

## Supporting Information

### How well do product labels indicate the presence of PFAS in consumer items used by children and adolescents?

Kathryn M. Rodgers<sup>1</sup>, Christopher H. Swartz<sup>1</sup>, James Occhialini<sup>2</sup>, Philip Bassignani<sup>2</sup>, Michelle McCurdy<sup>3</sup>, Laurel A. Schaidler<sup>1\*</sup>

<sup>1</sup>Silent Spring Institute, 320 Nevada Street, Suite 302, Newton, MA 02460, USA

<sup>2</sup>Alpha Analytical Laboratories, 320 Forbes Blvd, Mansfield, MA 02048, USA

<sup>3</sup>Galbraith Laboratories, Inc. 2323 Sycamore Drive, Knoxville, TN 37921, USA

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### **Standards and reagents**

For targeted PFAS analysis via LC/MS/MS, all native and isotopically labeled standards were purchased from Wellington Laboratories (ON, Canada). Reagents included methanol  $\geq 99.9\%$  (B&J Brand™) for HPLC and CHROMASOLV™ (Honeywell Research Chemicals) for LC-MS, 1M ammonium acetate at pH 5.0 (Waters), glacial acetic acid (BDH Chemicals), ammonium hydroxide (BDH Chemicals), sodium acetate buffer (Alfa Aesar), NaOH (BDH Chemicals), potassium persulfate (BDH Chemicals), 6N HCl (GFS Chemicals), and Ottawa sand (BDH Chemicals). Extraction media for sample preparation included Strata™-X-AW 33  $\mu\text{m}$  Polymeric Weak Anion, 500 mg cartridges and Strata GCB, 250 mg cartridges (Phenomenex). Sample preparation apparatus included SCP Digi Tubes (SCP Science), 25 mm 0.2- $\mu\text{m}$  syringe filters (Whatman), 5-mL PP syringes (Thermo Fisher), 50-mL centrifuge tubes (Bio Express), 700- $\mu\text{l}$  PP LC vials (Waters), 4-mL PP storage vials (Thermo Fisher), and vacuum manifolds (Waters).

### **LC/MS/MS procedures for PFAS analysis**

Extract cleanup was performed using a solid phase extraction (SPE) cartridge containing a mixed mode, weak anion exchange (WAX), reversed phase stacked onto a 250 mg graphitized carbon black cartridge. Cartridges were pre-conditioned by rinsing with 15 mL of methanol containing 2% ammonium hydroxide. Five mL of the methanol extract was transferred to an SPE cartridge and allowed to pass through it by gravity feed at a dropwise rate, ensuring adequate contact time with the cartridge sorbent. The cartridge was then rinsed with an additional 5 mL of methanol. Vacuum was applied only if the flow of solvent through the cartridge stopped. Eluates were concentrated to dryness under a gentle stream of nitrogen in a heated water bath (60-65 °C). 20  $\mu\text{L}$  of the isotope dilution recovery primary dilution standard was added to the collection vial, and the appropriate amount of 80:20% (vol/vol) methanol:water solution was added and to bring

the volume to 1 mL. The extract was then vortexed and two aliquots were each transferred with a plastic pipette into each of two polypropylene autosampler vials. Due to the possible volatility and suspect degradation of sulfonamides and sulfonamide ethanols when exposed to heat, a portion of the eluate was retained and analyzed independently with no evaporation for these analytes.

Individual PFAS were quantified in extracts using a Waters Acquity H class HPLC equipped with an LC BEH C<sub>18</sub> column (2.1 x 50 mm) packed with 1.7 µm d<sub>p</sub> C<sub>18</sub> solid phase particles coupled to a Waters Xevo TQ-S micro operating in the MS/MS mode. The mobile phases consisted of 2 mM ammonium acetate in 95:5 methanol water (mobile phase A) and methanol (mobile phase B).

### **Quality assurance/quality control (QA/QC) methods and results**

#### *Total fluorine*

Our quality control methods were informed by guidance for analysis of environmental data.<sup>1</sup> Product samples and replicates for six products (Table S6) were analyzed for total fluorine in eight batches, and the laboratory was blind to the identity of the replicate samples. Method blank samples were run after at least every 10 samples, as well as the beginning and end of each batch. Percent recovery ranged between 99-101%.

The limit of detection (LOD) for each total F measurement (in ppm) was calculated using the following equation:

$$LOD = \frac{((C_{std} \times DF) - C_{blank}) \times V_{sp} \times 1000}{M_s}$$

where:

C<sub>std</sub> = concentration of the lowest calibration standard, typically 0.1 or 0.5 mg/L

DF = dilution factor for the sample

C<sub>blank</sub> = concentration of the batch blank (mg/L)

V<sub>sp</sub> = sample preparation volume (amount of buffer used for sample), typically 20 or 100 mL

M<sub>s</sub> = sample mass, typically 200 mg

The calibration curve was checked using a potassium fluoride (KF) standard solution. For the calibration to be considered acceptable for a given batch, the calculated concentration was required to be within 90-110% of the expected value.

