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Perfluorinated compounds in whole fish homogenates from the Ohio, Missouri, and Upper Mississippi Rivers, USA

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Perfluorinated compounds were measured in whole fish homogenates collected from the Ohio, Missouri, and upper Mississippi Rivers in the east central US.

ARTICLE INFO

ABSTRACT

A method for the analysis of 10 perfluorinated compounds (PFCs) in whole fish homogenate is presented and applied to 60 fish samples collected from the Ohio, Missouri, and upper Mississippi Rivers in 2005. Method accuracy ranged between 86 and 125% with limits of quantitation between 0.2 and 10 ng/g wet weight. Intra- and inter-batch precision was generally ±20%. Perfluorooctane sulfonate (PFOS) was the predominant compound identified in these samples, contributing over 80% of total PFC composition in the fish from these rivers, with median PFOS concentrations of 24.4, 31.8, and 53.9 ng/g wet wt in the Missouri, Ohio, and Mississippi Rivers, respectively. Median PFOS levels were significantly ($p = 0.01$) elevated in piscivorous fish (88.0 ng/g) when compared with non-piscivorous fish (15.9 ng/g). The 10 samples with PFOS concentrations above 200 ng/g were broadly scattered across all three rivers, providing evidence of the widespread presence of this compound in these US waterways.

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1. Introduction

Perfluorinated compounds (PFCs) share a number of unique physical and chemical properties that have made them especially useful for a large number of applications, including polymers, paints, water repellents, lubricants, paper coatings, cosmetics, and fire-fighting foams (Prevedouros et al., 2006). While these compounds have been used worldwide for more than 50 years, it has only recently become apparent that perfluorooctane sulfonate (PFOS) and perfluorooctanoic acid (PFOA) are persistent, bio-accumulative, and toxic (Kudo and Kawashima, 2003; Seacat et al., 2003; Kennedy et al., 2004), raising concern about their presence in the environment and potential human exposure.

Preliminary risk assessments conducted for PFOS by the Organisation for Economic Co-operation and Development (2002) and for PFOA by the US Environmental Protection Agency (US EPA) (2002), indicate that these compounds may adversely affect wildlife and human health. Regulations and agreements limiting the production and emission of many PFCs have already been established (US EPA, 2006a), and guidelines for their presence in

drinking and surface water have been issued by authorities in some locations (Minnesota Department of Health, 2007a; State of New Jersey Department of Environmental Protection, 2007).

While a wide range of PFCs have been detected in various environmental and biological matrices, little is known about the distribution and accumulation of these materials in the environment. A limited number of studies have documented the presence of PFCs in fresh water (Hansen et al., 2002; Boulanger et al., 2004; Nakayama et al., 2007) and oceanic samples (Yamashita et al., 2005) and in fish in some locations around the world (Houde et al., 2006), but in general, literature on their distribution remains very sparse.

The majority of the work concerning PFCs in fresh water fish has been based on the analysis of blood or liver samples, which is where the PFCs have been found to concentrate (Giesy and Kannan, 2001; Houde et al., 2006). Although whole fish homogenates have been frequently used in ecological risk assessments involving persistent organic pollutants (Weisbrod et al., 2007), only a small number of studies, centered in the US Great Lakes Region, have used this approach to evaluate the distribution of the PFCs in fish (Martin et al., 2004; Furdui et al., 2007).

At present, limited information is available regarding the performance characteristics of analytical methods for quantifying PFCs in fish tissue. In general, studies have been hampered by analytical challenges such as low PFC concentrations, poor recoveries, lack of

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high quality analytical standards, use of external solvent-based calibration curves, and the unusual physical–chemical properties of perfluoroalkyl chemicals (van Leeuwen et al., 2006). Moreover, most of the work to date has been based on an ion-pair extraction method (Hansen et al., 2001) that may underestimate the actual concentration of many PFCs in unknown samples, most likely due to various matrix effects which inhibit ionization during mass spectrometry (van Leeuwen et al., 2006).

