate mass resolution		
(U)HPLC-HRMS Typical systems			
Typical systems (examples)	Accurate mass resolution High resolution MS: (Q-)TOF, (Q-)Orbitrap		
Typical systems			
Typical systems (examples) Mass resolution	High resolution MS: (Q-)TOF, (Q-)Orbitrap		
Typical systems (examples)	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range)		
Typical systems (examples) Mass resolution	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction		
Typical systems (examples) Mass resolution Acquisition	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ²		
Typical systems (examples) Mass resolution Acquisition Minimum number of ions	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ² 2 ions with mass accuracy ≤ 5 ppm ^{f), g)}		
Typical systems (examples) Mass resolution Acquisition Minimum number of ions Signal to noise (S/N) ratio	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ² 2 ions with mass accuracy ≤ 5 ppm ^{f), g)} ≥ 3		
Typical systems (examples) Mass resolution Acquisition Minimum number of ions	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ² 2 ions with mass accuracy ≤ 5 ppm ^{f), g)} ≥ 3 The ratio of the chromatographic RT of the analyte to that of the IS (i.e.		
Typical systems (examples) Mass resolution Acquisition Minimum number of ions Signal to noise (S/N) ratio	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ² 2 ions with mass accuracy ≤ 5 ppm ^{f), g)} ≥ 3 The ratio of the chromatographic RT of the analyte to that of the IS (i.e. relative RT of the analyte) shall correspond to that of the calibration		
Typical systems (examples) Mass resolution Acquisition Minimum number of ions Signal to noise (S/N) ratio	High resolution MS: (Q-)TOF, (Q-)Orbitrap ≥ 10 000 at 10 % valley (for the entire mass range) E.g. full scan, all ion fragmentation (AIF)/MS ^E , parallel reaction monitoring (PRM), data-dependent MS ² 2 ions with mass accuracy ≤ 5 ppm ^{f), g)} ≥ 3 The ratio of the chromatographic RT of the analyte to that of the IS (i.e. relative RT of the analyte) shall correspond to that of the calibration standard with a maximum deviation of 1 %. (NOTE: Only applicable for		

 e) applying identical MS/MS conditions, in particular collision energy and collision gas pressure, for each transition of an analyte

^{f)} preferably including the molecular ion, (de)protonated molecule or adduct ion <u>and</u> at least one fragment ion

^{g)} <1 mDa for m/z <200

NOTE: PFAS with only one specific MS/MS transition (e.g. PFBA, PFPeA) should be verified using a second chromatographic separation method (i.e. use of a secondary LC elution on a different analytical column and eluent) or another MS method (e.g. the use of high resolution MS).



2.5. Reporting of Results

The concentrations determined in test samples shall be expressed in units of μ g/kg wet weight for food or in μ g/kg product for feed (optionally, relative to a feed with a moisture content of 12 %). Results shall be reported as anions³ or neutral compounds⁴, respectively, and to two significant figures (see 2.5.1).

The uncertainty of measurement (see 2.5.2) should also be included as an aid to the interpretation of the data. The analytical results shall be reported as $x \pm U$ whereby x is the analytical result and U is the expanded measurement uncertainty using a coverage factor of 2 which gives a level of confidence of approximately 95 %.

PFOS should be reported as "total-PFOS" and additionally as L-PFOS and br-PFOS, if possible.

All target analytes should be reported as individual concentrations and PFOA, PFOS, PFNA and PFHxS additionally as lower bound summed concentration (∑PFOA, PFNA, PFOS, PFHxS).

NOTE: Only concentrations of PFAS, which have been confirmed by at least two MS/MS transitions should be reported. Otherwise a different chromatographic method or another MS method should be used to confirm the result (see 2.4.3). If this is not possible, the information that the result is not sufficiently confirmed must be included in the reporting format⁵.

Additional information that should (optionally) be included in the report:

- Information on the methods used for extraction and purification for PFAS should be included – this information can be basic, e.g. mention of the techniques used in the analysis.
- As an aid to the evaluation of the reported data, the recoveries of the individual internal standards can be included.
- Data for feed may also additionally be reported as µg/kg product relative to a feed with a moisture content of 12 % along with the determined moisture content.

2.5.1. Rounding of Results

Results shall be rounded to two significant figures.