A diluted p-fluorobenzoic acid standard was also used to check the calibration during each run and to monitor drift. This standard was run through the same preparation as the samples and the analysis criteria required 96.8–103.2% recovery of the standard for the calibration to remain valid. This standard was used for the low calibration specifically.

#### *Target PFAS and TOP assay analyses via LC/MS/MS*

Sample analysis for 36 target PFAS analytes was conducted in four batches, using LC/MS/MS with isotope dilution. Extracted Internal Standards (EIS) were spiked into pre-extracted samples and carried through the entire analytical process. Recoveries of extracted internal standards were used to assess extraction efficiency of the analytic method for each target analyte. Surrogate recovery standards were spiked into extracts after blow down. The concentrations of these recovery standards were determined using a calibration curve developed from a set of standards run separately. Recoveries for the surrogate standards were then used to correct the concentrations of the extracted internal standards. The sample concentrations were then calculated using the extracted internal standard recoveries.

For the TOP analysis, each extract was spiked with 20 µL of TOP pre-assay surrogates containing five negative control surrogates (to assess complete oxidation of precursors) and three positive control surrogates (to assess complete formation of terminal products). Positive control recoveries were within 50-150% and were considered adequate. Because TOP is based on oxidation of unknown precursors, we were not able to evaluate recoveries of all possible precursor analytes. We did observe that 13 of 18 internal standard surrogates run in the TOP assay had recoveries <50% in more than half of the samples that the 18 surrogates were measured in, indicating that measured PFAA precursor concentrations may have been biased low.

Method Detection Limits (MDL) (or LOD) for each analyte were determined by extracting and analyzing a minimum of seven low-level samples spiked at or just below the lowest calibration standard. The MDL is a statistically calculated value determined from the standard deviation of the spiked samples multiplied by a statistical constant student-t value at 99% confidence. The Reporting Limit (RL) is equal to the Limit of Quantitation (LOQ), which is set to the lowest calibration standard.

Precision and accuracy of the analytical method were evaluated using laboratory control samples. One field blank (10 mL of methanol with 2% ammonium hydroxide added to an empty centrifuge tube) was analyzed, and all target analyte concentrations were <LOD. Field blank surrogate recoveries were consistent with the surrogate recoveries for samples. Four to six method blanks were run for each target analyte in each of the four batches, and among these, only N-ethyl perfluorooctanesulfonamidoacetic acid (NEtFOSAA), perfluorobutanoic acid (PFBA), perfluorohexadecanoic acid (PFHxDA), perfluorohexanoic acid (PFHxA), perfluorooctanoic acid (PFOA), and perfluorotetradecanoic acid (PFTA) were detected, all below 1 ng/g. Because none of method blanks had any PFAS detected above the MRL, and following guidance from Udesky et al. on blank correction procedures for multiple blank detections,<sup>1</sup> we did not blank correct or censor the data.

Duplicate samples were analyzed for each of five products, with each duplicate conducted in a different analysis batch. The laboratory was blind to the identity of the duplicate samples. For each duplicate pair of a detected target analyte, the relative percent different (RPD) was below 20%, except for PFTrDA, PFPeS, and NMeFOSE. The measurements for each sample replicate are reported in Table S5.

**Table S2.** Reporting limits, method detection limits, and method reporting limits for PFAS target analytes and terminal PFAAs formed by the Total Oxidizable Precursor (TOP) assay. All concentrations in ng/g.

Analyte	TOP	CAS number	RL <sup>a</sup>	MDL <sup>a</sup>	MRL range <sup>a</sup>	MRL range (TOP)
Perfluorobutanoic acid (PFBA)	✓	375-22-4	0.5	0.0227	1.11-26.6	1.11-18.7
Perfluoropentanoic acid (PFPeA)	✓	2706-90-3	0.5	0.046	1.11-26.6	1.11-18.7
Perfluorohexanoic acid (PFHxA)	✓	307-24-4	0.5	0.0525	1.11-26.6	1.11-18.7
Perfluoroheptanoic acid (PFHpA)	✓	375-85-9	0.25	0.0451	1.11-26.6	1.11-18.7
Perfluorooctanoic acid (PFOA)	✓	335-67-1	0.25	0.0419	1.11-26.6	1.11-18.7
Perfluorononanoic acid (PFNA)	✓	375-95-1	0.25	0.075	1.11-26.6	1.11-18.7
Perfluorodecanoic acid (PFDA)	✓	335-76-2	0.25	0.067	1.11-26.6	1.11-18.7
Perfluoroundecanoic acid (PFUnA)	✓	2058-94-8	0.5	0.0468	1.11-26.6	1.11-18.7
Perfluorododecanoic acid (PFDoA)	✓	307-55-1	0.5	0.07	1.11-26.6	1.11-18.7
Perfluorotridecanoic acid (PFTrDA)	✓	72629-94-8	0.5	0.2045	1.11-26.6	1.11-18.7
Perfluorotetradecanoic acid (PFTA)	✓	376-06-7	0.5	0.054	1.11-26.6	1.11-18.7
Perfluorohexadecanoic acid (PFHxDA)		67905-19-5	2	0.12	2.22-53.2	
Perfluorooctadecanoic acid (PFODA)		16517-11-6	2	0.171	2.22-53.2	
Perfluorobutanesulfonic acid (PFBS)	✓	375-73-5	0.25	0.039	1.11-26.6	1.11-18.7
Perfluoropentanesulfonic acid (PFPeS)	✓	2706-91-4	1	0.0835	1.11-26.6	1.11-18.7
Perfluorohexanesulfonic acid (PFHxS)	✓	355-46-4	0.25	0.0605	1.11-26.6	1.11-18.7
Perfluoroheptanesulfonic acid (PFHpS)	✓	375-92-8	0.5	0.1365	1.11-26.6	1.11-18.7
Perfluorooctanesulfonic acid (PFOS)	✓	1763-23-1	0.25	0.13	1.11-26.6	1.11-18.7
Perfluorononanesulfonic acid (PFNS)	✓	68259-12-1	1	0.299	1.11-26.6	1.11-18.7
Perfluorodecanesulfonic acid (PFDS)	✓	335-77-3	0.5	0.153	1.11-129	1.11-18.7