In a recent inter-laboratory study evaluating these methodological shortcomings, agreement between the participating laboratories was shown to be poor for more complex and variable matrices such as fish muscle tissue, fish liver, and water (van Leeuwen et al., 2006). The low percentages of satisfactory PFOS results for fish tissue (17%) and fish liver (55%) suggest that improved extraction and cleanup procedures are needed to better characterize the concentrations of these compounds in fish. In an effort to provide more accurate and effective methods, Taniyasu et al. (2005) have compared the extraction efficiency of the conventional ion-pair method with an alkaline digestion/solid phase extraction (AD/SPE) method using human blood and beaver liver samples as test matrices. They found that compared to the conventional approach, the AD/SPE method led to a 3–5 fold increase in PFOS and other PFC concentrations recovered from the beaver liver samples, indicating substantially greater recoveries and more realistic estimates of the actual values. In this study, we apply a modification of this approach to whole fish homogenate samples in an effort to further characterize its performance and potential utility in dealing with this challenging matrix.

The USEPA has been conducting probabilistic monitoring research under the Environmental Monitoring and Assessment Program (EMAP) (US EPA, 2006b). In 2004–2006, EMAP conducted a probabilistic survey of three great rivers, the Ohio, Missouri, and upper Mississippi Rivers (EMAP GRE), to develop and demonstrate tools to assess the ecological conditions in these important freshwater resources. Fish samples were collected and whole body homogenates were prepared as part of this effort to provide material that could be used to evaluate the bioaccumulation of persistent toxic substances. These samples were also intended to estimate regional and river-wide scales of exposure to contaminants associated with fish consumption for higher trophic levels. Although whole-fish contamination is primarily an indicator of risk to piscivorous wildlife, whole-fish data may also be relevant for estimating human exposure to contaminants through fish consumption.

The objectives of this study were to refine an AD/SPE-based PFC measurement technique for use with whole fish homogenates and to test this method on a subset of samples collected during the 2005 EMAP GRE sampling season along the Ohio, Missouri, and upper Mississippi Rivers. The intent of this work was to validate this method and to provide data on the distribution of PFCs in fish collected from these great river systems.

2. Materials and methods

2.1. Reagents

Standards of perfluorobutane sulfonate (PFBS), perfluorohexane sulfonate (PFHxS), and perfluorooctane sulfonate (PFOS) were provided by 3M Corporation (Saint Paul, MN). Perfluorohexanoic acid (PFHxA), perfluoroheptanoic acid (PFHpA), perfluorooctanoic acid (PFOA), perfluorononanoic acid (PFNA), perfluorodecanoic acid (PFDA), perfluroundecanoic acid (PFUnA), and perfluorododecanoic acid (PFDoA) were purchased from Oakwood Products (West Columbia, SC). Stable isotopically labeled internal standards, $[^{18}O_2]$ammonium perfluorooctane sulfonate and $[1,2-^{13}C_2]$perfluorooctanoic acid, were obtained from Research Triangle Institute (Research Triangle Park, NC) and Perkin Elmer Life and Analytical Sciences (Boston, MA), respectively. The purity of all the analytical standards exceeded 90%. Methanol was purchased from Burdick & Jackson (Muskegon, MI). Sodium hydroxide, sodium acetate, glacial acetic acid, ammonium acetate and ammonium hydroxide were obtained from Sigma–Aldrich Chemical (St Louis, MO). A listing of the target

compounds, acronyms, and related information appears in the Supporting information (Table S1).

2.2. Fish collection

Fish were collected from 30 different sites on the Ohio, Missouri, and Upper Mississippi Rivers (10 sites for each river, Fig. 1) as part of the US EPA's EMAP-GRE program in 2005 (US EPA, 2006b). The 10 sites were selected from each of the three rivers in order to provide equidistant coverage over the entire sample area. Collectively, these sites represent 5120 river kilometers in a 12 state region of the east central United States. Samples were obtained by daytime electrofishing along two 500 m shoreline transects per site. Fish were composited by species, wrapped in foil, and placed in a double plastic bag on ice (US EPA, 2006b). At every sample site, two composite fish samples were collected: a small-fish sample (<120 mm for minnows, dace and shiners; <150 mm for gizzard shad) and a large-fish sample (>120 mm). The small-fish sample was a composite of 20–200 individuals of one species whose adults are small. The large-fish sample included 3–5 individuals of one species whose adults were larger. A listing of the species collected and the designations used in the following analyses appears in Supporting information (Table S2). Each sample was assumed to accurately represent the proportional abundance of the littoral fish assemblage at that site. After transport to the laboratory, the fish composites were frozen at −20 °C until further processing was completed.