The following general rules are proposed for rounding the result:

- a) If the digit following the digit to be rounded in the primary result is less than 5 (0, 1, 2, 3, 4), the previous digit will not change.
- b) If the digit following the digit to be rounded in the primary result is 5 or more (5, 6, 7, 8, 9), round the previous digit up by one unit.

³ if they exist as anions (e.g. PFCA, PFSA)

⁴ if they do not exist as anions (e.g. FOSA)

⁵ For reporting to EFSA the level of identification confirmation, the open text within the variable 'anmethText' can be used to report additional information regarding the analytical method or analysis.



- c) The expanded measurement uncertainty will be estimated by using the final rounded result.
- d) The value of the expanded uncertainty is always rounded up unless (after rounding of the second non-retained digit) the first non-retained digit would be 0. The value of the expanded uncertainty should be given with the same number of decimals as the rounded result.

Example:

- 1. Primary result = $0.5678 \,\mu g/kg$
- 2. Primary result rounded to two significant figures = **0.57 µg/kg** (final result)
- 3. Primary value for the expanded measurement uncertainty (e.g. 30 %) = 0.57 x 0.3 = $0.171 \mu g/kg$
- 4. Rounded value of the expanded measurement uncertainty = 0.18 μg/kg (two significant figures)
- 5. Reported result = $0.57 \ \mu g/kg \pm 0.18 \ \mu g/kg$ (k = 2; 95 %)

2.5.2. Measurement Uncertainty

As a first estimation for the individual combined uncertainty u the within-laboratory reproducibility standard deviation may be used. However, if possible, the uncertainty of the bias (after correction for a constant laboratory bias) should be included in the measurement uncertainty estimation, which can be derived from a) analysis of certified reference materials, b) participation in proficiency tests or c) fortification experiments. Further information on estimation can be found in [32–34].

According to Commission Implementing Regulation (EU) 2022/1428 a coverage factor of 2, which gives a level of confidence of approximately 95%, is used for the expanded uncertainty U [1].

The reporting of sum parameters and the possible comparison with legal limits requires the additional estimation of combined uncertainty u and expanded measurement uncertainty U for these sum parameters. For PFAS this is the case for the sum of PFOS, PFOA, PFNA and PFHxS and for e.g. total-PFOS, if calculated as the sum of L- and br-PFOS. [1]

In these cases the calculation of the combined uncertainty u of the sum parameter (as an absolute value) is calculated as the square root of the sum of squares of the individual absolute values of combined uncertainties (equation 2). For these calculations the rounded results and uncertainties of the individual substances can be used. Further rounding is then performed according to section 2.5.1.

$$u_{sum(abs)} = \sqrt{\sum_{i=1}^{n} u_{i(abs)}^2}$$

(**Eq. 2**)

with:

 $u_{sum(abs)}$ = absolute combined measurement uncertainty of sum parameter $u_{i(abs)}$ = absolute combined measurement uncertainty of individual parameters i



The expanded uncertainty for the sum parameter is then dependent on the contents of the individual substances contributing to the sum and cannot be standardised for the sum parameters.

Example:

Table 9: Example for calculation of measurement uncertainty for sum of PFOS, PFOA, PFNA and PFHxS

	Sum of PFOS	PFOA	PFNA	PFHxS
Content (rounded) [µg/kg]	5.5	0.66	0.35	0.12
Relative expanded uncertainty U _(rel)	20 %	20 %	20 %	20 %
Relative combined uncertainty u _(rel)	10 %	10 %	10 %	10 %
Absolute combined uncertainty u _(abs) (rounded) [µg/kg]	0.55	0.066	0.035	0.012

Content_{sum} = 5.5 + 0.66 + 0.35 + 0.12 = 6.6 [µg/kg] (rounded to two significant figures)

 $u_{sum(abs)} = \sqrt{0.55^2 + 0.066^2 + 0.035^2 + 0.012^2} = 0.555180 [\mu g/kg]$

 $U_{sum(abs)} = 0.555180 \times 2 = 1.11036 [\mu g/kg]$

 $U_{sum(abs)} = 1.1 [\mu g/kg]$ (rounded to two significant figures)

Reported result for sum of PFOS, PFOA, PFNA, PFHxS = $6.6 \mu g/kg \pm 1.1 \mu g/kg$ (k = 2; 95 %) Relative expanded uncertainty U for sum of PFOS, PFOA, PFNA, PFHxS: 18 %



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