Perfluorododecane sulfonic acid (PFDoDS)		79780-39-5	1	0.086	1.11-26.6	
1H,1H,2H,2H-perfluorohexanesulfonic acid (4:2 FTS)		757124-72-4	1	0.0645	1.11-26.6	
1H,1H,2H,2H-perfluorooctanesulfonic acid (6:2 FTS)		27619-97-2	0.5	0.1795	1.11-26.6	
1H,1H,2H,2H-Perfluorodecanesulfonic acid (8:2 FTS)		39108-34-4	0.5	0.287	1.11-26.6	
1H,1H,2H,2H-Perfluorododecanesulfonic acid (10:2 FTS)		120226-60-0	1	0.275	1.11-26.6	
Perfluorooctanesulfonamide (FOSA or PFOSA)		754-91-6	0.5	0.098	1.11-26.6	
N-Methyl perfluorooctane sulfonamide (NMeFOSA)		31506-32-8	1	0.379	1.11-19	
N-Ethyl perfluorooctane sulfonamide (NEtFOSA)		4151-50-2	1	0.407	1.11-19	
N-Methyl perfluorooctanesulfonamidoacetic acid (NMeFOSAA)		2355-31-9	0.5	0.2015	1.11-26.6	
N-Ethyl perfluorooctanesulfonamidoacetic acid (NEtFOSAA)		2991-50-6	0.5	0.0845	1.11-26.6	
N-Methyl perfluorooctane sulfonamidoethanol (NMeFOSE)		24448-09-7	2	0.52	2.23-39	
N-Ethyl perfluorooctane sulfonamidoethanol (NEtFOSE)		1691-99-2	2	0.73	2.22-39	
2,3,3,3-Tetrafluoro-2-[1,1,2,2,3,3,3-heptafluoropropoxy]-propanoic acid (HFPO-DA)		13252-13-6	10	3.81	1.11-266	
4,8-Dioxa-3h-perfluorononanoic acid (ADONA)		919005-14-4	1	0.0413	1.11-26.6	
9-Chlorohexadecafluoro-3-oxanone-1-sulfonic acid (9Cl-PF3ONS)		756426-58-1	1	0.0374	1.11-26.6	
11-Chloroeicosafluoro-3-oxaundecane-1-sulfonic acid (11Cl-PF3OUdS)		763051-92-9	1	0.0388	1.11-26.6	

<sup>a</sup> Applies to target PFAS. RL=reporting limit, MDL=method detection limit, MRL=method reporting limit

**Table S3.** Mass spectrometer settings

<b>ESI Conditions</b>	
Polarity	Negative ion
Capillary needle voltage	0.5 kV
Cone gas flow	25 L/hr
Nitrogen desolvation gas	1000 L/hr
Desolvation gas temperature	500 °C

**Table S4.** Liquid chromatography settings

<b>Time (min)</b>	<b>2 mM Ammonium acetate (5:95 MeOH/H<sub>2</sub>O)</b>	<b>100% Methanol</b>
Initial	100	0
1.0	100	0
2.2	85	15
11	20	80
11.4	0	100
12.4	100	0
15.5	100	0