2.3. Preparation of whole fish homogenates

After thawing the samples at 4 °C, bigger fish (>100 g) were cut into chunks with a knife, put through a Hobart Food Cutter 84145 (Hobart, Troy, OH), and then homogenized in a Waring Model 701HS Blender (Waring Products, Torrington, CT). Smaller fish were chopped with a knife and then further homogenized in the blender. The homogenates were put into polyethylene containers and refrozen at −20 °C. A subsample of these homogenates was sent to the analytical lab where it was thawed, weighed, diluted 1:1 with water (1.0 mL water per gram fish), and rehomogenized with a Polytron homogenizer (Brinkmann Instruments, Westbury, NY). A 1.0 mL subsample of this homogenate (containing 0.5 g of fish) was placed in a 15 mL polypropylene tube containing 9.0 mL of 10 mM sodium hydroxide/methanol solution and allowed to digest on an orbital shaker for 16 h at room temperature. After digestion, samples were centrifuged at $2000 \times g$ for 5 min using an IEC Centra CL12 centrifuge (Thermo Fisher Scientific Inc., Waltham, MA), and then 1.0 mL of the supernatant was added to 9.0 mL of deionized water prior to SPE cleanup. Oasis WAX 3cc SPE cartridges (Waters Corporation, Milford, MA) were preconditioned with 4.0 mL of 0.1% ammonium hydroxide in methanol followed by 4.0 mL of methanol and 4.0 mL of deionized water. Ten milliliters of the diluted digest supernatant was then loaded onto the cartridge and washed with 4.0 mL of 25 mM sodium acetate buffer (pH 4) followed by 4.0 mL of methanol. The target compounds were eluted with 4.0 mL of 0.1% ammonium hydroxide in methanol. This final eluate was concentrated to 0.5 mL under nitrogen gas using a Turbo-Vap LV evaporator (Caliper Life Sciences, Hopkinton, MA). Samples were prepared by combining 280 µL of the final eluate with 120 µL of 2 mM ammonium acetate buffer in autosampler vials.

2.4. HPLC/MS/MS analysis

PFC analysis was performed using a high-performance liquid chromatography–tandem mass spectrometry (HPLC/MS/MS) system consisting of an Agilent 1100 liquid chromatograph (Agilent Technologies, Palo Alto, CA) interfaced with a PE Sciex API 3000 triple quadrupole mass spectrometer (Applied Biosystems/MDS Sciex, Foster City, CA). A 10 µL aliquot of sample extract was injected and separation was achieved using a Luna C18 (2) 3 × 50 mm column (5 µm, fitted with a C18 guard column, Phenomenex, Torrance, CA) using an isocratic mobile phase of 30% 2 mM ammonium acetate and 70% methanol at a flow rate of 200 µL/min. The MS/MS was operated in multiple reaction monitoring (MRM) mode using the mass transitions listed in Table S1. All concentrations discussed below are presented as ng of PFC/g wet weight of fish.

2.5. Quality assurance and control

Double blanks (containing only mobile phase) were injected prior to running each batch and once every five samples to monitor for potential PFC contamination from the instrument. Method blanks (containing only deionized water) and matrix blanks (containing blank fish matrix) were treated the same as actual samples in order to monitor potential contamination during sample preparation. Matrix blanks were derived from whole body homogenates of ornamental Koi (Cyprinus carpio) from Carolina Biological Supply Company (Burlington, NC). This material was found to have only negligible levels of the target PFCs and was used for the construction of matrix spiked calibration curves and quality control (QC) pools.

