Spatial and Temporal Variation and Tissue Distribution of PFAS Concentrations in Smallmouth Bass

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Smallmouth Bass Health Monitoring and Assessment

- In response to fish kills, observed endocrine disruption (intersex) and population declines in certain areas of the Chesapeake Bay watershed, fish health was assessed at numerous sites from 2013-2021
- In addition to biological endpoints, tissues were saved for chemical analyses
- Initially analyzed archived plasma samples collected at the four integrator sites during 2018 to determine if PFAS might be a concern in these populations
 - 13 PFAS analytes were measured by SGS AXYS

Long Term Integrated Monitoring

• 2013-2019

- Biological monitoring of adults in spring and fall and young-of-year in early summer
- Surface water sampling for hormones, pesticides, phytoestrogens monthly (bimonthly in spring)
- Two in the Potomac (mouth of Antietam Creek and South Branch Potomac near Moorefield)
- Two in the Susquehanna (mouth of Pine Creek and mouth of West Branch Mahantango Creek)



Compounds Detected in Smallmouth Bass Plasma

- Four compounds were detected in every plasma sample
 - PFOS (perfluorooctane sulfonate) 8 carbon sulfonate
 - Ranged from 20 574 ng/ml
 - PFUnA (perfluoroundecanoate) 11 carbon carboxylic acid
 Ranged from 3 55 ng/ml
 - PFDA (perfluorodecanoate) 10 carbon carboxylic acid
 - Ranged from 2 37 ng/ml
 - PFDoA (perfluorododecanoate) 12 carbon carboxylic acid
 Ranged from 1 34 ng/ml
- Additional compounds were detected in some samples
 - PFOSA (perfluorooctane sulfamide) 8 carbon sulfonate
 - Ranged from BD 1.7 ng/ml
 - PFNA (perfluorononanoic acid) 9 carbon carboxylic acid
 - Ranged from BD 1.3 ng/ml

Site Locations Spatial Comparisons

A. Susquehanna River –Pennsylvania Pine Creek Chillisquaque Creek West Branch Mahantango Creek Swatara Creek

- B. West Virginia sites Cheat River South Branch Potomac Moorefield South Branch Potomac Petersburg
- C. Potomac River Maryland Antietam Creek

E. Delaware River - Pennsylvania Little Neshaminy Creek





Spatial Comparison



Land Cover/Land Use Attributes Upstream Catchment

Site	Developed	Agriculture	Forest	EPA - PFAS Facilities
Little Neshaminy	70.7	9.4	15.9	60
Swatara	13.9	39.8	42.2	39
Antietam	17.3	49.1	32.2	25
WB Mahantango	6.8	31.6	60.1	2
Chillisquaque	6.9	59.4	30.5	0
South Branch - P	3.7	15.8	78.7	2
South Branch -M	3.2	14.0	80.6	2
Pine	3.5	8.5	84.3	23
Cheat River	4.5	2.8	84.8	4

Temporal Comparisons



Sex Differences – Combined Years Four Temporal Sites

Males consistently had higher levels, except for PFOS at Antietam Creek

Site	PFOS	PFDA	PFUnA	PFDoA
Pine Creek	ns	ns	p < 0.001	p = 0.0006
WB Mahantango Creek	p = 0.0210	ns	p < 0.001	p < 0.0001
Antietam Creek	p = 0.0278	ns	p < 0.001	p = 0.0022
South Branch Potomac	ns	ns	p < 0.001	p < 0.0001



Tissue Comparisons at Swatara Creek Frequency (%) of Detections

Compound	Plasma	Blood	Liver	Gonad	Muscle
PFOS	100	100	100	100	100
PFDA	100	100	100	82	24
PFUnA	100	100	100	100	71
PFDoA	88	47	47	0	0
PFDS	94	24	18	18	0
PFTeDA	94	12	29	18	0
6:2 FTS	71	0	0	88	0

Tissue Comparisons at Little Neshaminy Creek Frequency (%) of Detections

Compound	Plasma	Blood	Liver	Gonad	Muscle
PFOS	100	100	100	100	100
PFDA	100	100	100	100	100
PFUnA	100	100	100	100	100
PFDoA	100	91	100	100	18
PFDS	100	91	100	100	45
PFTeDA	91	91	100	91	64
6:2 FTS	100	100	100	0	0

Mean Tissue Concentrations Swatara Creek Smallmouth Bass



Mean Tissue Concentrations Little Neshaminy Creek Smallmouth Bass



Conclusions

- Four of the 13 PFAS analyzed for were found in all plasma samples with PFOS having the highest concentrations
- There was spatial variation with the highest concentrations measured in watersheds with the highest percentage of developed and agricultural land cover and the presence of military installations and/or airports
- At four sites there was some temporal variation but often not significant
- Tissue comparisons (40 PFAS analytes) at two sites indicated site differences in the pattern of detections among sites that may be related to exposure routes, habitat, food sources etc.

Ongoing Analyses Identification of Adverse Effects

- **Biological Endpoints**
 - Visible abnormalities, condition factors, age, sex documented
 - Blood was collected for plasma vitellogenin and plasma aliquots frozen for future analyses
 - Histopathology of all the organs
 - Molecular analyses gene expression in liver, gonad, anterior kidney
 - Cheyenne Smith has immune function (macrophages and lymphocytes) and transcript abundance of > 60 genes in the anterior kidney at a subset of sites
 - Exposure to complex mixtures of bioaccumulative chemicals such as mercury, PCBs as well as varying concentrations of pesticides, phytoestrogens, hormones



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