**Table S5.** Target analyte, extracted internal standard, and surrogate standard quantitation parameters.

Analyte	Type	Quantitation transition	Qualifier transition	IS	CV	CE
PFBA	target analyte	213 > 169	N/A	3: M4PFBA	20	8
PFPeA	target analyte	263 > 219	N/A	5: M5PFPEA	18	8
PFHxA	target analyte	313 > 269	313>119	11: M5PFHxA	20	10
PFHpA	target analyte	363 > 319	363>169	14: M4PFHpA	18	8
L-PFOA	target analyte	413 > 369	413>219	23: M8PFOA	11	9
br-PFOA	target analyte	413 > 369	413>219	23: M8PFOA	11	9
PFOA (total)	target analyte	413 > 369	413>219	23: M8PFOA	11	9
PFNA	target analyte	463 > 419	463>219	33: M9PFNA	16	10
PFDA	target analyte	513 > 469	513>219	38: M6PFDA	21	9
PFUnA	target analyte	563 > 519	563>269	41: M7-PFUDA	22	18
PFDoA	target analyte	613 > 569	613>219	50: MPFDOA	32	10
PFTTrDA	target analyte	663 > 619	663>219	53: M2PFTEDA	30	10
PFTA	target analyte	713 > 669	713>219	53: M2PFTEDA	30	12
PFHxDA	target analyte	813>769	813>219	59: M2PFHxDA	35	15
PFODA	target analyte	913>869	913>219	59: M2PFHxDA	37	15
PFBS	target analyte	299 > 80	299 > 99	7: M3PFBS	28	20
PFPeS	target analyte	349 > 80	349>99	18: M3PFHxS	38	34
L-PFHxS	target analyte	399 > 80	399>99	18: M3PFHxS	20	38
br-PFHxS	target analyte	399 > 80	399>99	18: M3PFHxS	20	38
PFHxS (total)	target analyte	399 > 80	399>99	18: M3PFHxS	20	38
PFHpS	target analyte	449 > 80	449>99	33: M8PFOS	20	31
L-PFOS	target analyte	499 > 80	499>99	33: M8PFOS	18	50
br-PFOS	target analyte	499 > 80	499>99	33: M8PFOS	18	50
PFOS (total)	target analyte	499 > 80	499>99	33: M8PFOS	18	50
PFNS	target analyte	549 > 80	549>99	33:M8PFOS	18	42
PFDS	target analyte	599 > 80	599>99	33:M8PFOS	6	50
PFDoS	target analyte	699>80	699>99	33: M8PFOS	75	68
4:2 FTS	target analyte	327 > 307	327>307	9: M2-4:2FTS	15	20
6:2 FTS	target analyte	427 > 407	427>407	25: M2-6:2FTS	25	20
8:2 FTS	target analyte	527 > 507	527>507	35: M2-8:2FTS	25	20
10:2 FTS	target analyte	627>607	627>81	25: M2-8:2FTS	30	30
PFOSA	target analyte	498 > 78	498>169	29: M8FOSA	12	28
NMeFOSA	target analyte	512>169	512>219	63: d3-NMeFOSA	66	26

Analyte	Type	Quantitation transition	Qualifier transition	IS	CV	CE
NEtFOSA	target analyte	526>169	526>119	61: d5-NEtFOSA	50	28
br-NEtFOSAA	target analyte	584 > 419	584>483	48: d5-NEtFOSAA	41	20
L-NEtFOSAA	target analyte	584 > 419	584>483	48: d5-NEtFOSAA	41	20
NEtFOSAA (total)	target analyte	584 > 419	584>483	48: d5-NEtFOSAA	41	20
NMeFOSE	target analyte	616>59	N/A	66: d7-NMeFOSE	28	14
br-NMeFOSAA	target analyte	570 > 419	570>483	41: D3-NMeFOSAA	38	18
L-NMeFOSAA	target analyte	570 > 419	570>483	41: D3-NMeFOSAA	38	18
NMeFOSAA (total)	target analyte	570 > 419	570>483	41: D3-NMeFOSAA	38	18
NEtFOSE	target analyte	630>59	N/A	67: d9-NEtFOSE	10	12
HFPO-DA	target analyte	285>169	329>285	54: M3HFPO-DA	15	5
ADONA	target analyte	377>251	377>135	23: M8PFOA	22	10
9CIPF3ONS	target analyte	531>351	N/A	33: M8PFOS	18	22
11CIPF3OUdS	target analyte	631>451	N/A	33: M8PFOS	22	26
M2-4:2FTS	EIS	329 > 81	N/A	29:M4PFOS	12	
M2-6:2FTS	EIS	429 > 409	N/A	29:M4PFOS	25	20
M2-8:2FTS	EIS	529 > 509	N/A	29:M4PFOS	25	25
M4PFBA	EIS	217 > 172	N/A	1: M3PFBA	10	8
M5PFPEA	EIS	268 > 223	N/A	1: M3PFBA	10	6
M5PFHxA	EIS	318 > 273	N/A	19:M2PFOA	16	6
M4PFHpA	EIS	367 > 322	N/A	19:M2PFOA	20	8
M8PFOA	EIS	421 > 376	N/A	19: M2PFOA	20	8
M9PFNA	EIS	472 > 427	N/A	19: M2PFOA	18	4
M6PFDA	EIS	519 > 474	N/A	36: M2PFDA	21	9
M7-PFUDA	EIS	570 > 525	N/A	36: M2PFDA	25	11
M2PFTEDA	EIS	715 > 670	N/A	36: M2PFDA	27	17
M2PFHxDA	EIS	815>770	N/A	36:M2PFDA	30	14
MPFDOA	EIS	615 > 570	N/A	36: M2PFDA	31	11
M3HFPO-DA	EIS	287>169	N/A	19: M2PFOA	15	5
M3PFBS	EIS	302 > 80	N/A	29:M4PFOS	30	24
M3PFHxS	EIS	402 > 80	N/A	29:M4PFOS	24	34
M8PFOS	EIS	507 > 80	N/A	29: M4PFOS	49	47
M8FOSA	EIS	506 > 78	N/A	19: M2PFOA	23	33
d9-NEtFOSE	EIS	639>59	N/A	19: M2PFOA	22	74
d7-NMeFOSE	EIS	623>59	N/A	19: M2PFOA	46	52
d5-NEtFOSAA	EIS	589 > 419	N/A	36: M2PFOA	36	20
d5-NEtFOSA	EIS	531>169	N/A	19: M2PFOA	42	26