Calibration standards and QC pools were prepared by spiking PFC standards directly into Koi homogenate at levels intended to represent the concentrations anticipated in the unknown samples. Pilot tests indicated that PFOS concentrations were 1–3 orders of magnitude higher than the concentrations of the other PFC compounds of interest; therefore, the calibration range for PFOS was
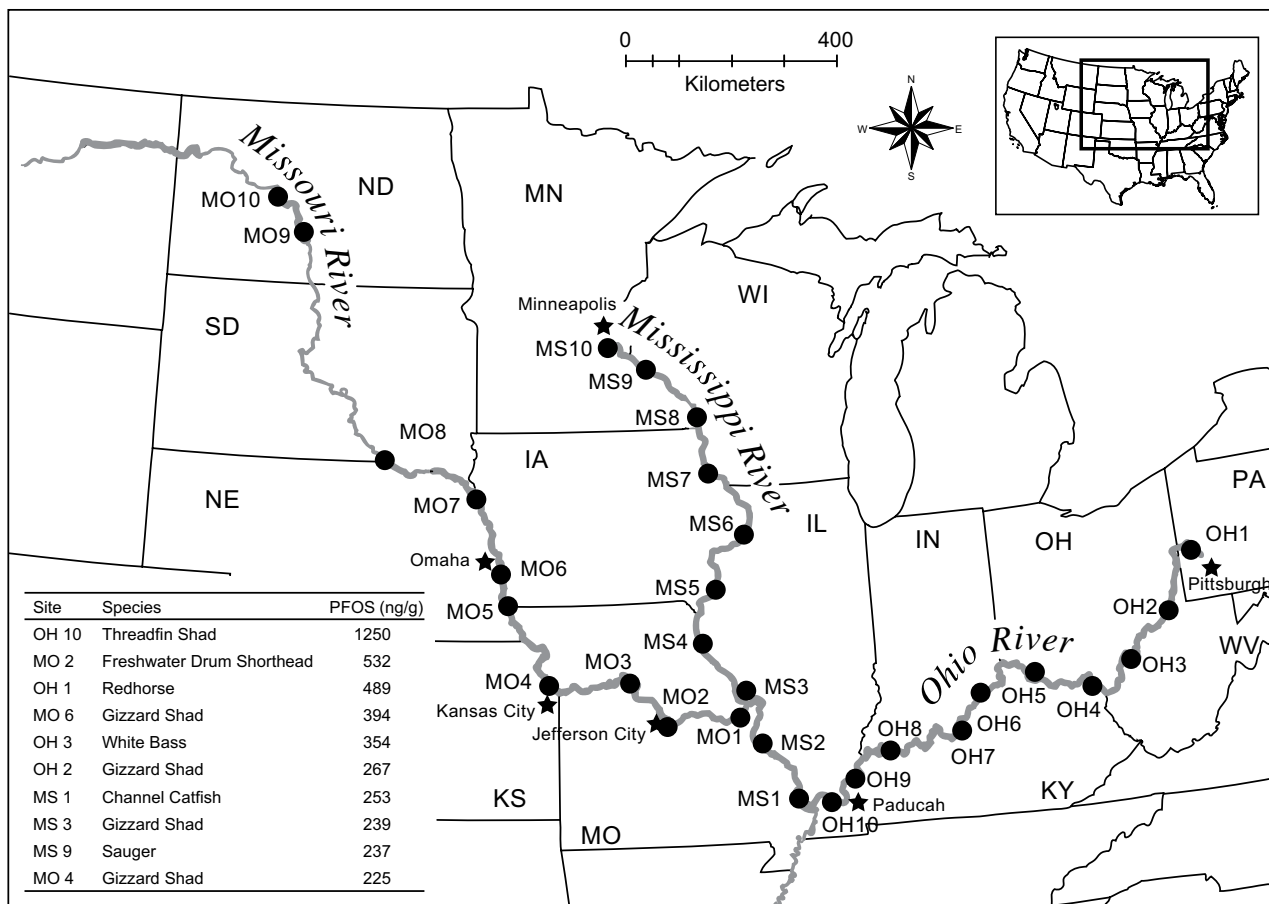
Site	Species	PFOS (ng/g)
OH 10	Threadfin Shad	1250
MO 2	Freshwater Drum Shorthead	532
OH 1	Redhorse	489
MO 6	Gizzard Shad	394
OH 3	White Bass	354
OH 2	Gizzard Shad	267
MS 1	Channel Catfish	253
MS 3	Gizzard Shad	239
MS 9	Sauger	237
MO 4	Gizzard Shad	225

Fig. 1. Sampling sites along the Ohio (OH), Missouri (MO), and Mississippi (MS) Rivers. The accompanying list indicates the species and location of samples with PFOS concentrations higher than 200 ng/g.

correspondingly increased. Calibration curves ranging from 10 to 600 ng/g were prepared daily for PFOS. After the initial analysis, five of the 60 samples had PFOS concentrations greater than 600 ng/g, therefore it was necessary to prepare a new overlapping calibration curve ranging from 200 to 4000 ng/g. For the other nine target PFCs, calibration curves were prepared from 0.2 to 100 ng/g. Separate curves were prepared for PFOS and the other PFCs for each analytical batch. Isotopically labeled PFOA ($^{13}C_2$-PFOA) and PFOS ($^{18}O_2$-PFOS) (5 ng of each) were used as internal standards for the perfluorinated carboxylates and perfluorinated sulfonates, respectively, being added to the fish homogenates before digestion to minimize analytical bias. Quantitation of samples was performed based on the relative response (peak area ratio) of each analyte to the internal standard using calibration curves as described above, adjusting for the known purity of the standards.

QC pools were prepared at 1 ng/g (low), 20 ng/g (medium) and 100 ng/g (high) levels. Separate PFOS QC pools were prepared at 25 ng/g (low), 100 ng/g (medium) and 500 ng/g (high) levels. Multiple QC samples of each concentration were run with the unknown samples so that they represented at least 10% of the samples run in every analytical batch. Determination of inter- and intra-batch precision was based upon these QC data. The limit of quantitation (LOQ) was set as the lowest point on the calibration curve having an accuracy of $100 \pm 30\%$, with accuracy being determined by comparing nominal concentrations of low level standards with their predicted concentrations using the standard curve.

2.6. Statistical analysis

The mean, median, and range were determined for each compound in each river. To allow inclusion of all samples in the statistical analyses, concentrations below the LOQ were assigned a value of one-half of the LOQ. Total PFC concentrations (summation of all concentrations measured) and the percent contribution of each compound to the total were calculated. A nonparametric analysis option in SPSS 13.0 software (SPSS Inc., Chicago, IL) using the Kruskal–Wallis and Mann–Whitney U-tests was used with the significance set at the 0.05 level to compare PFC concentrations between various subgroups of samples.

3. Results

3.1. Method performance characteristics

The LOQ, recovery, and precision for each of the target compounds are presented in Table 1. Mean accuracy for all analytes ranged from 86 to 125% at three different concentration levels. The LOQs ranged from 0.2–1.0 ng/g for all compounds except PFUnA (0.4–1.0 ng/g), PFBS (0.2–4.0 ng/g), PFHpA (0.2–4.0 ng/g), PFHxA (2–10 ng/g), and PFOS (10 ng/g). Intra-batch precision was <20% (coefficient of variation, CV) for all analytes except PFHpA (21.6%) and PFHxA (51.2%) at the lowest level (1 ng/g). Inter-batch precision was <30% (CV) for all compounds except PFDoA (40.1%), PFUnA (39.5%), and PFHxA (42.2%) at the 1 ng/g level. All blank samples (containing either blank mobile phase, deionized water, or blank fish matrix) consistently gave results <LOQ for each compound.

3.2. PFC concentrations in fish

Table 2 is a summary of the PFC concentrations found in the fish samples from each river in this study. One or more PFC was detected in all the samples except one (sample MO 10b), which was collected in the upper reaches of the Missouri River (Fig. 1). Out of 600 assays (60 fish samples × 10 PFCs), 279 (47%) were above the LOQ; 87 (44%) for the Missouri River, 92 (46%) for the Mississippi River, and 100 (50%) for the Ohio River. The least frequently detected compounds were PFBS (3/60 samples) and PFOA (6/60 samples), while the most frequently detected was PFOS (49/60