Analyte	Type	Quantitation transition	Qualifier transition	IS	CV	CE
d3-NMeFOSAA	EIS	573 > 419	N/A	36: M2PFOA	40	18
d3-NMeFOSA	EIS	515>169	N/A	19: M2PFOA	32	28
M3PFBA	surrogate	216>171	N/A		10	8
M2PFOA	surrogate	415 > 370	N/A		20	8
M4PFOS	surrogate	501 > 80	N/A		49	54
M2PFDA	surrogate	515 > 470	N/A		21	9

IS = internal standard, CV = collision voltage, CE = collision energy. Tables includes analytes, extracted internal standards, and surrogate recovery standards used in targeted PFAS and TOP assay analyses.

**Table S6.** Results for replicate total fluorine analyses of 6 products.

Sample ID	Concentration (ppm)	RSD (%) <sup>a</sup>
R7	<10, <10, <10, <10	0.0
S2	<10, <10, <10, <10	0.0
P3	728, 589, 411, 820, 837	26
C1	1820, 1770, 2120, 2080, 1810	8.7
U10	<10, <10, 12, 19	n.c.
C5	<10, <10, <10, 12	n.c.

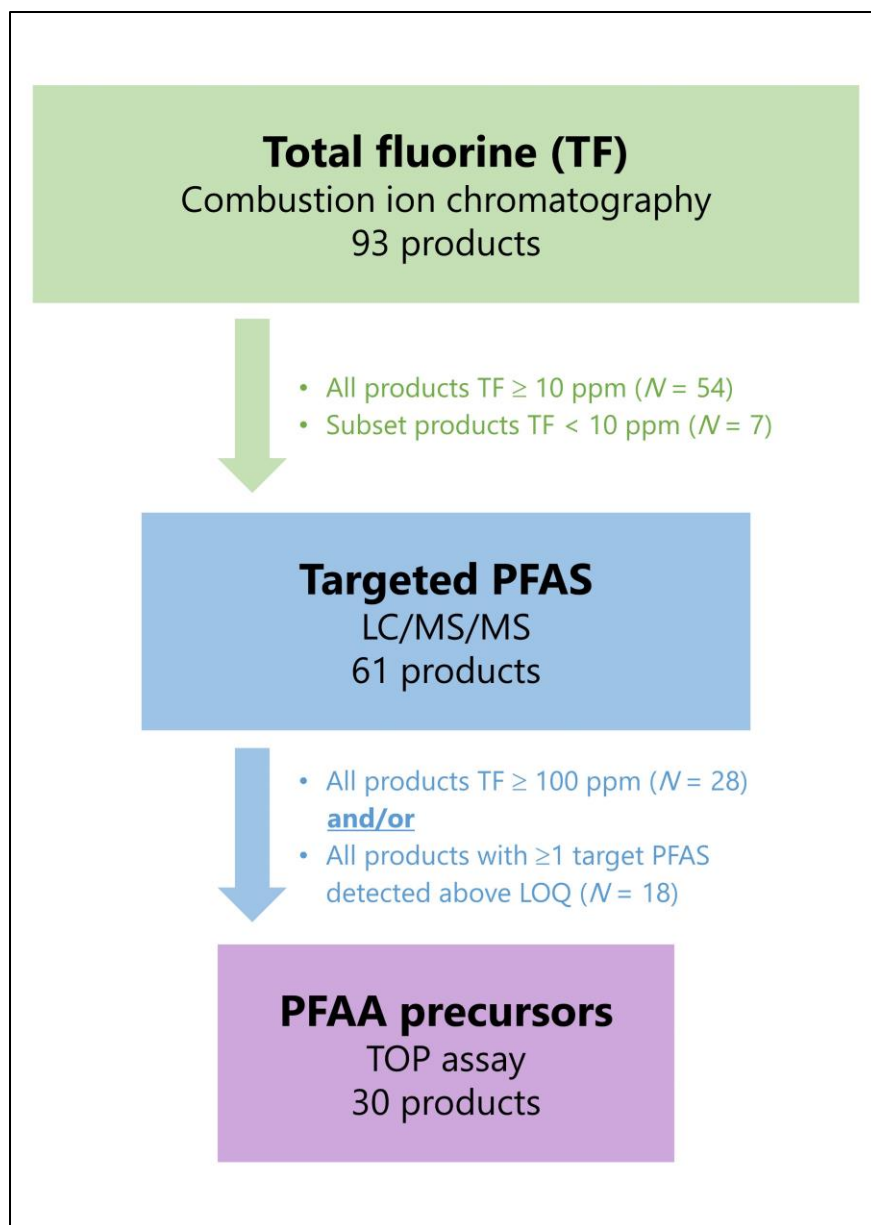
<sup>a</sup> RSD=relative standard deviation. n.c.=not calculated. RSD was calculated only for replicates that did not contain a mix of detect and non-detect values.