Table 1
Method performance characteristics of standard-spiked matrix

Analytes	LOQ (ng/g)	Validated range (ng/g)	Correlation coefficient (R)	Accuracy (%), mean ± SD			Average intra-assay precision (CV %)			Average inter-assay precision (CV %)		
				1 ng/g (n = 5)	20 ng/g (n = 5)	100 ng/g (n = 7)	1 ng/g (n = 5)	20 ng/g (n = 5)	100 ng/g (n = 7)	1 ng/g (n = 5)	20 ng/g (n = 5)	100 ng/g (n = 7)
PFOS[a]	10.0	10–4000	0.98	92.4 ± 12	86.8 ± 3.4	105 ± 6.3	5.7	3.1	3.2	13.7	2.8	6.6
PFHxS	0.2	0.2–100	0.99	105 ± 15	109.2 ± 8.4	92.2 ± 4.9	12.7	9.7	3.5	13.2	2.9	4.8
PFBS	0.2–0.4	0.2–100	0.97	103 ± 15	111 ± 5.9	98.3 ± 14	14.1	5.3	5.3	4.7	4.4	16.4
PFDoA	0.2–1.0	0.2–100	0.99	103 ± 33	106 ± 19	105 ± 22	15.8	15.1	13.8	40.1	11.7	17.9
PFUnA	0.4–1.0	0.4–100	0.98	111 ± 29	105 ± 22	107 ± 21	14.4	18.8	12.0	39.5	8.8	17.3
PFDA	0.2–1.0	0.2–100	0.99	104 ± 27	100 ± 13	109 ± 23	18.4	13.8	13.0	28.8	4.4	21.2
PFNA	0.2	0.2–100	0.99	98.7 ± 12	99.6 ± 7.2	111 ± 25	2.1	7.1	10.7	16.2	6.0	25.2
PFOA	0.2–1.0	0.2–100	0.99	85.8 ± 7.9	102 ± 5.1	113 ± 25	2.1	1.9	8.9	9.0	5.9	24.6
PFHpA	0.2–4.0	0.2–100	0.98	100 ± 24	102 ± 7.7	114 ± 24	21.6	5.7	8.7	19.3	7.6	23.7
PFHxA	2.0–10.0	2.0–100	0.94	111 ± 40	105 ± 19	125 ± 29	51.2	1.3	6.8	42.2	9.9	26.9

SD, standard deviation.

[a] PFOS spiked levels: low 25 ng/g, medium 100 ng/g, and high 500 ng/g.

samples) with an overall median concentration of 37.4 ng/g. The median PFOS level in the Mississippi River (53.9 ng/g) was higher than the Ohio River (31.8 ng/g) and Missouri River (24.4 ng/g), although the differences were not statistically significant. The highest concentration of PFOS, 1250 ng/g, was found in threadfin shad (*Dorosoma petenense*) from the Ohio River (sample site OH 10a). Fig. 1 indicates that elevated PFOS concentrations were broadly distributed in this survey, with 10 sites (17%) having samples above 200 ng/g (four samples on the Ohio River, three on the Mississippi River, and three on the Missouri River).

The other PFCs were all generally at least an order of magnitude lower than PFOS, with an overall median concentration for PFHxA at 3.71 ng/g, PFDA at 0.82 ng/g, and PFHxS at 0.36 ng/g. Notably high values of the other PFCs were also broadly dispersed among the rivers, including PFHxS at 8.14 ng/g on the Missouri River (sample MO 6b), PFUnA at 48.0 ng/g on the Mississippi River (sample MS 1a), and PFDA at 9.01 ng/g (samples OH 4a and OH 5a), and PFHxA at 18.4 ng/g (sample OH 4b), both on the Ohio River. It is also interesting to note that the median levels of PFDoA (1.87 ng/g), PFUnA (4.52 ng/g), and PFDA (4.10 ng/g) in fish from the Ohio River

were all significantly higher than those in the Mississippi and Missouri Rivers (*p* < 0.05). The current study is the first we found which determines short chain PFCs (PFHxA, PFHpA and PFBS) in whole fish homogenates. PFHxA was found in the fish from all three rivers, with concentrations ranging from <LOQ to 18.4 ng/g and a median of 3.71 ng/g.

Total PFCs (a summation of the PFC concentrations measured in individual fish) ranged from less than the LOQ to a maximum of 1270 ng/g, with the median individual sample having a total concentration of 53.4 ng/g PFCs. Median total PFC levels tended to be higher in the Mississippi River (66.4 ng/g) compared to the Ohio River (57.6 ng/g) and Missouri River (38.4 ng/g) but these differences were not statistically significant.

3.3. Composition profiles of various PFCs

PFOS was the predominant PFC contributing 86.3% of the total PFC in all rivers, with 72.9% from the Ohio River, 73.1% from the Missouri River, and 91.4% from the Mississippi River (Supporting information, Fig. S1). While PFOS dominated the composition

Table 2
PFC concentrations (ng/g wet weight) by river

River	Statistics	Analyte										Total PFC[c]
		PFOS	PFHxS	PFBS	PFDoA	PFUnA	PFDA	PFNA	PFOA	PFHpA	PFHxA	
MO	N > LOQ[a]	13	13	2	10	2	4	10	0	18	15	87
	Min	<10.0	<0.20	<0.40	<0.20	<1.00	<0.20	<0.20	<1.00	<1.00	<4.00	5.90
	Mean[b]	84.7	1.89	<0.40	0.49	<1.00	0.25	0.43	<1.00	1.53	5.15	95.2
	Median[b]	24.4	0.87	<0.40	<0.20	<1.00	<0.20	<0.20	<1.00	1.80	5.53	38.4
	Max	532	8.14	0.64	2.88	8.60	2.0	1.48	<1.00	4.03	12.6	536
MS	N > LOQ[a]	20	10	1	5	5	18	11	5	3	14	92
	Min	10.0	<0.20	<0.20	<0.40	<0.40	<0.40	<0.20	<0.20	<0.20	<2.00	14.4
	Mean[b]	83.1	0.42	<0.20	<0.40	3.38	1.24	0.78	<0.20	0.27	3.42	93.2
	Median[b]	53.9	0.18	<0.20	<0.40	<0.40	1.05	0.27	<0.20	<0.20	3.06	66.4
	Max	250	3.33	0.29	1.25	48.0	5.16	5.38	0.60	1.83	11.1	286
OH	N > LOQ[a]	16	12	0	15	17	17	14	1	0	8	100
	Min	<10.0	<0.20	<0.40	<1.00	<0.40	<1.00	<0.20	<1.00	<4.00	<10.0	11.3
	Mean[b]	147	0.52	<0.40	1.72	6.57	3.88	1.03	<1.00	<4.00	<10.0	166
	Median[b]	31.8	0.48	<0.40	1.87	4.52	4.10	0.42	<1.00	<4.00	<10.0	57.6
	Max	1250	1.96	<0.40	4.13	45.3	9.01	5.89	2.10	<4.00	18.4	1270
Total	N > LOQ[a]	49	35	3	30	24	39	35	6	21	37	279
	Min	<10.0	<0.20	<0.20	<0.20	<0.40	<0.20	<0.20	<0.20	<0.20	<4.00	5.90
	Mean[b]	105	0.94	<0.20	0.84	3.50	1.79	0.74	<0.20	0.64	4.67	118
	Median[b]	37.4	0.36	<0.20	<0.20	<0.40	0.82	0.30	<0.20	<0.20	3.71	53.4
	Max	1250	8.14	0.64	4.13	48.0	9.01	5.89	2.10	4.03	18.4	1270

MO, Missouri River; MS, Mississippi River; OH, Ohio River. Twenty samples from each river.

[a] Number of fish samples with concentration above LOQ.

[b] For samples under LOQ, half of LOQ was assigned to each compound for the calculation of mean and median.

[c] Total PFC assessed by individual fish, n = 20 for each river.

profiles for most individual sampling sites on each river, distinctive profiles were observed for each individual sample analyzed (Supporting information, Fig. S2). This may reflect differences in water quality and/or species specific uptake and elimination of the target compounds. In the Mississippi River, PFOS consistently contributed the most to total PFCs for all sampling sites. However, the profiles were much more variable on the Ohio and Missouri Rivers, suggesting less relative influence of PFOS or perhaps a greater variety of source inputs. Aside from PFOS, PFHxA was the most prevalent PFC contributing 16.6% of total in the Missouri River and 5.6% of in the Mississippi River. PFUnA and PFDA contributed 10.4 and 9.4%, respectively, of total PFC in the Ohio River.

4. Discussion

In Table S3 the mean PFC concentrations from this study are compared to mean concentrations from a small number of previously reported studies that have also used whole fish homogenates for analysis. The results from studies conducted in the US Great Lakes region (Martin et al., 2004; Kannan et al., 2005; Furdui et al., 2007) were substantially similar to the present work, with PFOS accounting for the majority of the PFCs measured. The highest PFOS concentration in these studies (450 ± 98 ng/g) was found in whole sculpin homogenate from Lake Ontario (Martin et al., 2004), approximately one-half the maximal value found in the present study for threadfin shad from the Ohio River (1250 ng/g). Taken together, these studies provide evidence that PFC contamination in freshwater fish is widespread in portions of the central and eastern US.