**Table S7.** Surrogate recoveries of laboratory control samples, product sample precision, and relative percent difference of PFAS concentrations for duplicate pairs of product samples.

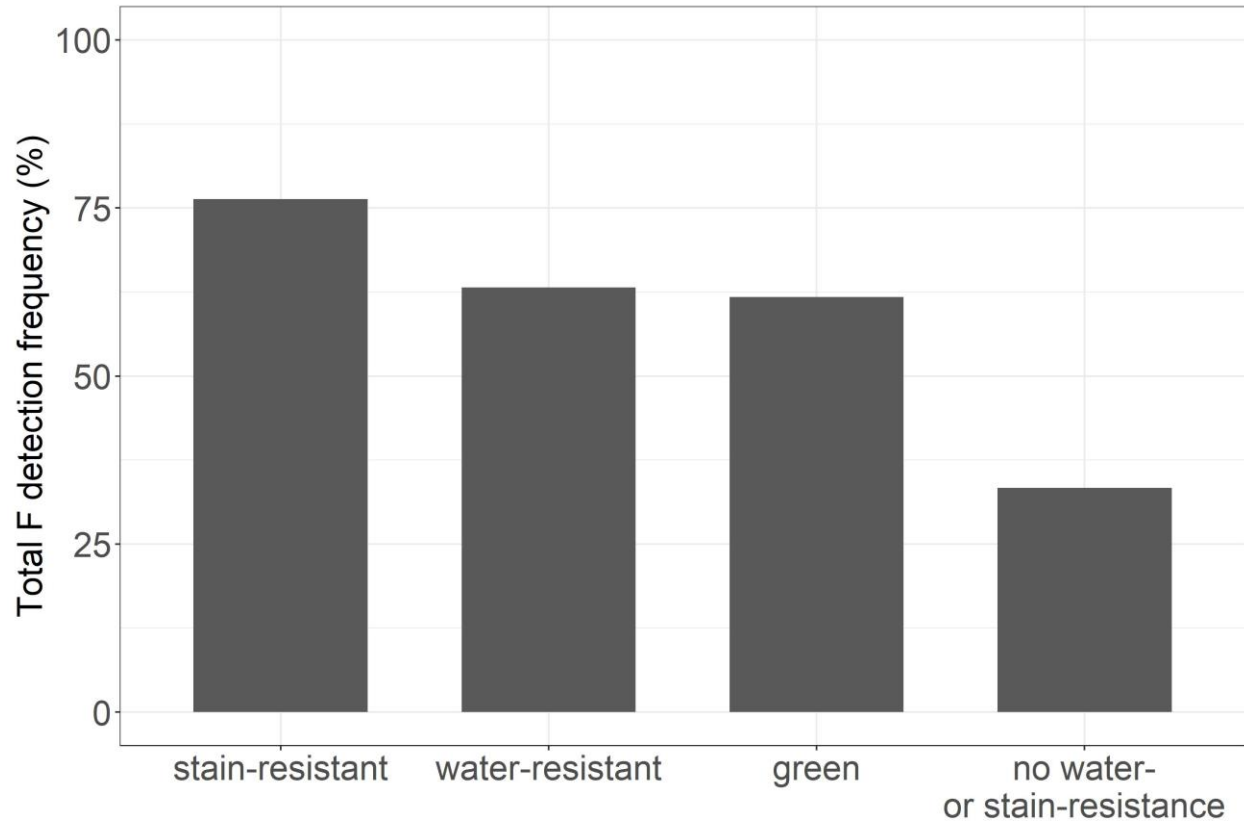
<b>Abbreviation</b>	<b>Avg. LCS<sup>a</sup> recovery (%)</b>	<b>Precision<sup>b</sup></b>	<b>Median RPD (n pairs)<sup>c</sup></b>	<b>Avg. surrogate recovery (%)</b>
<b>PFBA</b>	104.6	5.7	5.2 (4)	
<b>PFPeA</b>	103.8	12.4	6.8 (4)	
<b>PFHxA</b>	109.2	11.7	8.8 (4)	
<b>PFHpA</b>	104.2	4.7	7.1 (2)	
<b>PFOA</b>	114.6	4.5	6.5 (2)	
<b>PFNA</b>	107.2	6.3	11 (2)	
<b>PFDA</b>	103.6	5.7	9.5 (2)	
<b>PFUnA</b>	104.2	7	13 (2)	
<b>PFDoA</b>	104.2	18.2	17 (2)	
PFTrDA	114.6	17.3	21 (1)	
<b>PFTA</b>	106.4	14.3	7.2 (2)	
PFHxDA	108.5	15.4	12 (1)	
PFODA	78	15.8	14 (1)	
<b>PFBS</b>	104	4.7	1.2 (1)	
PFPeS	97.6	13	37 (1)	
PFHxS	104.8	11.4	2.9 (1)	
PFHpS	107.6	9.3	18 (1)	
<b>PFOS</b>	104.2	5.9	NA	
PFNS	109.2	5.9	NA	
PFDS	116.6	15.4	NA	
PFDoDS	135.5	15.4	NA	
4:2 FTS	112.8	13.9	NA	
6:2 FTS	121.2	8.8	NA	
8:2 FTS	112	5.9	NA	
10:2 FTS	158.5	15.4	NA	
FOSA	105.8	15.4	NA	
NMeFOSA	112.3	26.6	NA	
NEtFOSA	107.5	26.6	NA	
NMeFOSAA	109.2	5.9	NA	
NEtFOSAA	100.5	15.4	NA	
<b>NMeFOSE</b>	239.8	27.9	44 (1)	
<b>NEtFOSE</b>	166.3	33.1	NA	
HFPO-DA	111.2	14.1	NA	
ADONA	100	15.4	NA	
9CI-PF3ONS	118	15.4	NA	