Most of the other studies performed to date have measured PFC concentrations in the liver and blood (Taniyasu et al., 2003; Houde et al., 2006), making comparisons to the current study difficult. Liver and blood are the tissues where PFCs have been shown to accumulate, but because humans most often only eat fish fillets, the concentrations in muscle tissue are clearly the most important measurement for assessing potential human exposures. Because the relationship between fish body compartments remain unknown at this time, it is extremely difficult to compare and interpret the results from all of the existing studies. It also makes it difficult to make meaningful generalizations about geographical distributions and the interrelationships between various species. A study which examines the species-specific body disposition of the PFCs, comparing blood, liver, muscle, and whole body homogenate, would help to establish these relationships and make all of the data that have been published more useful.

These compounds are not known to be degraded or transformed in biological systems or the environment and transport between these widely separated drainage basins is unlikely. Considering the widespread occurrence and variable ranges of the PFCs documented in this study, it seems likely that there are many different sources of the PFCs present on these large North American rivers. The limited amount of work that has been done on this subject suggests that point source effluents may be responsible for locally elevated PFC concentrations found in some fish species. For example, in a report from Minnesota (Oliaei et al., 2006), elevated PFC levels were detected in wastewater, river water, sediment, and fish from the Mississippi River area near a chemical manufacturing facility in the Minneapolis–St. Paul area. The scattered nature of the high levels determined in the current study is consistent with this hypothesis, but much further work in this area must be completed to examine this fully.

Among the large fish listed in Table S2, the median PFOS concentration was significantly higher in piscivorous fish (88.0 ng/g) when compared to the non-piscivorous species (15.9 ng/g, $p = 0.01$). This observation is consistent with previous studies which also demonstrate higher levels of PFCs in top predator species in aquatic food chains (Martin et al., 2004; Kannan et al., 2005). It is also interesting to examine the differences between pelagic and benthic feeders. For example, the median PFOS level in shad (primarily benthic feeders) was 232 ng/g, which is almost nine times higher than the pelagic feeding shiner at 26.5 ng/g ($p = 0.08$). While this result only approaches significance in this analysis, it is consistent with a previous study which noted elevated PFC levels in benthic fauna (Martin et al., 2004). While these ecological observations are clearly tentative, being based on only a small number of observations in this study, they illustrate how application of a whole fish homogenate method is useful in evaluating trophic level dynamics and interspecies relationships.

At least one study has suggested that consumption of PFC-contaminated fish may be related to PFC levels in human blood (Falandysz et al., 2006). In recognition of the potential for human exposures via this route, the Minnesota Department of Health has recently issued fish consumption advisories for contaminated sections of the Mississippi River (Minnesota Department of Health, 2007b). This advisory suggests that people limit their intake of fish to no more than one meal a week if PFOS levels in fillets exceed 38 ng/g. It is therefore interesting to note that 48% of the whole fish homogenates examined in this study had PFOS levels that exceeded that threshold. As noted above, the relationship between measurements made with homogenates and fillets has not been examined, but the proportion of samples from this study that exceeded this advisory limit indicates that consumption of fish from this region may be a route of PFC exposure that needs further evaluation. A recent survey of people living in the eight US states that boarder the Great Lakes found that 84% included fish (from any source) in their diet, with the consumption of sport-caught fish being highest in Minnesota (44%) and Wisconsin (39%) (Imm et al., 2005). The same survey estimated that 2.9 million people in the Great Lakes region had more than 104 fish meals per year from all sources (commercial and sport-caught).

5. Conclusion

In summary, a method to measure PFCs in whole fish homogenate using alkaline digestion and solid phase extraction followed by LC/MS/MS analysis has been developed and tested. The performance characteristics described above demonstrate that this method is accurate, precise, and useful in studies intended to evaluate the distribution of PFCs in fish populations. This method was successfully applied to material collected as part of the USEPA's EMAP GRE study and provides evidence of the widespread presence of the PFCs in fish from the three largest rivers in the east central US. It is clear that a great deal of work remains to be done to adequately describe the distribution of these materials in aquatic ecosystems and to estimate potential human exposures resulting from the consumption of fish from this region.

been subjected to agency review and approved for publication. Mention of trade names or commercial products does not constitute endorsement or recommendation for use.

Appendix A. Supplementary data

Supplementary data can be found, in the online version, at doi: 10.1016/j.envpol.2008.03.014.

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