11Cl-PF3OUdS	124.2	15.4	NA	
M3HFPO-DA	52			42.07
MPFBA	88			74.3
M3PFBS	92			71.87
M5PFPEA	82			71.65
M2-4:2FTS	61.4			154.8
M5PFHXA	77.8			66.33
M3PFHXS	95.6			79.73
M4PFHPA	84.6			75.13
M2-6:2FTS	69			142.8
D9-NETFOSE	9.833			10.68
D7-NMEFOSE	12.67			12.62
D5-NETFOSAA	87.5			125.7
D3-NMEFOSAA	86.5			113.9
D5-NETFOSA	13.17			13.43
D3-NMEFOSA	10			11.04
M8FOSA	45.25			52.77
M8PFOS	93.6			78.63
M8PFOA	82.2			74.39
M9PFNA	80.2			76.83
M2-8:2FTS	81			153.9
M6PFDA	85.4			77.34
M7-PFUDA	89			83.66
M2PFTEDA	72.2			65.3
M2PFHXDA	100.8			79.67
MPFDOA	81			71.64

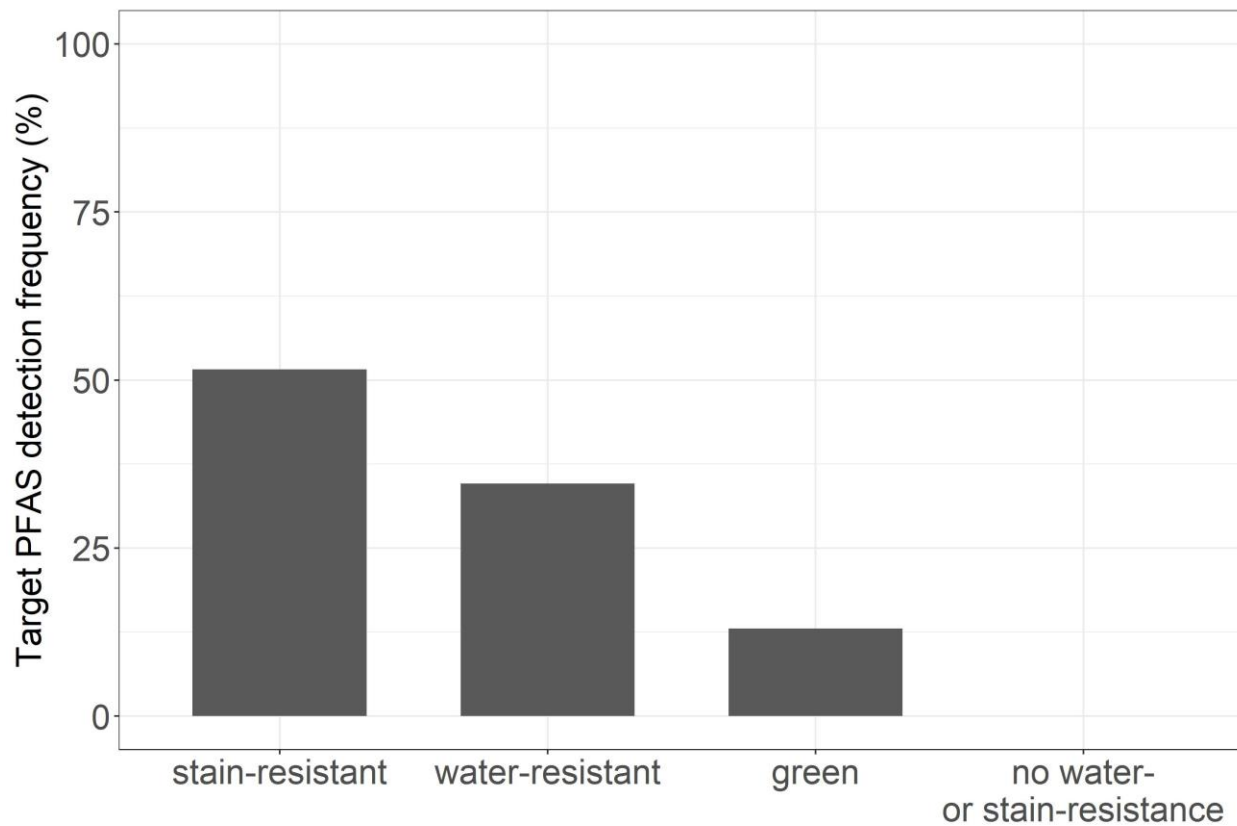
<sup>a</sup> Lab Control Sample recoveries. <sup>b</sup> Precision was calculated for each analyte by dividing the standard deviation of the results by the average of the samples. <sup>c</sup> Median relative percent difference for detected duplicate pairs. Analytes in bold were detected above the method reporting limit in at least one sample.



**Figure S1.** Criteria for selecting products for sample analyses.

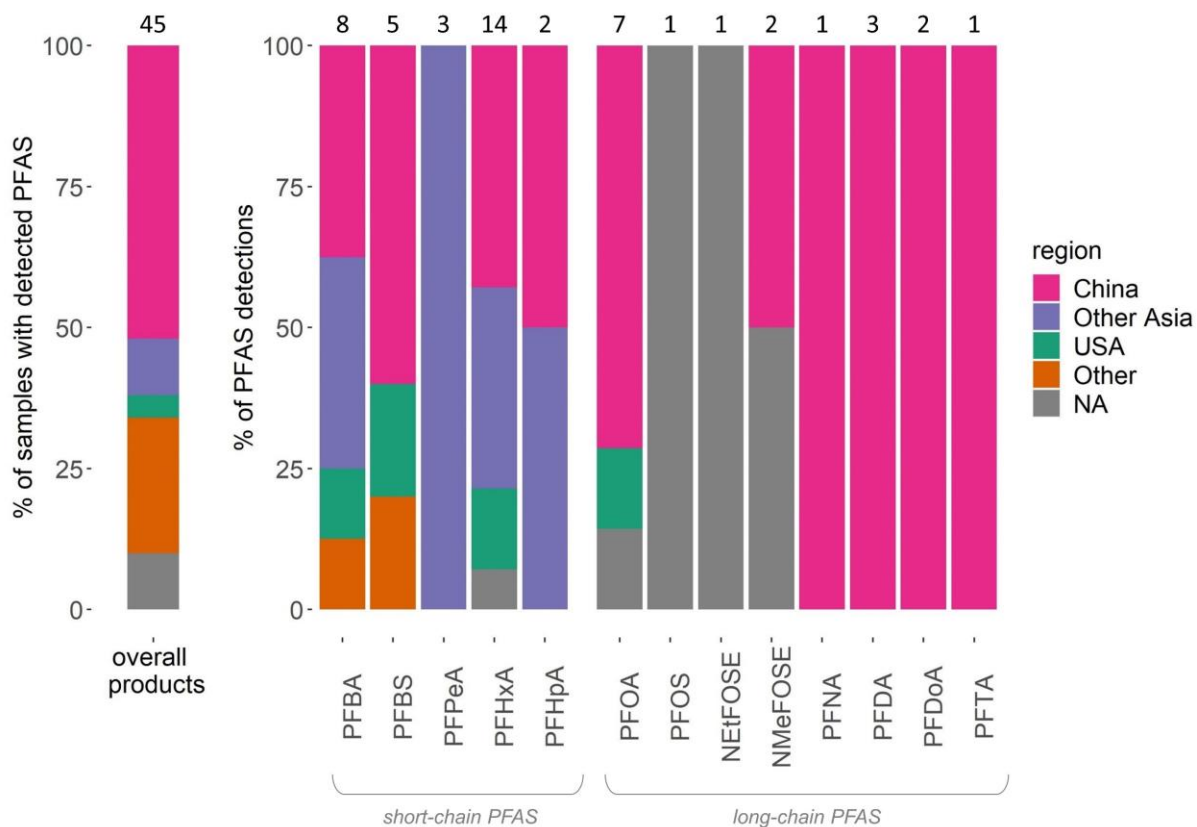


**Figure S2.** Proportion of 93 products with detectable total F ( $\geq 10$  ppm) according to whether or not product information contained any stain-resistant or water-resistant claims or “green” assurances or certifications.



**Figure S3.** Proportion of 61 products with detectable concentrations of at least one PFAS target analyte (above LOQ) according to whether or not product information contained any stain-resistant or water-resistant claims or “green” assurances or certifications.





**Figure S4.** Proportion of 45 products with at least one detectable PFAS target analyte ( $\geq$ LOQ) by region of origin (left) and proportion of each PFAS detected by region of origin (right). Number of samples represented by each bar is shown on top of each bar. All detected PFAS are shown, regardless of quality control measures. Other Asia=Bangladesh, India, Indonesia, Pakistan, Sri Lanka, Vietnam. Other=Egypt, Haiti, Honduras, Kenya, Peru, Turkey. NA = no country of origin indicated.

## REFERENCES

1. Udesky, J. O.; Dodson, R. E.; Perovich, L. J.; Rudel, R. A., Wrangling environmental exposure data: guidance for getting the best information from your laboratory measurements. *Environ Health* **2019**, *18*, (1), 99.